

APPENDIX 1

National Program 301 Plant Genetic Resources, Genomics, and Genetics Improvement

ACCOMPLISHMENT REPORT 2006 – 2011

Research Projects in National Program 301* [By Action Plan Component and Problem Statement]

Component 1: Plant and Microbial Genetic Resource Management

Problem Statement 1A: *Efficiently and Effectively Manage Plant and Microbial Genetic Resources.*

1230-21000-133-00D

EDUCATION AND VISITOR SERVICES AT THE U.S. NATIONAL ARBORETUM – Nancy Luria (P); U.S. National Arboretum, Washington, D.C.

1230-21000-135-00D

ESTABLISH PUBLIC DISPLAY GARDENS FOR WOODY AND HERBACEOUS LANDSCAPE PLANTS – Scott Aker (P); U.S. National Arboretum, Washington, D.C.

1230-21000-137-00D

GENETIC RESOURCES, EVALUATION, AND INFORMATION MANAGEMENT OF WOODY LANDSCAPE PLANT GERMPLASM – Richard Olsen (P), Kevin Conrad, and Mark Roh; U.S. National Arboretum, Washington, D.C.

1275-21000-226-00D

NATIONAL RHIZOBIUM GERMPLASM RESOURCE COLLECTION, GENETIC RESOURCE MANAGEMENT, SYSTEMATICS, AND BIOINFORMATICS – Peter Van Berkum (P); Beltsville, Maryland

1275-21000-227-00D

PLANT EXPLORATIONS TO ACQUIRE CROP GENETIC RESOURCES – Karen Williams (P) and Edward Garvey; Beltsville, Maryland

1275-21000-228-00D

EXCHANGE OF CROP GENETIC RESOURCES AND ASSOCIATED DOCUMENTATION FOR THE U.S. NATIONAL PLANT GERMPLASM SYSTEM – Edward Garvey (p), John Wiersema, and Karen Williams; Beltsville, Maryland

1275-21220-226-00D

OPERATING AND DEVELOPING THE GERMPLASM RESOURCES INFORMATION NETWORK FOR THE U.S. GENETIC RESOURCES PROGRAM – Gary Kinard (P); Beltsville, Maryland

1275-22000-268-00D

THE U.S. NATIONAL FUNGUS COLLECTIONS AND DEVELOPMENT OF ON-LINE RESOURCES ABOUT FUNGI – Amy Rossman (P); Beltsville, Maryland

* Because of the nature of their research, many NP 301 projects contribute to multiple Problem Statements and are listed accordingly. For the sake of consistency, projects are listed and organized in Appendix 1 and 2 according to the ARS project number used to track projects in the Agency's internal database. A (P) after a scientist's name indicates the project's principal investigator.

Problem Statement 1A: *Efficiently and Effectively Manage Plant and Microbial Genetic Resources*
[CONTINUED]

1910-21000-019-00D

CONSERVATION AND CHARACTERIZATION OF GERMPLASM OF SELECTED VEGETABLE CROPS – Larry Robertson (P), Joanne Labate, and Angelo Baldo; Geneva, New York

1910-21000-020-00D

CONSERVATION AND UTILIZATION OF THE GENETIC RESOURCES OF APPLES, GRAPES, AND TART CHERRIES – Gan-Yuan Zhong (P) and Angela Baldo; Geneva, New York

3611-21000-022-00D

CONSERVATION AND UTILIZATION OF MAIZE GENETIC STOCKS – Martin Sachs (P); Urbana, Illinois

3611-21000-023-00D

SOYBEAN GENETIC MANAGEMENT AND UTILIZATION – Steven Clough (P) and Randall Nelson; Urbana, Illinois

3620-22410-012-00D

MANAGEMENT AND GENETIC CHARACTERIZATION OF AGRICULTURAL AND BIOTECHNOLOGICAL MICROBIAL RESOURCES – David Labeda (P), Todd Ward, Stephen Peterson, Cletus Kurtzman, and Kerry Donnell; Peoria, Illinois

3625-21000-053-00D

PLANT GENETIC RESOURCE AND INFORMATION MANAGEMENT – Candice Gardner (P), Charles Block, and Mark Widrechner; Ames, Iowa

3655-21000-051-00D

CONSERVATION AND UTILIZATION OF POTATO GENETIC RESOURCES – John Bamberg (P), David Spooner, and Shelley Jansky; Sturgeon Bay, Wisconsin

5302-21000-010-00D

ARID LAND PLANT GENETIC RESOURCES CONSERVATION, EVALUATION, AND INFORMATION MANAGEMENT – John Preece (P) and Gabriela Romano; Davis, California

5306-21000-018-00D

MANAGEMENT OF GENETIC RESOURCES FOR VITIS, PRUNUS, JUGLANS, FICUS, OLEA, PISTACIA, PUNICA, DIOSPYROS, ACTINIDIA, AND MORUS – John Preece (P) and Mallikarjuna Aradhya; Davis, California

5310-21000-009-00D

MANAGEMENT OF CITRUS AND DATE GENETIC RESOURCES AND INFORMATION – Richard Lee (P) and Robert Krueger; Riverside, California

5320-21000-012-00D

ASSURING LONG TERM PRESERVATION OF TROPICAL CROPS VIA GENETIC RESOURCE MANAGEMENT – Francis Zee (P), Lisa Keith, Tracie Matsumoto Brower, and Dennis Gonsalves; Hilo, Hawaii

5341-21000-004-00D

ARCTIC AND SUBARCTIC PLANT GENETIC RESOURCES CONSERVATION, RESEARCH, AND INFORMATION MANAGEMENT – Nancy Robertson (P), Danny Barney, and Kim Hummer; Palmer, Alaska

5348-21000-022-00D

ACQUISITION, EVALUATION AND CONSERVATION OF TEMPERATE FORAGE LEGUME GENETIC RESOURCES – Stephanie Greene (P) and Jinguo Hu; Prosser, Washington

Problem Statement 1A: *Efficiently and Effectively Manage Plant and Microbial Genetic Resources*
[CONTINUED]

5348-21000-026-00D

RESEARCH, ACQUISITION, MANAGEMENT, AND DOCUMENTATION OF PLANT GENETIC RESOURCES – Jinguo Hu (P), Frank Dugan, Stephen Clement, Richard Johnson, Clarice Coyne, Barbara Hellier, Molly Welsh, Theodore Kisha, and Vicki Bradley; Pullman, Washington

5358-21000-038-00D

MANAGEMENT OF TEMPERATE FRUIT NUT AND SPECIALTY CROP GENETIC RESOURCES – Kim Hummer (P), Nahla Bassil, Barbara Reed, and Joseph Postman; Corvallis, Oregon

5366-21000-027-00D

MANAGEMENT OF NATIONAL SMALL GRAINS COLLECTION RESOURCES – Harold Bockelman (P), John Bonman, and Blair Goates; Aberdeen, Idaho

5402-21000-007-00D

PRESERVATION AND QUALITY ASSESSMENT OF PLANT GENETIC RESOURCES – David Ellis (P), David Dierig, and Maria Jenderek; Fort Collins, Colorado

5402-21000-012-00D

RESEARCH TO DEVELOP STRATEGIES AND TECHNOLOGIES FOR PRESERVING PLANT GENETIC DIVERSITY IN EX SITU GENE BANKS – Christina Walters (P), David Dierig, Gayle Volk, and Christopher Richards; Fort Collins, Colorado

6202-21000-031-00D

COTTON GENETIC RESOURCE MANAGEMENT – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6202-21000-032-00D

CHARACTERIZING AND CONSERVING THE GENETIC DIVERSITY OF CARYA POPULATIONS – Larry Grauke (P) and Tommy Thompson; College Station, Texas

6607-21000-010-00D

CONSERVATION, CHARACTERIZATION, AND EVALUATION OF CROP GENETIC RESOURCES AND ASSOCIATED INFORMATION – Gary Pederson (P), Ming Wang, John Morris, Roy Pittman, Melanie Harrison Dunn, and Robert Jarret; Griffin, Georgia

6631-21000-018-00D

CONSERVATION, CHARACTERIZATION, AND GENETIC IMPROVEMENT OF SUBTROPICAL AND TROPICAL ORNAMENTAL GERMPASM – Alan Meerow (P) and Tomas Ayala Silva; Miami, Florida

6631-21000-019-00D

CONSERVATION AND UTILIZATION OF SUBTROPICAL/TROPICAL FRUIT CROPS, SUGARCANE, AND TRIPSACUM GENETIC RESOURCES – Raymond Schnell (P), David Kuhn, and Tomas Ayala Silva; Miami, Florida

6635-21000-046-00D

SEED INCREASE AND PHYTOSANITARY ASSESSMENT OF QUARANTINED AND TROPICALLY-ADAPTED GENETIC RESOURCES – Ricardo Goenaga; Mayagüez, Puerto Rico

6635-21000-047-00D

GENETIC RESOURCE MANAGEMENT OF TROPICAL/SUBTROPICAL GENETIC RESOURCES AND ASSOCIATED INFORMATION – Brain Irish (P) and Ricardo Goenaga; Mayagüez, Puerto Rico

Problem Statement 1B: *Assess the Systematic Relationships and Genetic Diversity of Crop Genetic Resources.*

1230-21000-050-00D

TAXONOMY AND GENETIC DIVERSITY ASSESSMENT OF LANDSCAPE TREES AND SHRUBS – Alan Whittemore (P) and Joseph Kirkbride; U.S. National Arboretum, Washington, D.C.

1275-21000-228-00D

EXCHANGE OF CROP GENETIC RESOURCES AND ASSOCIATED DOCUMENTATION FOR THE U.S. NATIONAL PLANT GERMPLASM SYSTEM – Edward Garvey (p), John Wiersema, and Karen Williams; Beltsville, Maryland

1275-21000-264-00D

MOLECULAR CHARACTERIZATION AND DIVERSITY ASSESSMENT OF COCOA GERMPLASM IN THE AMERICAS – Dapeng Zhang (P) and Lyndel Meinhardt; Beltsville, Maryland

1910-21000-019-00D

CONSERVATION AND CHARACTERIZATION OF GERMPLASM OF SELECTED VEGETABLE CROPS – Larry Robertson (P), Joanne Labate, and Angelo Baldo; Geneva, New York

1910-21000-020-00D

CONSERVATION AND UTILIZATION OF THE GENETIC RESOURCES OF APPLES, GRAPES, AND TART CHERRIES – Gan-Yuan Zhong (P) and Angela Baldo; Geneva, New York

3611-21000-022-00D

CONSERVATION AND UTILIZATION OF MAIZE GENETIC STOCKS – Martin Sachs (P); Urbana, Illinois

3620-22410-012-00D

MANAGEMENT AND GENETIC CHARACTERIZATION OF AGRICULTURAL AND BIOTECHNOLOGICAL MICROBIAL RESOURCES – David Labeda (P), Todd Ward, Stephen Peterson, Cletus Kurtzman, and Kerry Donnell; Peoria, Illinois

3625-21000-053-00D

PLANT GENETIC RESOURCE AND INFORMATION MANAGEMENT – Candice Gardner (P), Charles Block, and Mark Widrlechner; Ames, Iowa

3655-21000-050-00D

SYSTEMATICS, GENETIC DIVERSITY ASSESSMENT, AND ACQUISITION OF POTATOES, CARROTS, AND THEIR RELATED WILD RELATIVES – David Spooner (P) And Philipp Simon; Madison, Wisconsin

3655-21000-051-00D

CONSERVATION AND UTILIZATION OF POTATO GENETIC RESOURCES – John Bamberg (P), David Spooner, and Shelley Jansky; Sturgeon Bay, Wisconsin

5306-21000-018-00D

MANAGEMENT OF GENETIC RESOURCES FOR VITIS, PRUNUS, JUGLANS, FICUS, OLEA, PISTACIA, PUNICA, DIOSPYROS, ACTINIDIA, AND MORUS – John Preece (P) and Mallikarjuna Aradhya; Davis, California

5310-21000-009-00D

MANAGEMENT OF CITRUS AND DATE GENETIC RESOURCES AND INFORMATION – Richard Lee (P) and Robert Krueger; Riverside, California

5320-21000-012-00D

ASSURING LONG TERM PRESERVATION OF TROPICAL CROPS VIA GENETIC RESOURCE MANAGEMENT – Francis Zee (P), Lisa Keith, Tracie Matsumoto Brower, and Dennis Gonsalves; Hilo, Hawaii

Problem Statement 1B: *Assess the Systematic Relationships and Genetic Diversity of Crop Genetic Resources.* [CONTINUED]

5341-21000-004-00D

ARCTIC AND SUBARCTIC PLANT GENETIC RESOURCES CONSERVATION, RESEARCH, AND INFORMATION MANAGEMENT – Nancy Robertson (P), Danny Barney, and Kim Hummer; Palmer, Alaska

5348-21000-022-00D

ACQUISITION, EVALUATION AND CONSERVATION OF TEMPERATE FORAGE LEGUME GENETIC RESOURCES – Stephanie Greene (P) and Jinguo Hu; Prosser, Washington

5348-21000-026-00D

RESEARCH, ACQUISITION, MANAGEMENT, AND DOCUMENTATION OF PLANT GENETIC RESOURCES – Jinguo Hu (P), Frank Dugan, Stephen Clement, Richard Johnson, Clarice Coyne, Barbara Hellier, Molly Welsh, Theodore Kisha, and Vicki Bradley; Pullman, Washington

5358-21000-038-00D

MANAGEMENT OF TEMPERATE FRUIT NUT AND SPECIALTY CROP GENETIC RESOURCES – Kim Hummer (P), Nahla Bassil, Barbara Reed, and Joseph Postman; Corvallis, Oregon

5366-21000-027-00D

MANAGEMENT OF NATIONAL SMALL GRAINS COLLECTION RESOURCES – Harold Bockelman (P), John Bonman, and Blair Goates; Aberdeen, Idaho

5402-21000-012-00D

RESEARCH TO DEVELOP STRATEGIES AND TECHNOLOGIES FOR PRESERVING PLANT GENETIC DIVERSITY IN EX SITU GENE BANKS – Christina Walters (P), David Dierig, Gayle Volk, and Christopher Richards; Fort Collins, Colorado

6202-21000-031-00D

COTTON GENETIC RESOURCE MANAGEMENT – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6202-21000-032-00D

CHARACTERIZING AND CONSERVING THE GENETIC DIVERSITY OF CARYA POPULATIONS – Larry Grauke (P) and Tommy Thompson; College Station, Texas

6402-21310-003-00D

Genomics and Bioinformatics Research in Agriculturally Important Organisms – Brian Scheffler (P); Stoneville, Mississippi

6607-21000-010-00D

CONSERVATION, CHARACTERIZATION, AND EVALUATION OF CROP GENETIC RESOURCES AND ASSOCIATED INFORMATION – Gary Pederson (P), Ming Wang, John Morris, Roy Pittman, Melanie Harrison Dunn, and Robert Jarret; Griffin, Georgia

6631-21000-018-00D

CONSERVATION, CHARACTERIZATION, AND GENETIC IMPROVEMENT OF SUBTROPICAL AND TROPICAL ORNAMENTAL GERMLASM – Alan Meerow (P) and Tomas Ayala Silva; Miami, Florida

6631-21000-019-00D

CONSERVATION AND UTILIZATION OF SUBTROPICAL/TROPICAL FRUIT CROPS, SUGARCANE, AND TRIPSACUM GENETIC RESOURCES – Raymond Schnell (P), David Kuhn, and Tomas Ayala Silva; Miami, Florida

Problem Statement 1B: *Assess the Systematic Relationships and Genetic Diversity of Crop Genetic Resources.* [CONTINUED]

6635-21000-047-00D

GENETIC RESOURCE MANAGEMENT OF TROPICAL/SUBTROPICAL GENETIC RESOURCES AND ASSOCIATED INFORMATION – Brain Irish (P) and Ricardo Goenaga; Mayagüez, Puerto Rico

6635-21000-049-00D

CONSERVATION, MANAGEMENT, AND GENETIC ENHANCEMENT OF SORGHUM GENETIC RESOURCES AND ASSOCIATED INFORMATION – Hugo Cuevas (P); Mayagüez, Puerto Rico

Component 2: Crop Informatics, Genomics, and Genetic Analyses

Problem Statement 2A: *Genome Database Stewardship and Informatics Tool Development.*

1907-21000-029-00D

DISSECTING COMPLEX TRAITS IN MAIZE AND BIOFUEL GRASSES BY APPLYING GENOMICS, BIOINFORMATICS, AND GENETIC RESOURCES – Edward Buckler IV (P) and Peter Bradbury; Ithaca, New York

1907-21000-031-00D

BIOINFORMATIC METHODS AND TOOLS TO PREDICT SMALL GRAIN FIELD PERFORMANCE – Jean Luc Jannink (P) and Peter Bradbury, Ithaca, New York

1907-21000-030-00D

IDENTIFICATION OF FUNCTIONAL SEQUENCE IN PLANT GENOMES THROUGH BIOINFORMATIC, GENOMIC, AND GENETIC APPROACHES – Doreen Ware (P); Ithaca, New York

3622-21000-032-00D

MODIFICATION OF SOYBEAN SEED COMPOSITION FOR FOOD, FEED, AND OTHER INDUSTRIAL USES – Jan Miernyk (P), Perry Gustafson, Jason Gillman, Melvin Oliver, Kristin Bilyeu, and Hari Krishnan; Columbia, Missouri

3625-21000-051-00D

THE MAIZE GENETICS AND GENOMICS DATABASE – Carolyn Lawrence (P), Craig Abel, and Taner Sen; Ames, Iowa

3625-21000-052-00D

CURATION AND DEVELOPMENT OF THE SOYBEAN BREEDER'S TOOLBOX AND ITS INTEGRATION WITH OTHER PLANT GENOME DATABASES – Randy Shoemaker (P), Rex Nelson, Craig Abel, David Grant, and Steven Cannon; Ames, Iowa

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

3625-21220-004-00D

FUNCTIONAL AND STRUCTURAL GENETIC ANALYSIS OF SOYBEAN AND OTHER LEGUMES – Randy Shoemaker (P), Craig Abel, and Michelle Graham; Ames, Iowa

5325-21000-014-00D

AN INTEGRATED DATABASE AND BIOINFORMATICS RESOURCE FOR SMALL GRAINS – Olin Anderson (P), Yong Gu, Gerard Lazo, and David Matthews; Albany, California

Problem Statement 2A: *Genome Database Stewardship and Informatics Tool Development.*

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6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6225-21220-004-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Stuttgart, Arkansas

6402-21310-003-00D

Genomics and Bioinformatics Research in Agriculturally Important Organisms – Brian Scheffler (P); Stoneville, Mississippi

Problem Statement 2B: *Structural Comparison and Analysis of Crop Genomes.*

1275-21000-263-00D

DEFINING THE GENETIC DIVERSITY AND STRUCTURE OF THE SOYBEAN GENOME AND APPLICATIONS TO GENE DISCOVERY IN SOYBEAN AND WHEAT GERMPLASM – Perry Cregan (P) and David Hyten; Beltsville, Maryland

1275-21220-229-00D

GENOMICS AND PROTEOMICS APPROACHES TO BROADENING RESISTANCE OF SOYBEAN TO PESTS AND PATHOGENS – Benjamin Matthews (P), Bret Cooper, and Mark Tucker; Beltsville, Maryland

1907-21000-029-00D

DISSECTING COMPLEX TRAITS IN MAIZE AND BIOFUEL GRASSES BY APPLYING GENOMICS, BIOINFORMATICS, AND GENETIC RESOURCES – Edward Buckler IV (P) and Peter Bradbury; Ithaca, New York

1907-21000-031-00D

BIOINFORMATIC METHODS AND TOOLS TO PREDICT SMALL GRAIN FIELD PERFORMANCE – Jean Luc Jannink (P) and Peter Bradbury, Ithaca, New York

1907-21000-030-00D

IDENTIFICATION OF FUNCTIONAL SEQUENCE IN PLANT GENOMES THROUGH BIOINFORMATIC, GENOMIC, AND GENETIC APPROACHES – Doreen Ware (P); Ithaca, New York

3622-21000-034-00D

GENETIC MECHANISMS AND MOLECULAR GENETIC RESOURCES FOR MAIZE – Michael McMullen (P), Mary Schaeffer, Melvin Oliver, and Sherry Flint Garcia; Columbia, Missouri

3625-21000-052-00D

CURATION AND DEVELOPMENT OF THE SOYBEAN BREEDER'S TOOLBOX AND ITS INTEGRATION WITH OTHER PLANT GENOME DATABASES – Randy Shoemaker (P), Rex Nelson, Craig Abel, David Grant, and Steven Cannon; Ames, Iowa

3625-21220-004-00D

FUNCTIONAL AND STRUCTURAL GENETIC ANALYSIS OF SOYBEAN AND OTHER LEGUMES – Randy Shoemaker (P), Craig Abel, and Michelle Graham; Ames, Iowa

3655-21000-052-00D

PHYSIOLOGICAL, BIOCHEMICAL AND GENETIC REGULATION OF CARBOHYDRATE METABOLISM IN CEREAL TISSUES – Cynthia Henson (P); Madison, Wisconsin

Problem Statement 2B: *Structural Comparison and Analysis of Crop Genomes*. [CONTINUED]

3655-21000-053-00D

METABOLISM AND ANALYSIS OF CEREAL PHYTOCHEMICALS – Mitchell Wise (P) and Cynthia Henson; Madison, Wisconsin

3655-21430-009-00D

IMPROVEMENT OF BARLEY SEED QUALITY THROUGH MOLECULAR AND FUNCTIONAL GENOMIC ANALYSIS OF GENE EXPRESSION – Ronald Skadsen (P) and Cynthia Henson; Madison, Wisconsin

5325-21000-014-00D

AN INTEGRATED DATABASE AND BIOINFORMATICS RESOURCE FOR SMALL GRAINS – Olin Anderson (P), Yong Gu, Gerard Lazo, and David Matthews; Albany, California

5325-21000-015-00D

GENOMICS AND MOLECULAR APPROACHES TO WHEAT TRAITS – Olin Anderson (P), Yong Gu, and Debbie Chingcuanco; Albany, California

5442-21000-033-00D

IMPROVEMENT OF HARD RED SPRING AND DURUM WHEAT FOR DISEASE RESISTANCE AND QUALITY USING GENETICS AND GENOMICS – Justin Faris (P), Steven Xu, Shunwen Lu, and Shiaoman Chao; Fargo, North Dakota

6202-21000-029-00D

PECAN GENETICS AND IMPROVEMENT – Tommy Thompson (P) and Larry Grauke; College Station, Texas

6202-21000-027-00D

GENETIC ANALYSES AND TRAIT MAPPING TO EXPLOIT UNTAPPED GENETIC DIVERSITY IN SORGHUM – Robert Klein (P); College Station, Texas

6208-21000-017-00D

MOLECULAR AND GENETIC ENHANCEMENT OF ABIOTIC STRESS TOLERANCE IN SORGHUM – Zhanguo Xin, Gloria Burow, and John Burke; Lubbock, Texas

6225-21220-004-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Stuttgart, Arkansas

6406-21000-011-00D

ENHANCING CORN WITH RESISTANCE TO AFLATOXIN CONTAMINATION AND INSECT DAMAGE – William Williams (P), Marilyn Warburton, William Henry, and Gary Windham; Mississippi State, Mississippi

6602-21000-021-00D

GENETIC AND GENOMIC APPROACHES TO IMPROVE PEANUT AND CORN RESISTANCE TO DISEASE AND AFLATOXIN CONTAMINATION – Baozhu Guo (P) and Brian Scully; Tifton, Georgia

6606-21000-002-00D

STONE FRUIT BREEDING AND DEVELOPMENT – Thomas Beckman (P); Byron, Georgia

6645-21220-013-00D

USING GENETIC DIVERSITY OF IMPROVE QUANTITATIVE DISEASE RESISTANCE AND AGRONOMIC TRAITS OF CORN – James Holland (P) and Peter Balint Kurti; Raleigh, North Carolina

Problem Statement 2C: *Genetic Analyses and Mapping of Important Traits.*

1275-21000-263-00D

DEFINING THE GENETIC DIVERSITY AND STRUCTURE OF THE SOYBEAN GENOME AND APPLICATIONS TO GENE DISCOVERY IN SOYBEAN AND WHEAT GERMPLASM – Perry Cregan (P) and David Hyten; Beltsville, Maryland

1907-21000-029-00D

DISSECTING COMPLEX TRAITS IN MAIZE AND BIOFUEL GRASSES BY APPLYING GENOMICS, BIOINFORMATICS, AND GENETIC RESOURCES – Edward Buckler IV (P) and Peter Bradbury; Ithaca, New York

1907-21000-031-00D

BIOINFORMATIC METHODS AND TOOLS TO PREDICT SMALL GRAIN FIELD PERFORMANCE – Jean Luc Jannink (P) and Peter Bradbury, Ithaca, New York

1910-21220-004-00D

IMPROVING GRAPE ROOTSTOCK AND SCION PEST AND DISEASE RESISTANCE – Peter Cousins (P), Lance Cadle Davidson, Angelo Baldo, and Gan-Yuan Zhong; Geneva, New York

1910-21220-005-00D

GENETICS AND GENOMICS OF GRAPE GROWTH, DEVELOPMENT, AND QUALITY – Christopher Owens (P), Gan-Yuan Zhong, and Angelo Baldo; Geneva, New York

3622-21000-034-00D

GENETIC MECHANISMS AND MOLECULAR GENETIC RESOURCES FOR MAIZE – Michael McMullen (P), Mary Schaeffer, Melvin Oliver, and Sherry Flint Garcia; Columbia, Missouri

3625-21000-051-00D

THE MAIZE GENETICS AND GENOMICS DATABASE – Carolyn Lawrence (P), Craig Abel, and Taner Sen; Ames, Iowa

3625-21000-052-00D

CURATION AND DEVELOPMENT OF THE SOYBEAN BREEDER'S TOOLBOX AND ITS INTEGRATION WITH OTHER PLANT GENOME DATABASES – Randy Shoemaker (P), Rex Nelson, Craig Abel, David Grant, and Steven Cannon; Ames, Iowa

3625-21220-004-00D

FUNCTIONAL AND STRUCTURAL GENETIC ANALYSIS OF SOYBEAN AND OTHER LEGUMES – Randy Shoemaker (P), Craig Abel, and Michelle Graham; Ames, Iowa

5306-21000-017-00D

GENETIC DISSECTION OF SEEDLING COLD TOLERANCE IN RICE (ORYZA SATIVA L.) – Thomas Tai (P); Davis, California

5325-21000-014-00D

AN INTEGRATED DATABASE AND BIOINFORMATICS RESOURCE FOR SMALL GRAINS – Olin Anderson (P), Yong Gu, Gerard Lazo, and David Matthews; Albany, California

5325-21000-015-00D

GENOMICS AND MOLECULAR APPROACHES TO WHEAT TRAITS – Olin Anderson (P), Yong Gu, and Debbie Chingcuanco; Albany, California

5325-21430-012-00D

PRODUCTION OF WHEAT GERMPLASM WITH ENHANCED BAKING QUALITY – Ann Blechl (P); Albany, California

Problem Statement 2C: *Genetic Analyses and Mapping of Important Traits*. [CONTINUED]

5325-43000-027-00D

MOLECULAR ANALYSIS OF EFFECTS OF ENVIRONMENT ON WHEAT FLOUR QUALITY AND ALLERGENIC POTENTIAL – William Hurkman (P), William Vensel, and Susan Altenbach; Albany, California

5335-21000-034-00D

POSITIONAL CLONING IN MAIZE OF GENES THAT REGULATE PLANT ARCHITECTURE– Sarah Hake (P); Albany, California

5366-21000-025-00D

PLANT AND SEED CHEMISTRY GENETICS – Victor Raboy (P) and John Bonman; Aberdeen, Idaho

5442-21000-032-00D

GENOMIC RELATIONSHIPS AND CYTOGENETIC MANIPULATION OF DURUM WHEAT BY CLASSICAL AND MOLECULAR TECHNIQUES – Prem Jauhar (P); Fargo, North Dakota

5442-21000-033-00D

IMPROVEMENT OF HARD RED SPRING AND DURUM WHEAT FOR DISEASE RESISTANCE AND QUALITY USING GENETICS AND GENOMICS – Justin Faris (P), Steven Xu, Shunwen Lu, and Shiaoman Chao; Fargo, North Dakota

5442-21220-035-00D

GENETIC IMPROVEMENT OF BARLEY – Lynn Dahleen (P); Fargo, North Dakota

6202-21000-027-00D

GENETIC ANALYSES AND TRAIT MAPPING TO EXPLOIT UNTAPPED GENETIC DIVERSITY IN SORGHUM – Robert Klein (P); College Station, Texas

6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6217-21220-006-00D

ENHANCING DISEASE RESISTANCE AND OIL QUALITY ATTRIBUTES OF PEANUT – Hassan Melouk (P) and Kelly Chamberlin; Stillwater, Oklahoma

6203-21000-001-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Shannon Pinson, and Ming Hsuan Chen; Beaumont, Texas

6208-21000-017-00D

MOLECULAR AND GENETIC ENHANCEMENT OF ABIOTIC STRESS TOLERANCE IN SORGHUM – Zhanguo Xin, Gloria Burow, and John Burke; Lubbock, Texas

6217-21000-007-00D

GENETIC AND GENOMIC APPROACHES TO IMPROVE INSECT RESISTANCE AND OTHER VALUE-ADDED TRAITS IN WHEAT, BARLEY, AND SORGHUM – Yinghua Huang (P), Gary Puterka, and Dolores Mornhinweg; Stillwater, Oklahoma

6225-21000-009-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Rolfe Bryant, Robert Fjellstrom, Helen Miller, and Wengui Yan; Stuttgart, Arkansas

Problem Statement 2C: *Genetic Analyses and Mapping of Important Traits*. [CONTINUED]

6225-21220-004-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Stuttgart, Arkansas

6402-21310-003-00D

Genomics and Bioinformatics Research in Agriculturally Important Organisms – Brian Scheffler (P); Stoneville, Mississippi

6402-21000-031-00D

COMBINING CONVENTIONAL AND MOLECULAR TECHNIQUES TO IMPROVE SEED COTTON VALUE – Jodi Scheffler (P), Lawrence Young, and Rickie Turley; Stoneville, Mississippi

6402-21000-033-00D

GENETIC-PHYSIOLOGICAL TEAM RESEARCH TO IMPROVE PRODUCTION, FIBER QUALITY AND COMPETITIVE ABILITY OF COTTON – Lawrence Young (P), Efrem Bechere, and Linghe Zeng; Stoneville, Mississippi

6404-21000-008-00D

SMALL FRUIT AND ORNAMENTAL GENETIC RESEARCH FOR THE MID-SOUTH – Timothy Rinehart (P), Cecil Pounders, James Spiers, Stephen Stringer, and Hamidou Sakhanokho; Poplarville, Mississippi

6435-21000-016-00D

MOLECULAR ANALYSIS OF DEVELOPMENT TO IMPROVE COTTON FIBER – David Fang (P) and Hee Kim; New Orleans, Louisiana

6602-21000-022-00D

DEVELOPMENT OF IMPROVED PEANUT GERMPLASM AND RESISTANCE TO DISEASE AND NEMATODE PESTS – C. Holbrook Jr (P); Tifton, Georgia

Component 3: Genetic Improvement of Crops

Problem Statement 3A: *Genetic Theory and Methods of Crop Improvement*.

1275-21000-195-00D

GENETIC ENHANCEMENT OF QUALITY CONSTITUENTS IN SOLANACEOUS VEGETABLES – John Stommel (P); Beltsville, Maryland

1910-21000-023-00D

DEVELOPMENT OF PEST, DISEASE RESISTANCE, AND STRESS TOLERANCE IN APPLE ROOTSTOCKS – Gennaro Fazio (P) and Angelo Baldo; Geneva, New York

1907-21000-029-00D

DISSECTING COMPLEX TRAITS IN MAIZE AND BIOFUEL GRASSES BY APPLYING GENOMICS, BIOINFORMATICS, AND GENETIC RESOURCES – Edward Buckler IV (P) and Peter Bradbury; Ithaca, New York

1907-21000-031-00D

BIOINFORMATIC METHODS AND TOOLS TO PREDICT SMALL GRAIN FIELD PERFORMANCE – Jean Luc Jannink (P) and Peter Bradbury; Ithaca, New York

3611-21000-023-00D

SOYBEAN GENETIC MANAGEMENT AND UTILIZATION – Steven Clough (P) and Randall Nelson; Urbana, Illinois

Problem Statement 3A: *Genetic Theory and Methods of Crop Improvement*. [CONTINUED]

3622-21000-034-00D

GENETIC MECHANISMS AND MOLECULAR GENETIC RESOURCES FOR MAIZE – Michael McMullen (P), Mary Schaeffer, Melvin Oliver, and Sherry Flint Garcia; Columbia, Missouri

3622-21000-035-00D

FUNCTIONAL GENOMICS FOR IDENTIFICATION OF SOYBEAN QUALITY TRAIT GENES – Yong Qiang An (P) and Melvin Oliver; Columbia, Missouri

3625-21000-054-00D

IDENTIFICATION OF PLANT GENETIC MECHANISMS THAT INFLUENCE INSECT-POLLINATOR PREFERENCE – Reid Palmer (P) and Craig Abel; Ames, Iowa

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

3635-21220-014-00D

BREEDING SELECTION AND MOLECULAR CHARACTERIZATION FOR IMPROVED SUGAR BEET GERMPLASM – Jon McGrath (P), Renfu Lu, and Linda Hanson; East Lansing, Michigan

5325-21000-015-00D

GENOMICS AND MOLECULAR APPROACHES TO WHEAT TRAITS – Olin Anderson (P), Yong Gu, and Debbie Chingcuanco; Albany, California

5325-21420-004-00D

MOLECULAR GENETIC TOOLS FOR POTATO AND FRUIT TREE IMPROVEMENT – William Belknap (P) and Kent McCue; Albany, California

5347-21410-005-00D

COMMERCIALIZATION OF NEW INDUSTRIAL CROP GERMPLASM AND CROPPING SYSTEMS – Terry Coffelt (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

5347-21000-011-00D

PHYSIOLOGICAL AND GENETIC BASIS OF COTTON ACCLIMATION TO ABIOTIC STRESS – Michael Gore (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

5442-21000-034-00D

SUNFLOWER GERMPLASM DIVERSIFICATION AND CHARACTERIZATION UTILIZING WILD SUNFLOWER SPECIES, CYTOGENETICS, AND APPLIED GENOMICS – Brady Vick (P), Lili Qi, Chao-Chien Jan, and Gerald Seiler; Fargo, North Dakota

5442-21220-024-00D

SUNFLOWER GERMPLASM DEVELOPMENT FOR IMPROVED INSECT AND DISEASE RESISTANCE – Thomas Gulya Jr (P) and Brent Hulke; Fargo, North Dakota

6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6203-21000-001-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Shannon Pinson, and Ming Hsuan Chen; Beaumont, Texas

Problem Statement 3A: *Genetic Theory and Methods of Crop Improvement*. [CONTINUED]

6208-21000-018-00D

CHARACTERIZATION AND ENHANCEMENT OF PLANT RESISTANCE TO WATER-DEFICIT AND THERMAL STRESSES – John Burke (P), James Mahan, Jeffrey Velten, Zhanguo, Paxton Payton, and Junping Chen; Lubbock, Texas

6217-21220-006-00D

ENHANCING DISEASE RESISTANCE AND OIL QUALITY ATTRIBUTES OF PEANUT – Hassan Melouk (P) and Kelly Chamberlin; Stillwater, Oklahoma

6402-21000-031-00D

COMBINING CONVENTIONAL AND MOLECULAR TECHNIQUES TO IMPROVE SEED COTTON VALUE – Jodi Scheffler (P), Lawrence Young, and Rickie Turley; Stoneville, Mississippi

6402-21000-033-00D

GENETIC-PHYSIOLOGICAL TEAM RESEARCH TO IMPROVE PRODUCTION, FIBER QUALITY AND COMPETITIVE ABILITY OF COTTON – Lawrence Young (P), Efrem Bechere, and Linghe Zeng; Stoneville, Mississippi

6402-21220-010-00D

REDUCING SOYBEAN YIELD LOSSES THROUGH GENETIC IMPROVEMENT – Jeffery Ray (P), Lawrence Young, Shuxian Li, and James Smith; Stoneville, Mississippi

6406-21000-012-00D

GERMPLASM ENHANCEMENT AND GENETIC IMPROVEMENT OF COTTON – Johnie Jenkins (P), Sukumar Saha, Jack McCarthy Jr., and Martin Wubben; Mississippi State, Mississippi

6402-21310-003-00D

Genomics and Bioinformatics Research in Agriculturally Important Organisms – Brian Scheffler (P); Stoneville, Mississippi

6602-21000-022-00D

DEVELOPMENT OF IMPROVED PEANUT GERMPLASM AND RESISTANCE TO DISEASE AND NEMATODE PESTS – C. Holbrook Jr (P); Tifton, Georgia

6604-21000-003-00D

PHYSIOLOGICAL MECHANISMS THAT DETERMINE CROP RESPONSE TO IRRIGATION, DISEASE, AND PRODUCTION PRACTICES – Charles Chen (P), Marshall Lamb, and Phat Dang; Dawson, Georgia

6645-21000-028-00D

PHYSIOLOGICAL/BIOCHEMICAL MECHANISMS ASSOCIATED WITH GENETIC ALTERATION OF SOYBEAN QUALITY AND PRODUCTIVITY – Robert Upchurch (P), Earl Taliercio, Thomas Carter Jr., and Joseph Burton; Raleigh, North Carolina

6645-21220-012-00D

INCREASING THE COMPETITIVE POSITION OF U.S. SOYBEANS IN GLOBAL MARKETS THROUGH GENETIC DIVERSITY AND PLANT BREEDING – Thomas Carter (P), Joseph Burton, and Robert Upchurch; Raleigh, North Carolina

6657-21000-006-00D

GENETIC AND CULTURAL PRACTICE IMPROVEMENT FOR SUSTAINABLE COTTON PRODUCTION – Philip Bauer (P) and Benjamin Campbell; Florence, South Carolina

Problem Statement 3B: *Capitalizing on Untapped Genetic Diversity.*

1910-21000-019-00D

CONSERVATION AND CHARACTERIZATION OF GERMPLASM OF SELECTED VEGETABLE CROPS – Larry Robertson (P), Joanne Labate, and Angelo Baldo; Geneva, New York

1910-21000-020-00D

CONSERVATION AND UTILIZATION OF THE GENETIC RESOURCES OF APPLES, GRAPES, AND TART CHERRIES – Gan-Yuan Zhong (P) and Angela Baldo; Geneva, New York

1910-21000-023-00D

DEVELOPMENT OF PEST, DISEASE RESISTANCE, AND STRESS TOLERANCE IN APPLE ROOTSTOCKS – Gennaro Fazio (P) and Angelo Baldo; Geneva, New York

1910-21220-005-00D

GENETICS AND GENOMICS OF GRAPE GROWTH, DEVELOPMENT, AND QUALITY – Christopher Owens (P), Gan-Yuan Zhong, and Angelo Baldo; Geneva, New York

1931-21000-017-00D

GENETIC IMPROVEMENT OF FRUIT CROPS THROUGH FUNCTIONAL GENOMICS AND BREEDING – Ralph Scorza (P), Richard Bell, Ann Callahan, Zongrang Liu, Christopher Dardick, and Chinnathambi Srinivasan; Kearneysville, West Virginia

3602-22000-016-00D

MOLECULAR AND GENETIC MECHANISMS OF HESSIAN FLY RESISTANCE IN SOFT WINTER WHEAT – Richard Shukle (P), Brandon Schemerhorn, and Christopher Williams; West Lafayette, Indiana

3607-21000-012-00D

IMPROVEMENT OF PEST RESISTANCE AND QUALITY TRAITS OF SOYBEAN – Rouf Mian (P) and Margaret Redinbaugh; Wooster, Ohio

3611-21000-022-00D

CONSERVATION AND UTILIZATION OF MAIZE GENETIC STOCKS – Martin Sachs (P); Urbana, Illinois

3611-21000-023-00D

SOYBEAN GENETIC MANAGEMENT AND UTILIZATION – Steven Clough (P) and Randall Nelson; Urbana, Illinois

3620-42000-041-00D

ENHANCED INSECT RESISTANCE AND CROP MANAGEMENT FOR REDUCTION OF MYCOTOXINS IN MIDWEST CORN – Patrick Dowd (P), Eric Johnson, And Alejandro Rooney; Peoria, Illinois

3622-21000-030-00D

GENOMIC CHARACTERIZATION AND MANIPULATION OF VALUE-ADDED TRAITS FROM RYE FOR UTILIZATION IN WHEAT IMPROVEMENT – Perry Gustafson (P); Columbia, Missouri

3625-21000-053-00D

PLANT GENETIC RESOURCE AND INFORMATION MANAGEMENT – Candice Gardner (P), Charles Block, and Mark Widrlechner; Ames, Iowa

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

3635-21430-009-00D

IMPROVED FOOD QUALITY IN DRY BEANS USING GENETIC AND MOLECULAR APPROACHES – Karen Cichy (P); East Lansing, Michigan

Problem Statement 3B: *Capitalizing on Untapped Genetic Diversity*. [CONTINUED]

3625-21000-055-00D

GENETIC ANALYSIS OF SELECTION RESPONSE IN MAIZE POPULATIONS – Marvin Scott (P), Craig Abel, and Jode Edwards; Ames, Iowa

3640-21000-025-00D

GENOMICS, GERMPLASM EVALUATION, AND GENETIC IMPROVEMENT OF OATS – St. Paul, Minnesota

3640-21000-027-00D

IMPROVING RUST AND FHB RESISTANCE IN HARD RED SPRING WHEAT THROUGH GENETICS AND GENOMICS – David Garvin (P) and Carroll Vance; St. Paul, Minnesota

3655-21000-048-00D

ALLIUM, CUCUMIS, AND DAUCUS GERMPLASM ENHANCEMENT, GENETICS, AND BIOCHEMISTRY – Philipp Simon (P), Yiqun Weng, and Michael Havey; Madison, Wisconsin

3655-21000-049-00D

POTATO GENETICS, CYTOGENETICS, DISEASE RESISTANCE, AND PRE-BREEDING UTILIZING WILD AND CULTIVATED SPECIES – Shelley Jansky (P), Philipp Simon, Michael Havey, Paul Bethke, and Dennis Halterman; Madison, Wisconsin

3655-21000-051-00D

CONSERVATION AND UTILIZATION OF POTATO GENETIC RESOURCES – John Bamberg (P), David Spooner, and Shelley Jansky; Sturgeon Bay, Wisconsin

5305-21000-013-00D

SUGARBEET GERMPLASM ENHANCEMENT, BREEDING, AND GENETICS – Kelley Richardson (P) and Frank Martin; Salinas, California

5306-21000-017-00D

GENETIC DISSECTION OF SEEDLING COLD TOLERANCE IN RICE (ORYZA SATIVA L.) – Thomas Tai (P); Davis, California

5306-21000-018-00D

MANAGEMENT OF GENETIC RESOURCES FOR VITIS, PRUNUS, JUGLANS, FICUS, OLEA, PISTACIA, PUNICA, DIOSPYROS, ACTINIDIA, AND MORUS – John Preece (P) and Mallikarjuna Aradhya; Davis, California

5310-21000-009-00D

MANAGEMENT OF CITRUS AND DATE GENETIC RESOURCES AND INFORMATION – Richard Lee (P) and Robert Krueger; Riverside, California

5320-21000-012-00D

ASSURING LONG TERM PRESERVATION OF TROPICAL CROPS VIA GENETIC RESOURCE MANAGEMENT – Francis Zee (P), Lisa Keith, Tracie Matsumoto Brower, and Dennis Gonsalves; Hilo, Hawaii

5335-21000-034-00D

POSITIONAL CLONING IN MAIZE OF GENES THAT REGULATE PLANT ARCHITECTURE – Sarah Hake (P); Albany, California

5341-21000-004-00D

ARCTIC AND SUBARCTIC PLANT GENETIC RESOURCES CONSERVATION, RESEARCH, AND INFORMATION MANAGEMENT – Nancy Robertson (P), Danny Barney, and Kim Hummer; Palmer, Alaska

Problem Statement 3B: *Capitalizing on Untapped Genetic Diversity*. [CONTINUED]

5347-21410-005-00D

COMMERCIALIZATION OF NEW INDUSTRIAL CROP GERMPLASM AND CROPPING SYSTEMS – Terry Coffelt (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

5348-21000-022-00D

ACQUISITION, EVALUATION AND CONSERVATION OF TEMPERATE FORAGE LEGUME GENETIC RESOURCES – Stephanie Greene (P) and Jinguo Hu; Prosser, Washington

5348-21000-023-00D

CHARACTERIZATION OF STRESS RESISTANCE GENES AND MECHANISMS, AND IMPROVEMENT AND GENOTYPING OF WHEAT AND BARLEY GERMPLASM FOR THE WESTERN U.S. – Camille Steber (P), Deven See, Daniel Skinner, and Kimberly Garland Campbell; Pullman, Washington

5348-21000-026-00D

RESEARCH, ACQUISITION, MANAGEMENT, AND DOCUMENTATION OF PLANT GENETIC RESOURCES – Jinguo Hu (P), Frank Dugan, Stephen Clement, Richard Johnson, Clarice Coyne, Barbara Hellier, Molly Welsh, Theodore Kisha, and Vicki Bradley; Pullman, Washington

5348-21430-004-00D

GENETIC IMPROVEMENT OF FREEZING TOLERANCE IN WINTER WHEAT – Daniel Skinner (P); Pullman, Washington

5358-21000-037-00D

PHYSIOLOGY AND GENETIC IMPROVEMENT OF SMALL FRUIT CROPS – David Bryla (P) and Chad Finn; Corvallis, Oregon

5358-21000-038-00D

MANAGEMENT OF TEMPERATE FRUIT NUT AND SPECIALTY CROP GENETIC RESOURCES – Kim Hummer (P), Nahla Bassil, Barbara Reed, and Joseph Postman; Corvallis, Oregon

5366-21000-027-00D

MANAGEMENT OF NATIONAL SMALL GRAINS COLLECTION RESOURCES – Harold Bockelman (P), John Bonman, and Blair Goates; Aberdeen, Idaho

5366-21000-028-00D

SMALL GRAINS GENETICS AND GERMPLASM ENHANCEMENT – John Bonman (P), Donald Obert, Paul Bregitzer, and Eric Jackson; Aberdeen, Idaho

5430-21000-006-00D

GENETIC ENHANCEMENT FOR RESISTANCE TO BIOTIC AND ABIOTIC STRESSES IN HARD WINTER WHEAT – Robert Bowden (P), Ming Shum, Guihua Bai, and John Fellers; Manhattan, Kansas

5440-21000-029-00D

GENETIC IMPROVEMENT AND EVALUATION OF HARD WINTER AND SPRING WHEATS – Robert Graybosch (P); Lincoln, Nebraska

5440-21220-027-00D

ENHANCEMENT OF SORGHUM FOR BIOENERGY, FEED, AND FOOD VALUE – Jeffrey Pedersen (P), Scott Sadler, and Deanna Funnel-Harris; Lincoln, Nebraska

5442-21000-034-00D

SUNFLOWER GERMPLASM DIVERSIFICATION AND CHARACTERIZATION UTILIZING WILD SUNFLOWER SPECIES, CYTOGENETICS, AND APPLIED GENOMICS – Brady Vick (P), Lili Qi, Chao-Chien Jan, and Gerald Seiler; Fargo, North Dakota

Problem Statement 3B: *Capitalizing on Untapped Genetic Diversity*. [CONTINUED]

5442-21220-024-00D

SUNFLOWER GERMPLASM DEVELOPMENT FOR IMPROVED INSECT AND DISEASE RESISTANCE – Thomas Gulya Jr (P) and Brent Hulke; Fargo, North Dakota

6202-21000-029-00D

PECAN GENETICS AND IMPROVEMENT – Tommy Thompson (P) and Larry Grauke; College Station, Texas

6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6202-21000-031-00D

COTTON GENETIC RESOURCE MANAGEMENT – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6202-21000-032-00D

CHARACTERIZING AND CONSERVING THE GENETIC DIVERSITY OF CARYA POPULATIONS – Larry Grauke (P) and Tommy Thompson; College Station, Texas

6203-21000-001-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Shannon Pinson, and Ming Hsuan Chen; Beaumont, Texas

6225-21000-009-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Rolfe Bryant, Robert Fjellstrom, Helen Miller, and Wengui Yan; Stuttgart, Arkansas

6225-21220-004-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Stuttgart, Arkansas

6402-21000-031-00D

COMBINING CONVENTIONAL AND MOLECULAR TECHNIQUES TO IMPROVE SEED COTTON VALUE – Jodi Scheffler (P), Lawrence Young, and Rickie Turley; Stoneville, Mississippi

6402-21000-033-00D

GENETIC-PHYSIOLOGICAL TEAM RESEARCH TO IMPROVE PRODUCTION, FIBER QUALITY AND COMPETITIVE ABILITY OF COTTON – Lawrence Young (P), Efrem Bechere, and Linghe Zeng; Stoneville, Mississippi

6406-21000-012-00D

GERMPLASM ENHANCEMENT AND GENETIC IMPROVEMENT OF COTTON – Johnie Jenkins (P), Sukumar Saha, Jack McCarthy Jr., and Martin Wubben; Mississippi State, Mississippi

6404-21000-008-00D

SMALL FRUIT AND ORNAMENTAL GENETIC RESEARCH FOR THE MID-SOUTH – Timothy Rinehart (P), Cecil Pounders, James Spiers, Stephen Stringer, and Hamidou Sakhanokho; Poplarville, Mississippi

6406-21000-011-00D

ENHANCING CORN WITH RESISTANCE TO AFLATOXIN CONTAMINATION AND INSECT DAMAGE – William Williams (P), Marilyn Warburton, William Henry, and Gary Windham; Mississippi State, Mississippi

Problem Statement 3B: *Capitalizing on Untapped Genetic Diversity*. [CONTINUED]

6602-21000-022-00D

DEVELOPMENT OF IMPROVED PEANUT GERMPLASM AND RESISTANCE TO DISEASE AND NEMATODE PESTS – C. Holbrook Jr (P); Tifton, Georgia

6602-21220-014-00D

GENETIC IMPROVEMENT OF MAIZE AND PEARL MILLET FOR RESISTANCE TO INSECTS AND AFLATOXIN – Jeffrey Wilson (P), C Holbrooke Jr., and Xinzhi Ni; Tifton, Georgia

6604-21000-003-00D

PHYSIOLOGICAL MECHANISMS THAT DETERMINE CROP RESPONSE TO IRRIGATION, DISEASE, AND PRODUCTION PRACTICES – Charles Chen (P), Marshall Lamb, and Phat Dang; Dawson, Georgia

6607-21000-010-00D

CONSERVATION, CHARACTERIZATION, AND EVALUATION OF CROP GENETIC RESOURCES AND ASSOCIATED INFORMATION – Gary Pederson (P), Ming Wang, John Morris, Roy Pittman, Melanie Harrison Dunn, and Robert Jarret; Griffin, Georgia

6615-21000-010-00D

FUNCTIONAL GENOMICS OF AGRONOMIC TRAITS IN DEVELOPING SEED AND POLLEN IN MAIZE AND SORGHUM – Prem Chourey (P), Alisa Huffaker, and Peter Teal; Gainesville, Florida

6631-21000-017-00D

DEVELOPMENT OF AN INTERNATIONAL MARKER-ASSISTED SELECTION PROGRAM FOR CACAO – Raymond Schnell (P) and David Kuhn; Miami, Florida

6631-21000-018-00D

CONSERVATION, CHARACTERIZATION, AND GENETIC IMPROVEMENT OF SUBTROPICAL AND TROPICAL ORNAMENTAL GERMPLASM – Alan Meerow (P) and Tomas Ayala Silva; Miami, Florida

6631-21000-019-00D

CONSERVATION AND UTILIZATION OF SUBTROPICAL/TROPICAL FRUIT CROPS, SUGARCANE, AND TRIPSACUM GENETIC RESOURCES – Raymond Schnell (P), David Kuhn, and Tomas Ayala Silva; Miami, Florida

6635-21000-047-00D

GENETIC RESOURCE MANAGEMENT OF TROPICAL/SUBTROPICAL GENETIC RESOURCES AND ASSOCIATED INFORMATION – Brain Irish (P) and Ricardo Goenaga; Mayagüez, Puerto Rico

6635-21000-049-00D

CONSERVATION, MANAGEMENT, AND GENETIC ENHANCEMENT OF SORGHUM GENETIC RESOURCES AND ASSOCIATED INFORMATION – Hugo Cuevas (P); Mayagüez, Puerto Rico

6645-21220-012-00D

INCREASING THE COMPETITIVE POSITION OF U.S. SOYBEANS IN GLOBAL MARKETS THROUGH GENETIC DIVERSITY AND PLANT BREEDING – Thomas Carter (P), Joseph Burton, and Robert Upchurch; Raleigh, North Carolina

6645-22000-016-00D

EVALUATION AND IMPROVEMENT OF CEREAL GERMPLASM FOR DISEASE RESISTANCE AND WINTER-HARDINESS – Gina Brown Guedira (P), Christina Cowger, David Marshall, and David Livingstone; Raleigh, North Carolina

6657-21000-006-00D

GENETIC AND CULTURAL PRACTICE IMPROVEMENT FOR SUSTAINABLE COTTON PRODUCTION – Philip Bauer (P) and Benjamin Campbell; Florence, South Carolina

Problem Statement 3C: *Germplasm Enhancement/Release of Improved Genetic Resources and Varieties.*

1230-21000-051-00D

GENETICS, GENETIC RESOURCE EVALUATION, AND GENETIC IMPROVEMENT OF LANDSCAPE TREES AND SHRUBS – Margaret Pooler (P); U.S. National Arboretum, Washington, D.C.

1230-21000-052-00D

GENETICS, GENETIC IMPROVEMENT, AND IMPROVED PRODUCTION EFFICIENCY OF NURSERY CROPS – Sandra Reed (P) and Donna Fare; McMinnville, Tennessee

1230-21000-053-00D

GENETIC IMPROVEMENT OF LANDSCAPE TREES FOR DISEASE AND PEST TOLERANCE, NON-INVASIVENESS, AND ORNAMENTAL TRAITS – Richard Olsen (P); U.S. National Arboretum, Washington, D.C.

1275-21000-190-00D

GENETIC ENHANCEMENT OF POTATOES FOR NUTRITIONAL AND PROCESSING QUALITY AND FOR RESISTANCE TO DISEASES – Kathleen Haynes (P); Beltsville, Maryland

1275-21220-189-00D

ENHANCEMENT OF SMALL FRUIT GERMPLASM THROUGH GENOMIC CHARACTERIZATION AND GENETIC IMPROVEMENT WITH EMPHASIS ON DISEASE RESISTANCE – Mark Ehlenfeldt (P), Kimberly Lewers, James Polashock, And Shiow Wang; Beltsville, Maryland

1275-22000-261-00D

DISCOVERING AND USING DISEASE RESISTANCE GENES IN PHASEOLUS VULGARIS FOR THE CONTROL OF RUSTS – Marcial Pastor Corrales (P); Beltsville, Maryland

1910-21000-023-00D

DEVELOPMENT OF PEST, DISEASE RESISTANCE, AND STRESS TOLERANCE IN APPLE ROOTSTOCKS – Gennaro Fazio (P) and Angelo Baldo; Geneva, New York

1910-21220-004-00D

IMPROVING GRAPE ROOTSTOCK AND SCION PEST AND DISEASE RESISTANCE – Peter Cousins (P), Lance Cadle Davidson, Angelo Baldo, and Gan-Yuan Zhong; Geneva, New York

1931-21000-017-00D

GENETIC IMPROVEMENT OF FRUIT CROPS THROUGH FUNCTIONAL GENOMICS AND BREEDING – Ralph Scorza (P), Richard Bell, Ann Callahan, Zongrang Liu, Christopher Dardick, and Chinnathambi Srinivasan; Kearneysville, West Virginia

3607-21000-012-00D

IMPROVEMENT OF PEST RESISTANCE AND QUALITY TRAITS OF SOYBEAN – Rouf Mian (P) and Margaret Redinbaugh; Wooster, Ohio

3611-21000-023-00D

SOYBEAN GENETIC MANAGEMENT AND UTILIZATION – Steven Clough (P) and Randall Nelson; Urbana, Illinois

3625-21000-050-00D

GERMPLASM ENHANCEMENT OF MAIZE PROJECT (GEM) – Michael Blanco (P) and Candice Gardner; Ames, Iowa

3625-21000-056-00D

BREEDING HIGH-QUALITY CORN FOR LOW-INPUT AND ORGANIC FARMING SYSTEMS –Craig Abel; Ames, Iowa

Problem Statement 3C: *Germplasm Enhancement/Release of Improved Genetic Resources and Varieties*. [CONTINUED]

3655-21220-001-00D

CRANBERRY GENETIC IMPROVEMENT AND INSECT PEST MANAGEMENT— Philipp Simon (P), Juan Zalapa, and Shawn Steffan; Madison, Wisconsin

5302-21220-005-00D

IMPROVEMENT OF PRUNUS AND VITIS SCIONS FOR FRUIT QUALITY AND PEST RESISTANCE – Craig Ledbetter (P) and David Ramming; Parlier, California

5305-21000-013-00D

SUGARBEET GERMPLASM ENHANCEMENT, BREEDING, AND GENETICS – Kelley Richardson (P) and Frank Martin; Salinas, California

5305-21220-004-00D

EVALUATION, ENHANCEMENT, GENETICS AND BREEDING OF LETTUCE, SPINACH, AND MELON – James McCreight (P), Beiquan Mou, Ivan Simko, and Ryan Hayes; Salinas, California

5347-21410-005-00D

COMMERCIALIZATION OF NEW INDUSTRIAL CROP GERMPLASM AND CROPPING SYSTEMS – Terry Coffelt (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

5348-21000-024-00D

COOL SEASON GRAIN LEGUME GENETIC ENHANCEMENT AND PATHOLOGY – George Vandemark (P), Kevin McPhee, and Weidong Chen; Pullman, Washington

5354-21000-013-00D

TESTING OF ADVANCED POTATO GERMPLASM FOR BIOTIC AND ABIOTIC RESISTANCES, AND PROFITABILITY COMPONENTS – Charles Brown (P); Prosser, Washington

5354-21220-010-00D

POTATO GERMPLASM ENHANCEMENT THROUGH TRAIT DISCOVERY, GENETIC EVALUATION AND INCORPORATION – Duroy Navarre (P), James Crosslin, and Charles Brown; Prosser, Washington

5354-21220-016-00D

DEVELOP STRESS-RESISTANT DRY BEAN GERMPLASM AND SUSTAINABLE PEST MANAGEMENT STRATEGIES FOR EDIBLE LEGUMES – Phillip Miklas (P), Lydon Porter, and Richard Porter; Prosser, Washington

5358-21000-037-00D

PHYSIOLOGY AND GENETIC IMPROVEMENT OF SMALL FRUIT CROPS – David Bryla (P) and Chad Finn; Corvallis, Oregon

5358-21000-040-00D

GENOMICS, GERMPLASM DEVELOPMENT AND IPM OF HOP – John Henning (P), David Gent, and Gary Banowetz; Corvallis, Oregon

5366-21000-026-00D

DEVELOPMENT OF POTATO VARIETIES AND GERMPLASM WITH IMPROVED RESISTANCES, PRODUCTION EFFICIENCIES, AND TUBER QUALITIES FOR THE WESTERN U.S. – Richard Novy (P), Jonathan Whitworth, and John Bonman; Aberdeen, Idaho

5368-21220-003-00D

SUGARBEET GERMPLASM DEVELOPMENT AND INNOVATIVE GENETIC AND MANAGEMENT APPROACHES TO REDUCING LOSSES CAUSED BY PATHOGENS – Carl Strausbaugh (P) and Imad Eujayl; Kimberly, Idaho

Problem Statement 3C: *Germplasm Enhancement/Release of Improved Genetic Resources and Varieties*. [CONTINUED]

5366-21000-028-00D

SMALL GRAINS GENETICS AND GERMPLASM ENHANCEMENT – John Bonman (P), Donald Obert, Paul Bregitzer, and Eric Jackson; Aberdeen, Idaho

5402-21220-007-00D

NONCHEMICAL PEST CONTROL AND ENHANCED SUGAR BEET GERMPLASM VIA TRADITIONAL AND MOLECULAR TECHNOLOGIES – Leonard Panella (P) and Kimberly Webb; Fort Collins, Colorado

5430-21000-006-00D

GENETIC ENHANCEMENT FOR RESISTANCE TO BIOTIC AND ABIOTIC STRESSES IN HARD WINTER WHEAT – Robert Bowden (P), Ming Shum, Guihua Bai, and John Fellers; Manhattan, Kansas

5440-21000-029-00D

GENETIC IMPROVEMENT AND EVALUATION OF HARD WINTER AND SPRING WHEATS – Robert Graybosch (P); Lincoln, Nebraska

5440-21220-027-00D

ENHANCEMENT OF SORGHUM FOR BIOENERGY, FEED, AND FOOD VALUE – Jeffrey Pedersen (P), Scott Sadler, and Deanna Funnel-Harris; Lincoln, Nebraska

5442-21000-034-00D

SUNFLOWER GERMPLASM DIVERSIFICATION AND CHARACTERIZATION UTILIZING WILD SUNFLOWER SPECIES, CYTOGENETICS, AND APPLIED GENOMICS – Brady Vick (P), Lili Qi, Chao-Chien Jan, and Gerald Seiler; Fargo, North Dakota

5442-21220-024-00D

SUNFLOWER GERMPLASM DEVELOPMENT FOR IMPROVED INSECT AND DISEASE RESISTANCE – Thomas Gulya Jr (P) and Brent Hulke; Fargo, North Dakota

6202-21000-029-00D

PECAN GENETICS AND IMPROVEMENT – Tommy Thompson (P) and Larry Grauke; College Station, Texas

6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

6203-21000-001-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Shannon Pinson, and Ming Hsuan Chen; Beaumont, Texas

6203-21220-001-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Beaumont, Texas

6208-21000-018-00D

CHARACTERIZATION AND ENHANCEMENT OF PLANT RESISTANCE TO WATER-DEFICIT AND THERMAL STRESSES – John Burke (P), James Mahan, Jeffrey Velten, Zhanguo, Paxton Payton, and Junping Chen; Lubbock, Texas

6217-21220-006-00D

ENHANCING DISEASE RESISTANCE AND OIL QUALITY ATTRIBUTES OF PEANUT – Hassan Melouk (P) and Kelly Chamberlin; Stillwater, Oklahoma

Problem Statement 3C: *Germplasm Enhancement/Release of Improved Genetic Resources and Varieties*. [CONTINUED]

6225-21000-009-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Rolfe Bryant, Robert Fjellstrom, Helen Miller, and Wengui Yan; Stuttgart, Arkansas

6402-21000-031-00D

COMBINING CONVENTIONAL AND MOLECULAR TECHNIQUES TO IMPROVE SEED COTTON VALUE – Jodi Scheffler (P), Lawrence Young, and Rickie Turley; Stoneville, Mississippi

6402-21000-032-00D

NATIONAL COTTON VARIETY TEST PROGRAM – Lawrence Young (P); Stoneville, Mississippi

6402-21000-033-00D

GENETIC-PHYSIOLOGICAL TEAM RESEARCH TO IMPROVE PRODUCTION, FIBER QUALITY AND COMPETITIVE ABILITY OF COTTON – Lawrence Young (P), Efrem Bechere, and Linghe Zeng; Stoneville, Mississippi

6402-21000-034-00D

DEVELOPMENT OF SOYBEAN GENEOTYPES AND MANAGEMENT SYSTEMS FOR EARLY SEASON AND STRESS ENVIRONMENTS – Lawrence Young (P), Anne Gillen, and Nacer Bellaloui; Stoneville, Mississippi

6402-21220-010-00D

REDUCING SOYBEAN YIELD LOSSES THROUGH GENETIC IMPROVEMENT – Jeffery Ray (P), Lawrence Young, Shuxian Li, and James Smith; Stoneville, Mississippi

6404-21000-008-00D

SMALL FRUIT AND ORNAMENTAL GENETIC RESEARCH FOR THE MID-SOUTH – Timothy Rinehart (P), Cecil Pounders, James Spiers, Stephen Stringer, and Hamidou Sakhanokho; Poplarville, Mississippi

6406-21000-011-00D

ENHANCING CORN WITH RESISTANCE TO AFLATOXIN CONTAMINATION AND INSECT DAMAGE – William Williams (P), Marilyn Warburton, William Henry, and Gary Windham; Mississippi State, Mississippi

6406-21000-012-00D

GERMPLASM ENHANCEMENT AND GENETIC IMPROVEMENT OF COTTON – Johnie Jenkins (P), Sukumar Saha, Jack McCarthy Jr., and Martin Wubben; Mississippi State, Mississippi

6410-21000-014-00D

GENETIC IMPROVEMENT OF SUGARCANE BY CONVENTIONAL AND MOLECULAR APPROACHES – Thomas Tew (P), Anna Hale, Edward Richard Jr, Yong Bao Pan, and William White; Houma, Louisiana

6602-21000-022-00D

DEVELOPMENT OF IMPROVED PEANUT GERMPLASM AND RESISTANCE TO DISEASE AND NEMATODE PESTS – C. Holbrook Jr (P); Tifton, Georgia

6602-21220-014-00D

GENETIC IMPROVEMENT OF MAIZE AND PEARL MILLET FOR RESISTANCE TO INSECTS AND AFLATOXIN – Jeffrey Wilson (P), C Holbrooke Jr., and Xinzhi Ni; Tifton, Georgia

6604-21000-003-00D

PHYSIOLOGICAL MECHANISMS THAT DETERMINE CROP RESPONSE TO IRRIGATION, DISEASE, AND PRODUCTION PRACTICES – Charles Chen (P), Marshall Lamb, and Phat Dang; Dawson, Georgia

Problem Statement 3C: *Germplasm Enhancement/Release of Improved Genetic Resources and Varieties*. [CONTINUED]

6606-21000-002-00D

STONE FRUIT BREEDING AND DEVELOPMENT – Thomas Beckman (P); Byron, Georgia

6618-21000-013-00D

GENETIC IMPROVEMENT OF CITRUS – Kim Bowman (P), Randall Niedz, Eddie Stover, and Thomas McCollum; Fort Pierce, Florida

6625-21000-003-00D

ENHANCEMENT OF SUGARCANE GERMPLASM FOR DEVELOPMENT OF CULTIVARS AND SUSTAINABLE PRODUCTION – Barry Glaz (P), Jack Comstock, Serge Edme, Neil Glynn, and Duli Zhao; Canal Point, Florida

6631-21000-017-00D

DEVELOPMENT OF AN INTERNATIONAL MARKER-ASSISTED SELECTION PROGRAM FOR CACAO – Raymond Schnell (P) and David Kuhn; Miami, Florida

6635-21000-048-00D

CHARACTERIZATION, CONVERSION, AND IMPROVEMENT OF COMMON BEAN GERMPLASM – Timothy Porch Clay (P); Mayagüez, Puerto Rico

6645-21000-027-00D

GERMPLASM ENHANCEMENT OF MAIZE (GEM) – Matthew Krakowsky (P), Raleigh, North Carolina

6645-21220-012-00D

INCREASING THE COMPETITIVE POSITION OF U.S. SOYBEANS IN GLOBAL MARKETS THROUGH GENETIC DIVERSITY AND PLANT BREEDING – Thomas Carter (P), Joseph Burton, and Robert Upchurch; Raleigh, North Carolina

6645-21220-013-00D

USING GENETIC DIVERSITY OF IMPROVE QUANTITATIVE DISEASE RESISTANCE AND AGRONOMIC TRAITS OF CORN – James Holland (P) and Peter Balint Kurti; Raleigh, North Carolina

6657-21000-006-00D

GENETIC AND CULTURAL PRACTICE IMPROVEMENT FOR SUSTAINABLE COTTON PRODUCTION – Philip Bauer (P) and Benjamin Campbell; Florence, South Carolina

6659-21000-016-00D

GENETIC IMPROVEMENT OF SOUTHERNPEAS AND PEPPERS – Richard Fery (P); Charleston, South Carolina

6659-21000-017-00D

GENETICS AND GENETIC IMPROVEMENT OF DISEASE RESISTANCE AND QUALITY TRAITS IN WATERMELON, BROCCOLI, AND LEAFY GREEN BRASSICAS – Mark Farnham (P) and Amnon Levi; Chaleston, South Carolina

Component 4: Plant Biological and Molecular Processes

Problem Statement 4A: *Functional Utilization of Plant Genomes – Translating Plant Genomics into Crop Improvement*.

1265-21000-141-00D

MOLECULAR APPROACHES TO ENHANCE PLANT NUTRIENT CONTENT, SHELF-LIFE, AND STRESS TOLERANCE – Autar Mattoo (P); Beltsville, Maryland

Problem Statement 4A: *Functional Utilization of Plant Genomes – Translating Plant Genomics into Crop Improvement*. [CONTINUED]

1275-21000-185-00D

ENHANCEMENT OF BLUEBERRY, STRAWBERRY, AND BRAMBLES THROUGH MOLECULAR APPROACHES – Lisa Rowland (P), Kimberly Lewers, Janet Slovin, Mark Ehlenfeldt, and James Polashock; Beltsville, Maryland

1275-21220-229-00D

GENOMICS AND PROTEOMICS APPROACHES TO BROADENING RESISTANCE OF SOYBEAN TO PESTS AND PATHOGENS – Benjamin Matthews (P), Bret Cooper, and Mark Tucker; Beltsville, Maryland

1907-21000-032-00D

DETERMINATION AND CHARACTERIZATION OF UNINTENDED EFFECTS IN GENETICALLY MODIFIED FOOD CROPS – Owen Hoekenga (P), James Giovannoni, and Leon Kochian; Ithaca, New York

1907-21000-033-00D

GENOMICS APPROACHES FOR IMPROVING NUTRITIONAL QUALITY OF FOOD CROP SPECIES – James Giovannoni (P), Leon Kochian, Yong Yang, Li Li, and Theodore Thannhauser; Ithaca, New York

1907-21000-034-00D

GENOMIC APPROACHES TO IMPROVING TRANSPORT AND DETOXIFICATION OF SELECTED MINERAL ELEMENTS IN CROP PLANTS – Leon Kochian (P), Yong Yang, and Theodore Thannhauser; Ithaca, New York

1931-21000-020-00D

USING FUNCTIONAL AND APPLIED GENOMICS TO IMPROVE STRESS AND DISEASE RESISTANCE IN FRUIT TREES – Michael Wisniewski (P), John Norelli, Timothy Artlip, and Carole Bassett; Kearneysville, West Virginia

3607-21000-012-00D

IMPROVEMENT OF PEST RESISTANCE AND QUALITY TRAITS OF SOYBEAN – Rouf Mian (P) and Margaret Redinbaugh; Wooster, Ohio

3622-21000-032-00D

MODIFICATION OF SOYBEAN SEED COMPOSITION FOR FOOD, FEED, AND OTHER INDUSTRIAL USES – Jan Miernyk (P), Perry Gustafson, Jason Gillman, Melvin Oliver, Kristin Bilyeu, and Hari Krishnan; Columbia, Missouri

3622-21000-033-00D

GENETIC ENHANCEMENT OF SOYBEAN SEED VALUE BY BIOTECHNOLOGY – Ivan Baxter (P), Douglas Allen, and Melvin Oliver; Columbia, Missouri

3622-21000-035-00D

FUNCTIONAL GENOMICS FOR IDENTIFICATION OF SOYBEAN QUALITY TRAIT GENES – Yong Qiang An (P) and Melvin Oliver; Columbia, Missouri

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

3625-21220-004-00D

FUNCTIONAL AND STRUCTURAL GENETIC ANALYSIS OF SOYBEAN AND OTHER LEGUMES – Randy Shoemaker (P), Craig Abel, and Michelle Graham; Ames, Iowa

3640-21000-028-00D

FUNCTIONAL GENOMICS FOR IMPROVING NUTRIENTS AND QUALITY IN ALFALFA AND SOYBEAN – Carroll Vance (P); St. Paul, Minnesota

Problem Statement 4A: *Functional Utilization of Plant Genomes – Translating Plant Genomics into Crop Improvement*. [CONTINUED]

3655-21430-009-00D

IMPROVEMENT OF BARLEY SEED QUALITY THROUGH MOLECULAR AND FUNCTIONAL GENOMIC ANALYSIS OF GENE EXPRESSION – Ronald Skadsen (P) and Cynthia Henson; Madison, Wisconsin

5320-21000-013-00D

MOLECULAR RESOURCES FOR THE IMPROVEMENT OF TROPICAL ORNAMENTAL AND FRUIT CROPS – Dennis Gonsalves (P), Roxana Cabos, and Jon Suzuki; Hilo, Hawaii

5325-21430-012-00D

PRODUCTION OF WHEAT GERMPLASM WITH ENHANCED BAKING QUALITY – Ann Blechl (P); Albany, California

5335-21000-026-00D

GENOMIC AND GENETIC ANALYSIS OF THE CEREAL CIRCADIAN SYSTEM AND CROP PRODUCTION – Frank Harmon (P); Albany, California

5335-21000-032-00D

MOLECULAR MECHANISMS OF PHYTOCHROME SIGNALING AND GENE REGULATION – Sarah Hake (P); Albany, California

5335-21000-034-00D

POSITIONAL CLONING IN MAIZE OF GENES THAT REGULATE PLANT ARCHITECTURE – Sarah Hake (P); Albany, California

5335-21000-036-00D

MOLECULAR DEVELOPMENTAL GENETICS OF POLLEN AND POLLEN-PISTIL INTERACTIONS IN CROP PLANTS – Shelia McCormick (P); Albany, California

5335-21000-038-00D

FUNCTIONAL GENOMICS OF PLANT ARCHITECTURE – Jennifer Fletcher (P); Albany, California

5335-21430-008-00D

MOLECULAR GENETICS OF ETHYLENE BIOSYNTHESIS – Sarah Hake (P); Albany, California

5348-21000-023-00D

CHARACTERIZATION OF STRESS RESISTANCE GENES AND MECHANISMS, AND IMPROVEMENT AND GENOTYPING OF WHEAT AND BARLEY GERMPLASM FOR THE WESTERN U.S. – Camille Steber (P), Deven See, Daniel Skinner, and Kimberly Garland Campbell; Pullman, Washington

5366-21000-025-00D

PLANT AND SEED CHEMISTRY GENETICS – Victor Raboy (P) and John Bonman; Aberdeen, Idaho

5442-21000-032-00D

GENOMIC RELATIONSHIPS AND CYTOGENETIC MANIPULATION OF DURUM WHEAT BY CLASSICAL AND MOLECULAR TECHNIQUES – Prem Jauhar (P); Fargo, North Dakota

6435-21000-016-00D

MOLECULAR ANALYSIS OF DEVELOPMENT TO IMPROVE COTTON FIBER – David Fang (P) and Hee Kim; New Orleans, Louisiana

6602-21000-021-00D

GENETIC AND GENOMIC APPROACHES TO IMPROVE PEANUT AND CORN RESISTANCE TO DISEASE AND AFLATOXIN CONTAMINATION – Baozhu Guo (P) and Brian Scully; Tifton, Georgia

Problem Statement 4A: *Functional Utilization of Plant Genomes – Translating Plant Genomics into Crop Improvement.* [CONTINUED]

6615-21000-010-00D

FUNCTIONAL GENOMICS OF AGRONOMIC TRAITS IN DEVELOPING SEED AND POLLEN IN MAIZE AND SORGHUM – Prem Chourey (P), Alisa Huffaker, and Peter Teal; Gainesville, Florida

Problem Statement 4B: *Biological Processes that Improve Crop Productivity and Quality.*

1230-21000-048-00D

GENETIC ENGINEERING OF FLORAL BULB CROPS FOR VIRUS AND NEMATODE RESISTANCE – Kathryn Kamo (P); U.S. National Arboretum, Washington, D.C.

1265-21000-141-00D

MOLECULAR APPROACHES TO ENHANCE PLANT NUTRIENT CONTENT, SHELF-LIFE, AND STRESS TOLERANCE – Autar Mattoo (P); Beltsville, Maryland

1275-21000-185-00D

ENHANCEMENT OF BLUEBERRY, STRAWBERRY, AND BRAMBLES THROUGH MOLECULAR APPROACHES – Lisa Rowland (P), Kimberly Lewers, Janet Slovin, Mark Ehlenfeldt, and James Polashock; Beltsville, Maryland

1275-21220-230-00D

MOLECULAR GENETIC APPROACHES TO PEST AND PATHOGEN RESISTANCE IN SUGAR BEET, BETA VULGARIS L. – Anna Smigocki (P) and L. David Kuykendall; Beltsville, Maryland

1907-21000-033-00D

GENOMICS APPROACHES FOR IMPROVING NUTRITIONAL QUALITY OF FOOD CROP SPECIES – James Giovannoni (P), Leon Kochian, Yong Yang, Li Li, and Theodore Thannhauser; Ithaca, New York

1907-21000-034-00D

GENOMIC APPROACHES TO IMPROVING TRANSPORT AND DETOXIFICATION OF SELECTED MINERAL ELEMENTS IN CROP PLANTS – Leon Kochian (P), Yong Yang, and Theodore Thannhauser; Ithaca, New York

1931-21000-020-00D

USING FUNCTIONAL AND APPLIED GENOMICS TO IMPROVE STRESS AND DISEASE RESISTANCE IN FRUIT TREES – Michael Wisniewski (P), John Norelli, Timothy Artlip, and Carole Bassett; Kearneysville, West Virginia

3602-21000-006-00D

IDENTIFICATION AND CHARACTERIZATION OF GENES IMPORTANT DURING SEED DEVELOPMENT IN LEGUMES – Karen Hudson (P); West Lafayette, Indiana

3611-21000-024-00D

IDENTIFYING AND MANIPULATING DETERMINANTS OF PHOTOSYNTHATE PRODUCTION AND PARTITIONING – Donald Ort (P), Elizabeth Ainsworth, Carl Bernacchi, and Steven Huber; Urbana, Illinois

3622-21000-033-00D

GENETIC ENHANCEMENT OF SOYBEAN SEED VALUE BY BIOTECHNOLOGY – Ivan Baxter (P), Douglas Allen, and Melvin Oliver; Columbia, Missouri

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

3640-21000-028-00D

FUNCTIONAL GENOMICS FOR IMPROVING NUTRIENTS AND QUALITY IN ALFALFA AND SOYBEAN – Carroll Vance (P); St. Paul, Minnesota

Problem Statement 4B: *Biological Processes that Improve Crop Productivity and Quality*. [CONTINUED]

3655-21000-052-00D

PHYSIOLOGICAL, BIOCHEMICAL AND GENETIC REGULATION OF CARBOHYDRATE METABOLISM IN CEREAL TISSUES – Cynthia Henson (P); Madison, Wisconsin

3655-21000-053-00D

METABOLISM AND ANALYSIS OF CEREAL PHYTOCHEMICALS – Mitchell Wise (P) and Cynthia Henson; Madison, Wisconsin

3655-21430-009-00D

IMPROVEMENT OF BARLEY SEED QUALITY THROUGH MOLECULAR AND FUNCTIONAL GENOMIC ANALYSIS OF GENE EXPRESSION – Ronald Skadsen (P) and Cynthia Henson; Madison, Wisconsin

5320-21000-013-00D

MOLECULAR RESOURCES FOR THE IMPROVEMENT OF TROPICAL ORNAMENTAL AND FRUIT CROPS – Dennis Gonsalves (P), Roxana Cabos, and Jon Suzuki; Hilo, Hawaii

5335-21000-034-00D

POSITIONAL CLONING IN MAIZE OF GENES THAT REGULATE PLANT ARCHITECTURE – Sarah Hake (P); Albany, California

5325-21430-012-00D

PRODUCTION OF WHEAT GERMPLASM WITH ENHANCED BAKING QUALITY – Ann Blechl (P); Albany, California

5335-21430-008-00D

MOLECULAR GENETICS OF ETHYLENE BIOSYNTHESIS – Sarah Hake (P); Albany, California

5348-21000-023-00D

CHARACTERIZATION OF STRESS RESISTANCE GENES AND MECHANISMS, AND IMPROVEMENT AND GENOTYPING OF WHEAT AND BARLEY GERMPLASM FOR THE WESTERN U.S. – Camille Steber (P), Deven See, Daniel Skinner, and Kimberly Garland Campbell; Pullman, Washington

5366-21000-025-00D

PLANT AND SEED CHEMISTRY GENETICS – Victor Raboy (P) and John Bonman; Aberdeen, Idaho

5442-21430-007-00D

SUCROSE ACCUMULATION AND RETENTION IN SUGARBEETS – Karen Fugate (P) and Larry Campbell; Fargo, North Dakota

6408-21410-008-00D

DISCOVERY AND DEVELOPMENT OF NATURAL PRODUCT FOR PHARMACEUTICAL AND AGRICHEMICAL APPLICATIONS – Stephen Duke (P); University, Mississippi

6435-21000-016-00D

MOLECULAR ANALYSIS OF DEVELOPMENT TO IMPROVE COTTON FIBER – David Fang (P) and Hee Kim; New Orleans, Louisiana

6615-21000-010-00D

FUNCTIONAL GENOMICS OF AGRONOMIC TRAITS IN DEVELOPING SEED AND POLLEN IN MAIZE AND SORGHUM – Prem Chourey (P), Alisa Huffaker, and Peter Teal; Gainesville, Florida

6645-21000-028-00D

PHYSIOLOGICAL/BIOCHEMICAL MECHANISMS ASSOCIATED WITH GENETIC ALTERATION OF SOYBEAN QUALITY AND PRODUCTIVITY – Robert Upchurch (P), Earl Taliercio, Thomas Carter Jr., and Joseph Burton; Raleigh, North Carolina

Problem Statement 4C: *Plant Biotechnology Risk Assessment.*

1275-21000-224-00D

EVALUATION OF THE QUALITY AND SAFETY OF TRANSGENIC SOYBEANS – Savithiry Natarajan (P); Beltsville, Maryland

1907-21000-032-00D

DETERMINATION AND CHARACTERIZATION OF UNINTENDED EFFECTS IN GENETICALLY MODIFIED FOOD CROPS – Owen Hoekenga (P), James Giovannoni, and Leon Kochian; Ithaca, New York

1931-21000-017-00D

GENETIC IMPROVEMENT OF FRUIT CROPS THROUGH FUNCTIONAL GENOMICS AND BREEDING – Ralph Scorza (P), Richard Bell, Ann Callahan, Zongrang Liu, Christopher Dardick, and Chinnathambi Srinivasan; Kearneysville, West Virginia

3655-21000-054-00D

GENETIC EXCHANGE AND GENE FLOW RISKS FROM PLANTS IN AGRICULTURE – Johanne Brunet (P) and Philipp Simon; Madison, Wisconsin

5320-21000-013-00D

MOLECULAR RESOURCES FOR THE IMPROVEMENT OF TROPICAL ORNAMENTAL AND FRUIT CROPS – Dennis Gonsalves (P), Roxana Cabos, and Jon Suzuki; Hilo, Hawaii

5325-21000-018-00D

MOLECULAR TOOLS TO MINIMIZE RISK IN GENETICALLY ENGINEERED CROPS – Ann Blechl (P), Roger Thilmony, and James Thomson; Albany, California

5335-21000-040-00D

TRANSGENE MANAGEMENT THROUGH SITE-SPECIFIC RECOMBINATION – Sarah Hake (P); Albany, California

APPENDIX 2

National Program 301 Plant Genetic Resources, Genomics, and Genetics Improvement

ACCOMPLISHMENT REPORT 2006 – 2011

Publications by Research Project*

1230-21000-048-00D

GENETIC ENGINEERING OF FLORAL BULB CROPS FOR VIRUS AND NEMATODE RESISTANCE – Kathryn Kamo (P); U.S. National Arboretum, Washington, D.C.

- Kamo, K., Jordan, R., Guaragna, M.A., Hsu, H.T., Ueng, P. 2010. Resistance to cucumber mosaic virus in Gladiolus plants transformed with either a defective replicase or coat protein subgroup II gene from cucumber mosaic virus. *Plant Cell Reports*. 29:695-704.
- Kamo, K., Joung, Y.H., Green, K. 2009. GUS expression in Gladiolus plants controlled by two Gladiolus ubiquitin promoters. *Floriculture and Ornamental Biotechnology*. 3:10-14.

1230-21000-050-00D

TAXONOMY AND GENETIC DIVERSITY ASSESSMENT OF LANDSCAPE TREES AND SHRUBS – Alan Whittemore (P) and Joseph Kirkbride; U.S. National Arboretum, Washington, D.C.

- A. T. Whittemore and R. Olsen. 2011. *Ulmus americana* is a polyploid complex. *American Journal of Botany* 98: 754-760.
- R. Arias, R. S., N. Tehen, T. A. Rinehart, R. T. Olsen, J. H. Kirkbride, Jr., and B. E. Scheffler. 2011. Development of simple sequence repeat markers for *Chionanthus retusus* (Oleaceae) and effective discrimination of closely related taxa. *HortScience* 46(1):23-29.
- Whittemore, A.T. 2009. What is *Ranunculus gelidus*? *Journal of Botanical Research Institute of Texas*. 3:245-250.
- Delprete, P., Kirkbride, J.H. 2008. Clarification of *Borreria gymnocephala*, *Diodia gymnocephala*, *Diodia schumannii*, *Borreria flavoriens*, and *Spermacoce schumannii* (Rubiaceae). *Journal of Botanical Research Institute of Texas*. 3:245-250.
- Whittemore, A.T. 2008. Exotic species of *Celtis* (Cannabaceae) in the flora of North America. *Journal of the Botanical Research Institute of Texas*. 2:627-632.
- Whittemore, A.T. 2008. Proposal to reject the name *Ulmus campestris* (Ulmaceae). *Taxon*. 57:656-657.
- Whittemore, A.T. Ranunculaceae: *Aquilegia*, *Ceratocephala*, *Myosurus*, and *Ranunculus*. *Flora of the San Juan Basin*.
- Ramirez, P., Whittemore, A.T., Churchill, S.P. 2007. The bryophytes of the Departments of Norina and Putumayo, Columbia: II. Hepatics and hornworts. *The Bryologist*. 111:363-549.
- Kirkbride, J.H.Jr., Wiersema, J.H. 2007. Proposal to reject the name *Arundo vallisoria* (Poaceae). *Taxon*. 56(1):258.
- Wiersema, J.H., Dahlberg, J. 2007. The nomenclature of *Sorghum bicolor* (L.) Moench (Gramineae). *Taxon*. 56(3):941-946.
- Kirkbride, J.H., Wiersema, J.H. 2007. Proposal to conserve the name *Zinnia elegans* against *Z. violacea* (Compositae). *Taxon*. 56(3):958-959.
- Kirkbride, J.H. Jr., Wiersema, J.H., Turland, N.J. 2006. Proposal to conserve the name *Ziziphus Jujuba* against *z. zizyphus* (Rhamnaceae). *Taxon*. 55(4):1049-1050.

* For the sake of consistency, projects are listed and organized in Appendixes 1 and 2 according to the ARS project number used to track projects in the Agency's internal database. A (P) after a scientist's name indicates the project's principal investigator.

1230-21000-051-00D

GENETICS, GENETIC RESOURCE EVALUATION, AND GENETIC IMPROVEMENT OF LANDSCAPE TREES AND SHRUBS – Margaret Pooler (P); U.S. National Arboretum, Washington, D.C.

- Pooler, M.R. 2011. 'Anacostia' Camellia. *HortScience*. 46:139-140.
- Pooler, M.R. 2011. 'Cree' and 'Nantucket' Viburnum. *HortScience*. 45:1384-1385.
- Ma, H., Olsen, R., Pooler, M., Kramer, Matthew. 2009. Evaluation of Flowering Cherry Species, Hybrids, and Cultivars Using Simple Sequence Repeat Markers. *Journal of the American Society for Horticultural Science*. 134:435-444.
- Ma, H., Pooler, M.R., Griesbach, R.J. 2009. Anthocyanin regulatory/structural gene expression in Phalaenopsis. *Journal of the American Society for Horticultural Science*. 134:88-96.
- Pooler, M.R. 2008. 'Betsy Ross', 'Old Glory', and 'Declaration' lilacs. *HortScience*. 43(2):544-545.
- Pooler, M.R., Nicholson, R., Vandergrift, A. 2008. Clonal fidelity in large colonies of box huckleberry (*Gaylussacia brachycea* Gray) assessed by DNA fingerprinting. *Northeastern Naturalist*. 15(1):67-74.
- Pooler, M.R. 2007. 'Dream Cater' and 'First Lady' Flowering Cherry. *HortScience*. 42(1):174-175.

1230-21000-052-00D

GENETICS, GENETIC IMPROVEMENT, AND IMPROVED PRODUCTION EFFICIENCY OF NURSERY CROPS – Sandra Reed (P) and Donna Fare; McMinnville, Tennessee

- Oliver, J.B., Fare, D.C., Youssef, N., Scholl, S., Reding, M.E., Ranger, C.M., Moyseenko, J.J., Halcomb, M.A. 2010. Evaluation of a single application of Neonicotinoid and multi-application contact insecticides for flatheaded borer management in field grown *Acer rubrum* L. cultivars. *Journal of Environmental Horticulture*. 28(3):135-149.
- Reed, S.M., Rinehart, T.A. 2009. SSR marker analysis of genetic relationships within *Hydrangea paniculata*. *HortScience*. 44(1):21-31.
- Wadl, P.A., Skinner, J.A., Dunlap, J.R., Reed, S.M., Rinehart, T.A., Trigiano, R.N. 2009. Honeybee-mediated controlled pollinations in *Cornus florida* and *C. kousa* intra- and inter-specific crosses. *HortScience*. 44:1527-1533.
- Oliver, J.B., Fare, D.C., Youssef, N., Halcomb, M.A., Reding, M.E., Ranger, C.M. 2009. Evaluation of systemic insecticides for potato leafhopper control in field-grown red maple. *Journal of Environmental Horticulture*. 27(1):17-23.
- Reed, S.M., Bachman, G.R. and Davis, W.E. 2008. 'Duet' beautyberry. *HortScience* 43:933-934.
- Reed, S.M., Jones, K., Rinehart, T.A. 2008. Production and characterization of intergeneric hybrids between *Dichroa febrifuga* and *Hydrangea macrophylla*. *Journal of the American Society for Horticultural Science*. 133:84-91.
- Li, Y., M.T. Windham, R.N. Trigiano, S.M. Reed, J.M. Spiers, and T.A. Rinehart. 2008. Effects of shading on *Cercospora* leaf spot in bigleaf hydrangea. *Proc. Southern Nursery Assoc. Res. Conf.* 53:379-380.
- Reed, S.M. and Rinehart, T.A. 2007. SSR marker analysis of genetic relationships within *Hydrangea macrophylla*. *J. Amer. Soc. Hort. Sci.* 132:341-351.
- Jones, K.D., Reed, S.M. and Rinehart, T.A. 2007. Analysis of ploidy level and its effects on guard cell length, pollen diameter and fertility in *Hydrangea macrophylla*. *HortScience*. 42:483-488.
- Reed, S.M. and Rinehart, T.A. 2006. *Hydrangea macrophylla* and *serrata* - Should we lump 'em or split 'em? *Proceedings of the Southern Nursery Association Research Conference*. 51:573-576.
- Jones, K.D., Reed, S.M. and Rinehart, T.A. 2006. Wide crosses in the *Hydrangeaceae*: *Dichroa febrifuga* x *Hydrangea macrophylla*. *Proceedings of the Southern Nursery Association Research Conference*. 51:577-579.

1230-21000-053-00D

GENETIC IMPROVEMENT OF LANDSCAPE TREES FOR DISEASE AND PEST TOLERANCE, NON-INVASIVENESS, AND ORNAMENTAL TRAITS – Richard Olsen (P); U.S. National Arboretum, Washington, D.C.

- Trueblood, C.E., Ranney, T.G., Lynch, N.P., Neal, J.C., Olsen, R.T. 2010. Evaluating fertility of triploid clones of *Hypericum androsaemum* L. for use as non-invasive landscape plants. *HortScience*. 45:1026-1028.
- Montgomery, M.E., Bentz, S.E., Olsen, R.T. 2009. Evaluation of hemlock (*Tsuga*) species and hybrids for resistance to *Adelges tsugae* (Hemiptera: Adelgidae) using artificial inoculation. *Journal of Economic Entomology*. 102:1247-1254.
- Olsen, R.T. and Whittemore, A.T. 2009. Validation of the hybrid flowering cherry *Prunus xincam* (Rosaceae). *Novon*. 19:490-493.

1230-21000-137-00D

GENETIC RESOURCES, EVALUATION, AND INFORMATION MANAGEMENT OF WOODY LANDSCAPE PLANT

GERMPLASM – Richard Olsen (P), Kevin Conrad, and Mark Roh; U.S. National Arboretum, Washington, D.C.

- Joung, Y., Suh, J., Lee, N., Eum, S., Roh, M.S. 2011. Identification of Hydrangeaceae accessions of wild origin from Jeju, Korea using molecular markers. *FLORA*. 8(3):235-241.
- Yuan, T., Wang, L.Y., Roh, M.S. 2010. Confirmation of Clematis hybrids using molecular markers. *Scientia Horticulturae*. 125:136-145.
- Roh, M.S., and Lawson, R. 2009. Source of boron in Curcuma for burn symptoms at leaf margins. *J. Plant Nutrition*. 32:798-808.
- Roh, M.S., Lee, A.K., Suh, J.K., Bordelon, C.M. 2009. Interspecific variations in seed germination of Corylopsis. *Scient. Hort.* 118:347-350.
- Roh, M.S., Lee, A.K., Suh, J.K., Joung, Y.H., Choi, I.Y., Lee, N.S., Park, J.O., Koshioka, M., Yeau, S.H., Slovin, J.E. 2009. Evaluation, characterization, and identification of woody landscape plants. *Acta Hort.* 769:327-338.
- Roh, M., Joung, Young Hee, Suh, Jung Keun, and Lee, Ae-Kyung. 2008. Production of quality woody and floral crops using innovative production techniques. *Acta Hort.* 766:29-44.
- Roh, M.S., Cheong, E.J., Choi, I-Y and Young, Y.H. 2007. Characterization of wild Prunus yedoensis analyzed by inter-simple sequence repeat and chloroplast DNA. *Scientia Hort.* doi:10.1016/J.scientia.2007.06.005.
- Roh, M.S., Lee, A.K. and Suh, J.K. 2007. Induction of bulb maturity of Ornithogalum thyrsoides. *Scientia Hort.* doi:10.1016/j.scientia.2007.06.004.
- Roh, M.S., Lee, A.K., Choe, I-Y., Kim, J-Y., Joung, Y.H., Lee, S.H. and Such, J.K. 2007. Characterization of the Corylopsis coreana using Molecular Markers. *Hort. Environ. Biotechnol.* 48:176-187.

1265-21000-141-00D

MOLECULAR APPROACHES TO ENHANCE PLANT NUTRIENT CONTENT, SHELF-LIFE, AND STRESS TOLERANCE –

Autar Mattoo (P); Beltsville, Maryland

- Sobolev, A.P., Capitani, D., Giannino, D., Nicolodi, C., Testone, G., Santoro, F., Frugis, G., Iannelli, M.A., Mattoo, A.K., Brosio, E., Gianferri, R., Amico, I.D., Mannina, L. 2010. NMR-Metabolic Methodology in the Study of GM Foods. *Nutrients*. 2:1-15.
- Handa, A.K., Mattoo, A.K. 2010. Differential and functional interactions emphasize the multiple roles of polyamines in plants. *Plant Physiology and Biochemistry*. 48:540-546.
- Nambessan, S., Datsenka, T., Ferruzzi, M.G., Malladi, A., Mattoo, A.K., and Handa, A.K. 2010. Overexpression of yeast spermidine synthase impacts ripening, senescence and decay symptoms in tomato. *Plant J.* 63:836-847.
- Srivastava, A., Gupta, A.K., Datsenka, T., Mattoo, A.K., and Handa, A.K. 2010. Maturity and ripening-stage specific modulation of tomato (Solanum lycopersicum) fruit transcriptome. *GM Crops* 1:1-13.
- Mattoo, A.K., Minocha, S., Minocha, R., Handa, A.K. 2009. Polyamines and cellular metabolism in plants: Transgenic approaches reveal different responses to diamine putrescine versus higher polyamines spermidine and spermine. *Amino Acids*. Available: <http://www.springerlink.com/content/05h3202610227w87/fulltext.html>

1275-21000-185-00D

ENHANCEMENT OF BLUEBERRY, STRAWBERRY, AND BRAMBLES THROUGH MOLECULAR APPROACHES –

Lisa Rowland (P), Kimberly Lewers, Janet Slovin, Mark Ehlenfeldt, and James Polashock; Beltsville, Maryland

- Bell, D.J., Stommel, J.R., Rowland, L.J., Drummond, F. 2010. Yield variation among clones of lowbush blueberry as a function of kinship and self-compatibility. *Journal of the American Society for Horticultural Science*. 135:259-270.
- Liu, C., Callow, P., Rowland, L.J., Hancock, J.F., Song, G. 2010. Adventitious shoot regeneration from leaf explants of southern highbush blueberry cultivars. *Plant Cell Tissue And Organ Culture*. 103:137-144.
- Polashock, J.J., Arora, R., Peng, Y., Dhananjay, N., Rowland, L.J. 2010. Functional identification of a blueberry CBF/DREB-like element associated with cold acclimation and freezing tolerance. *Journal of the American Society for Horticultural Science*. 135(1)40-48.
- Rowland, L.J., Alkharouf, N., Bassil, N.V., Beers, L., Bell, D.J., Buck, E., Drummond, F.A., Finn, C.E., Graham, J., Hancock, J., McCallum, S., Olmstead, J. 2010. Generating genomic tools for blueberry improvement -- an update of our progress. *HortScience*. 2010:45(8)S197.

- Spigler, R.B., Ashman, T., Lewers, K.S. 2010. Genetic architecture of sexual dimorphism in a subdioecious plant with a proto-sex chromosome. *Evolution*. DOI: 10.1111/j.1558-5646.2010.01189.x.
- Bell, D.J., Rowland, L.J., Smagula, J.M., Drummond, F.A. 2009. Recent Advances in the Biology and Genetics of Lowbush Blueberry. *Experiment Station Bulletins. MAFES Technical Bulletin* 203:1-28
- Bell, D.J., Rowland, L.J., Zhang, D., Drummond, F.A. 2009. The spatial genetic structure of lowbush blueberry, *Vaccinium angustifolium* Ait., in four fields in Maine. *Botany*. 87:932-946
- Lewers, K.S., Saski, C.A., Cuthbertson, B.J., Henry, D.C., Staton, M.E., Main, D.S., Dhanaraj, A.L., Rowland, L.J., Tomkins, J.P. 2008. A blackberry (*Rubus* L.) expressed sequence tag library for the development of simple sequence repeat markers. *Plant Science*. 8:69-76.
- Bell, D.J., Rowland, L.J., Polashock, J.J., Drummond, F.A. 2008. Suitability of EST-PCR markers developed in highbush blueberry (*Vaccinium corymbosum* L.) for genetic fingerprinting and relationship studies in lowbush blueberry (*V. angustifolium* Ait.). *Journal of the American Society for Horticultural Science*. 133:701-707.
- Alkharouf, N.W., Dhanaraj, A.L., Naik, D., Overall, C., Matthews, B.F., Rowland, L.J. 2007. Bbgd: an online database for blueberry genomic data.. *Biomed Central (BMC) Plant Biology*. 7:05.
- Naik, D., Dhanaraj, A.L., Arora, R., Rowland, L.J. 2007. Identification of cold-responsive genes in blueberry (*Vaccinium corymbosum* L.) using a hybridization approach. *Plant Science*. 173: 213-222
- Polashock, J.J., Saftner, R.A., Ehlenfeldt, M.K. 2007. Organoleptic Analysis of Blueberry Cultivars. *New Jersey Annual Vegetable Meeting Proceedings*, pp. #152-153.
- Dhanaraj, A.L., Alfharouf, N.W., Beard, H.S., Chouikha, I.B., Matthews, B.F., Wei, H., Arora, R., Rowland, L.J. 2006. Gene expression profiles during cold acclimation in blueberry under field and cold room conditions using cdna microarrays. *Planta*. DOI10.1007/s00425-006-0382-1. 225:735-751.

1275-21000-190-00D

GENETIC ENHANCEMENT OF POTATOES FOR NUTRITIONAL AND PROCESSING QUALITY AND FOR RESISTANCE TO DISEASES – Kathleen Haynes (P); Beltsville, Maryland

- Haynes, K.G., Clevidence, B.A., Rao, D.D., Vinyard, B.T., White, J.M. 2010. Genotype x environment interactions for potato tuber carotenoid content. *Journal of the American Society for Horticultural Science*. 135:250-258.
- Clough, M.E., Yencho, C.G., Christ, B., Delong, W., Halseth, D., Haynes, K.G., Henninger, M., Hutchinson, C., Kleinhenz, M., Porter, G., Veilleux, R. 2010. An Interactive Online Database for Potato Varieties Evaluated in the Eastern U.S.. *HortTechnology*. 20:245-249.
- Haynes, K.G., Wanner, L.A., Thill, C.A., Bradeen, J.M., Miller, J., Novy, R.G., Whitworth, J.L., Corsini, D.L., Vinyard, B.T. 2010. Common Scab Trials of Potato Varieties and Advanced Selections at Three U.S. Locations. *American Journal of Potato Research*. 86:261-276.
- Haynes, K.G., Christ, B., Burkhart, C., Vinyard, B.T. 2009. Heritability of Resistance to Common Scab in Diploid Potatoes. *American Journal of Potato Research*. 86:165-170.
- Santa Cruz, J., Haynes, K.G., Chris, B.J. 2009. Effects of one cycle of recurrent selection for early blight resistance in a diploid hybrid solanum phureja-S. stenotomum population. *American Journal of Potato Research*. 86:490-498.
- Wickramasinghe, W., Qu, S., Costanzo, S., Haynes, K.G., Christ, B. 2009. Development of PCR-Based Markers Linked to Quantitative Resistance to Late Blight in a Diploid Hybrid Potato Population of Solanum phureja x S. stenotomum. *American Journal of Potato Research*. 86:188-195.
- Mccord, P., Yencho, G., Haynes, K.G., Sterrett, S. 2008. Internal Heat Necrosis of Potato A Review. *American Journal of Potato Research*. 85:69-76.
- Haynes, K.G. 2008. Heritability of Chip Color and Specific Gravity in a Long-Day Adapted Solanum phureja-S. stenotomum Population. *American Journal of Potato Research*. 85:361-366.
- Haynes, K.G., Christ, B.J., Vinyard, B.T. 2008. Determining the Importance of Combining Ability for Late Blight Resistance in Early Generations of Potato Breeding When Susceptible Clones are Discarded. *American Journal of Potato Research*. 85:445-454.
- Haynes, K.G., Goth, R.W., Lambert, D.H., Christ, B.J. 2007. Evaluation of a short-day adapted tetraploid potato population with horizontal resistance to phythphora infestans under long-day conditions in northern Maine. *American Journal of Potato Research*. 84:459-466.
- Burkhart, C.R., Christ, B.J., Haynes, K.G. 2007. Heritability of resistance to fusarium dry rot in a diploid hybrid potato population. *American Journal of Potato Research*. 84:199-204.

1275-21000-195-00D

GENETIC ENHANCEMENT OF QUALITY CONSTITUENTS IN SOLANACEOUS VEGETABLES – John Stommel (P); Beltsville, Maryland

- Lightbourn, G., Winkel, B., Griesbach, R.J., Stommel, J.R. 2009. Transcription Factor Families Regulate the Anthocyanin Biosynthetic Pathway in Capsicum. *Journal of the American Society for Horticultural Science*. 134:244-251.
- Stommel, J.R., Griesbach, R.J. 2008. Capsicum Annuum L. Lil' Pumpkin and Pepper Jack. *HortScience*. 43:935-938.
- Stommel, J.R., Griesbach, R.J. 2008. Capsicum Annuum L. Midnight Creeper and Solar Eclipse. *HortScience*. 43:939-942.
- Stommel, J.R., Griesbach, R.J. 2008. Inheritance of fruit, foliar and plant habit attributes in Capsicum L. *Journal of the American Society for Horticultural Science*. 133:396-407.
- Lightbourn, G., Griesbach, R., Stommel, J.R. 2007. Genetic basis for genotype-environment interactions influencing flavonoid gene expression in Capsicum. *Molecular Genetics and Genomics*. 132:824-829.

1275-21000-224-00D

EVALUATION OF THE QUALITY AND SAFETY OF TRANSGENIC SOYBEANS – Savithiry Natarajan (P); Beltsville, Maryland

- Natarajan, S.S., Krishnan, H.B., Lakshman, S., Garrett, W.M. 2009. An efficient extraction method to enhance analysis of low abundant proteins from soybean seed. *Analytical Biochemistry*. 394(2):259-268.

1275-21000-226-00D

NATIONAL RHIZOBIUM GERMLASM RESOURCE COLLECTION, GENETIC RESOURCE MANAGEMENT, SYSTEMATICS, AND BIOINFORMATICS – Peter Van Berkum (P); Beltsville, Maryland

- Bromfield, E.S., Tambong, J.T., Cloutier, S., Prevost, D., Laguerre, G., Van Berkum, P.B., Tran Thi, T.V., Assabgui, R., Barran, L.R. 2010. Ensifer, Phyllobacterium and Rhizobium species occupy nodules of Medicago sativa (alfalfa) and Melilotus alba (sweet clover) grown at a Canadian site without a history of cultivation. *Microbiology*. 156:505-520.
- Van Berkum, P.B., Elia, P.E., Eardly, B.D. 2010. Application of Multilocus Sequence Typing To Study the Genetic Structure of Megaplasmids in Medicago-Nodulating Rhizobia. *Applied and Environmental Microbiology*. 76:3967-3977.
- Zurdo-Pineiro, J.L., Garcia-Fraile, P., Rivas, R., Leon-Barrios, M., Willems, A., Mateos, P.F., Martinez-Molina, E., Velazquez, E., Van Berkum, P.B. 2009. Rhizobia from Lanzarote, the Canary Islands, that nodulate Phaseolus vulgaris have characteristics in common with LMW RNA group II Sinorhizobium meliloti of Medicago, Melilotus and Trigonella from soils of mainland Spain. *Applied and Environmental Microbiology*. 75:2354-2359.
- Van Berkum, P.B., Fuhrmann, J.J. 2009. Evidence from ITS sequence analysis of 31 and 110 serogroup soybean strains that extant members of the genus Bradyrhizobium are likely the products of reticulate evolutionary events. *Applied and Environmental Microbiology*. 75:78-82.
- Del Villar, M., Rivas, R., Peix, A., Mateos, P.F., Martinez-Molina, E., Van Berkum, P.B., Willems, A., Velazquez, E. 2008. Two different stable low molecular weight RNA (LMW RNA) profiles within Sinorhizobium meliloti and within Sinorhizobium medicae. *FEMS Microbiology Letters*. 282:273-281.
- Zribi, K., Badri, Y., Van Berkum, P.B., Aouani, M.E. 2007. Medicago ciliaris growing in Tunisian soils is preferentially nodulated by Sinorhizobium medicae. *Australian Journal of Soil Science*. 45:473-477.
- Martir, M., Tlusty, B., Van Berkum, P.B., Graham, P.H. 2007. The genetic diversity of rhizobia associated with Dalea purpurea Vent. in fragmented grasslands of West-Central Minnesota. *Canadian Journal of Microbiology*. 53:351-363.
- Van Berkum, P.B., Abou-Shanab, R.A., Angle, J.S. 2007. Heavy metal resistance and genotypic analysis of metal resistance genes in gram-positive and gram-negative bacteria present in serpentine soil and in the rhizosphere of alyssum murale. *Chemosphere*. v. 68, p. 360-367.
- Van Berkum, P.B., Badri, Y., Elia, P.E., Aouani, M.E., Eardly, B.D. 2007. Chromosomal and symbiotic relationships of rhizobia nodulating Medicago truncatula and M. laciniata. *Applied and Environmental Microbiology*. 73(23):7597-7604.
- Van Berkum, P.B., Badri, Y., Zribi, K., Badri, M., Aouani, M.E., Huguet, T. 2007. Comparison of rhizobia that nodulate medicago laciniata and m. truncatula present in a single tunisian arid soil. *Canadian Journal of Microbiology*. 53:277-283.

- Van Berkum, P.B., Beyhaut, E., Tlustý, B., Graham, P.H. 2006. *Rhizobium giardinii* is the microsymbiont of illinois bundleflower (*desmanthus illinoensis* (michx) macmillan in midwestern prairies. *Canadian Journal of Microbiology*. 52:903-907.
- Van Berkum, P.B., Elia, P.E., Eardly, B.D. 2006. Multi-locus sequence typing (mlst), a basis for the population analysis of medicago-nodulating rhizobia. *Journal of Bacteriology*. 188:5570-5577.

1275-21000-228-00D

EXCHANGE OF CROP GENETIC RESOURCES AND ASSOCIATED DOCUMENTATION FOR THE U.S. NATIONAL PLANT GERMPLASM SYSTEM – Edward Garvey (p), John Wiersema, and Karen Williams; Beltsville, Maryland

- Dorr, L., Wiersema, J.H. 2010. Names of American vascular plants published in Loeffling's *Iter Hispanicum* (1758) and its German translation(1766). *Taxon*. 59(4):1245-1262
- Dorr, L.J., Wiersema, J.H. 2010. (1947-1958) Proposals to reject twelve names emanating from Loeffling's *Iter Hispanicum* (1758), *Ayenia sidiformis* (Malvaceae), *Cofer* (Symplocaceae), *Cruzeta* and *C. hispanica* (Amaranthaceae), *Edechia inermis* and *E. spinosa* (Rubiaceae), *Justicia putata* (Acanthaceae), *Menais* and *M. topiaria* (? Boraginaceae), *Muco* (Capparaceae), *Samyda parviflora* (Salicaceae), and *Spermacoce suffruticosa* (Rubiaceae). *Taxon*. 59(4):1280-1282.
- Reveal, J.L., Gandhi, K.N., Wiersema, J.H. 2009. Proposal to conserve the name *Thaspium* (Umbelliferae/Apiaceae). *Taxon* 58(3): 14.
- Lohne, C., Borsch, T., Jacobs, S.W.L., Hellquist, C.B., Wiersema, J.H. 2008. Nuclear and plastid DNA sequences reveal complex reticulate patterns in Australian water-lilies (*Nymphaea* subgenus *Anecphyta*, *Nymphaeaceae*). *Australian Systematic Botany*. 21(4):229-250.
- Lohne, C., Yoo, M., Borsch, T., Wiersema, J.H., Wilde, V., Bell, C.D., Barthlott, W., Soltis, D.E., Soltis, P.S. 2008. Biogeography of *Nymphaeales*: Extant patterns and historical events. *Taxon* 57(4):1123-1146.
- Ward, D.B., Wiersema, J.H. 2008. (1807) Proposal to reject the name *Juglans alba* (*Carya alba*)(*Juglandaceae*). *Taxon*. 57(1):308-310.
- Borsch, T., Hilu, K.W., Wiersema, J.H., Lohne, C., Barthlott, W., Wilde, V. 2007. Phylogeny analysis of *Nymphaea* (*Nymphaeaceae*): Evidence from substitutions and microstructural changes in the chloroplast trnT-trnF region. *International Journal of Plant Sciences*. 168:639-671.
- Lohne, C., Borsch, T., Wiersema, J.H. 2007. Phylogenetic analysis of *Nymphaeales* using fast-evolving and non-coding chloroplast markers. *Botanical Journal of the Linnean Society*. 154:141-163.
- Wiersema, J.H., Dahlberg, J. 2007. The nomenclature of *Sorghum bicolor* (L.) Moench (*Gramineae*). *Taxon*. 56(3):941-946.

1275-21000-263-00D

DEFINING THE GENETIC DIVERSITY AND STRUCTURE OF THE SOYBEAN GENOME AND APPLICATIONS TO GENE DISCOVERY IN SOYBEAN AND WHEAT GERMPLASM – Perry Cregan (P) and David Hyten; Beltsville, Maryland

- Hyten, D.L., Cannon, S.B., Song, Q., Weeks, N.T., Fickus, E.W., Shoemaker, R.C., Specht, J.E., May, G.D., Cregan, P.B. 2010. High-Throughput SNP Discovery through Deep Resequencing of a Reduced Representation Library to Anchor and Orient Scaffolds in the Soybean Whole Genome Sequence. *Biomed Central (BMC) Genomics*. 11:38.
- Hyten, D.L., Choi, I., Song, Q., Specht, J.E., Carter Jr, T.E., Shoemaker, R.C., Hwang, E., Matukumalli, L.K., Cregan, P.B. 2010. A high density integrated genetic linkage map of soybean and the development of a 1,536 Universal Soy Linkage Panel for QTL mapping. *Crop Science*. 50:960-968.
- Hyten, D.L., Song, Q., Fickus, E.W., Choi, I., Quigley, C.V., Hwang, E., Pastor Corrales, M.A., Cregan, P.B. 2010. High-throughput SNP discovery and assay development in Common Bean. *Biomed Central (BMC) Genomics*. 11:475.
- Song, Q., Jia, G., Zhu, Y., Hwang, E., Hyten, D.L., Cregan, P.B., Grant, D.M., Nelson, R. 2010. Abundance of SSR motifs and development of candidate polymorphic SSR markers (BARCSOYSSR_1.0) in soybean. *Crop Science*. 50:1950-1960.
- Hyten, D.L., Smith, J.R., Frederick, R.D., Tucker, M.L., Song, Q., Cregan, P.B. 2009. Bulk Segregate Analysis using the GoldenGate Assay to Locate the Rpp3 Locus that Confers Resistance to *Phakopsora pachyrhizi* (Soybean Rust) in Soybean. *Crop Science*. 49:265-271.
- Wu, X., Blake, S., Sleper, D.A., Shannon, J.G., Cregan, P.B., Nguyen, H.T. 2008. QTL and Additive and Epistatic Effects for SCN Resistance in PI 437654. *Theoretical and Applied Genetics*. 118:1093-1105.
- Gaitan-Solis, E., Choi, I., Quigley, C.V., Cregan, P.B., Tohme, J. 2008. Single nucleotide polymorphisms in common bean: their discovery and genotyping using a multiplex detection system. *The Plant Genome*. 1:125-134.

- Hyten, D.L., Song, Q., Choi, I., Yoon, M., Specht, J.E., Matukumalli, L.K., Shoemaker, R.C., Young, N.D., Cregan, P.B. 2008. High-throughput genotyping with the GoldenGate assay in the highly complex genome of soybean. *Theoretical and Applied Genetics*. 116:945-952.
- Wu, X., Zhong, G., Findley, S., Cregan, P.B., Stacey, G., Nguyen, H.T. 2008. Genetic marker anchoring by six-dimensional pools for development of a soybean physical map. *Biomed Central (BMC) Genomics*. 9:28.
- Choi, I., Hyten, D.L., Matukumalli, L.K., Song, Q., Chaky, J.M., Quigley, C.V., Chase, K., Lark, K.G., Reiter, R.S., Yoon, M., Hwang, E., Yi, S., Young, N.D., Shoemaker, R.C., Van Tassell, C.P., Specht, J.E., Cregan, P.B. 2007. A soybean transcript map: discovery and mapping of single nucleotide polymorphisms in soybean genes. *Genetics*. 176:685-696.
- Hyten, D.L., Choi, I., Song, Q., Shoemaker, R.C., Nelson, R.L., Costa, J.M., Specht, J.E., Cregan, P.B. 2007. Highly Variable Patterns of Linkage Disequilibrium in Multiple Soybean Populations. *Genetics*. 175:1937-1944.
- Hyten, D.L., Hartman, G.L., Nelson, R.L., Frederick, R.D., Concibido, V.C., Cregan, P.B. 2007. Map location of the Rpp1 locus that confers resistance to *Phakopsora pachyrhizi* (soybean rust) in soybean. *Crop Science*. 47:837-840.

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MOLECULAR CHARACTERIZATION AND DIVERSITY ASSESSMENT OF COCOA GERMLASM IN THE AMERICAS – Dapeng Zhang (P) and Lyndel Meinhardt; Beltsville, Maryland

- Motilal, L., Zhang, D., Umaharan, P., Mischke, B.S., Mooleedhar, V., Meinhardt, L.W. 2010. The relic Criollo cacao in Belize- genetic diversity and relationship with Trinitario and other cacao clones held in the International Cocoa Genebank, Trinidad. *Plant Genetic Resources: Characterization and Utilization*. Available at <http://dx.doi.org/10.1017/S1479262109990232>.
- Tiburcio, R.A., Costa, G.G., Carazzolle, M.F., Schuster, S.C., Carlson, J.E., Guiltinan, M.J., Bailey, B.A., Argueso, J.L., Mieckowski, P., Meinhardt, L.W., Pereira, G.A. 2009. Genes acquired by horizontal transfer are potentially involved in the evolution of phytopathogenicity in *Moniliophthora perniciosa* and *Moniliophthora roreri*, two of the major pathogens of cacao. *Journal of Molecular Evolution*. 70:85-97.
- Zhang, D., Boccara, M., Motilal, L., Mischke, B.S., Johnson, E.S., Butler, D., Bailey, B.A., Meinhardt, L.W. 2009. Molecular characterization of an earliest cacao (*Theobroma cacao* L.) collection from Peruvian Amazon using microsatellite DNA markers. *Tree Genetics and Genomes*. <http://dx.doi.org/10.1007/s11295-009-0212-2>.
- Johnson, E.S., Aime, M.C., Crozier, J., Flood, J., Schnell II, R.J. A new morpho-type of phytophthora palmivora on cacao in central america. *Proceedings of the International Cocoa Producer's Conference*. pp. 31-39.
- Johnson, E.S., Bekele, F., Brown, J.S., Song, J.H., Motamayor, J., Zhang, D., Schnell II, R.J., Meinhardt, L.W. 2009. Population Structure and Genetic Diversity of the Trinitario Cacao (*Theobroma Cacao* L.) from Trinidad and Tobago. *Crop Science*. 49:564-572.
- Guiltinan, M.J., Verica, J., Zhang, D., Figueira, A. 2008. Genomics of *Theobroma cacao*, "the Food of the Gods". In: Moore, P.H. & Ming, R. editors *Genomics of Tropical Crop Plants*. 1st edition. New York, NY: Springer. p. 145-170.
- Khan, N., Motilal, L.A., Sukha, D.A., Bekele, F.L., Iwaro, A.D., Bidaisee, P., Umaharan, L.H., Grierson, L.H., Zhang, D. 2008. Variability of butterfat content in cacao (*Theobroma cacao* L.): combination and correlation with other seed-derived traits at the International Cocoa Genebank, Trinidad. *Plant Genetic Resources*. <http://dx.doi.org/10.1017/S1479262108994132>.
- Lachenaud, P., Zhang, D. 2008. Genetic diversity and population structure in wild stands of cacao trees (*Theobroma cacao* L.) in French Guiana. *Forest Science*. 65(3):310.
- Meinhardt, L.W., Rincones, J., Bailey, B.A., Aime, M.C., Griffith, C., Zhang, D., Pereira, G. 2008. *Moniliophthora perniciosa*, the causal agent of witches' broom disease of cacao: What's new from this old foe. *Molecular Plant Pathology*. 9(5):577-588.
- Mondego, J.M., Carazzolle, M.F., Costa, G., Formighieri, L.P., Parizzi, L.P., Rincones, J., Cotomacci, C., Vidal, R.O., Estrela, R.C., Garcia, O., Castro, L.A., Gramacho, K.P., Goncalves, M.S., Neto-Goes, A., Barbosa, L.V., Guiltinan, M.J., Bailey, B.A., Meinhardt, L.W., Cascardo, J.C., Pereira, G.A. 2008. A genome survey of *Moniliophthora perniciosa* gives new insights into Witches' Broom Disease of cacao. *Biomed Central (BMC) Genomics*. 9:548.
- Motilal, L., Zhang, D., Umaharan, P., Mischke, B.S., Boccara, M., Pinney, S.M. 2008. Increasing accuracy and throughput in large-scale microsatellite fingerprinting of cacao field germplasm collections. *Tropical Plant Biology*. 2:23-37.
- Rincones, J., Scarpari, L.M., Mondego, J.M., Carazzolle, M.F., Formighieri, E.F., Costa, G.G., Barau, J.G., Vilas-Boas, L.A., Sabha, M., Azevedo, R.A., Dias, R., Cascardo, J.M., Meinhardt, L.W., Periera, G.A. 2008. Differential gene expression between the biotrophic-like and saprotrophic mycelia of the Witches' broom pathogen *Moniliophthora perniciosa*. *Molecular Microbiology*. 21:891-908.
- Zhang, D., Amores, F., Cedeno, A., Hermann, M., Meinhardt, L.W. 2008. Genetic diversity and population structure of cacao landraces in Northern and Central coastal Ecuador. *Conservation Genetics*. 9(2) 327-337.

- Formighieri, E.F., Tiburcio, R.A., Armas, E.D., Shimo, H.M., Carels, N., Goes-Neto, A., Araujo, M.R., Cotomacci, C., Carazzolle, M.F., Thomazella, D.P., Sardinha-Pinto, N., Rincones, J., Digiampietri, L.A., Carraro, D.M., Azeredo-Espin, A., Reis, S.F., Deckman, A.C., Gramacho, K., Goncalves, M.S., Moura-Neto, J.P., Luciana, V.B., Meinhardt, L.W., Cascardo, J.C., Pereira, G.A. 2008. The phytopathogenic basidiomycete *Moniliophthora perniciosa* senesces and contains a kalilo-like plasmid stably integrated in its large mitochondrial genome. *Mycological Research*. <http://apsjournals.apsnet.org/doi/pdf/10.1094/MPMI-21-7-0891>.
- Zhang, D., Mischke, B.S., Johnson, E.S., Mora, A., Phillips-Mora, W., Meinhardt, L.W. 2008. Molecular characterization of an International cocoa collection using microsatellite markers. *Tree Genetics & Genomics*. doi: <http://dx.doi.org/10.1007/s11295-008-0163-z>.
- Garcia, O., Macedo, J., Tiburcio, R., Zaparoli, G., Bittencourt, L.M., Ceita, G., Rincones, J., Micheli, F., Gesteira, A., Mariano, A., Schiavinato, M.A., Medrano, F., Meinhardt, L.W., Pereira, G.A., Cascardo, J.C. 2007. Characterization of Necrosis and Ethylene inducing Proteins (NEP) in the Basidiomycete *Moniliophthora perniciosa*, the Causal Agent of Witches' Broom in *Theobroma cacao*. *Mycological Research*. 111:443-455.
- Ceita, G.D., Macedo, J.N., Santos, T.B., Alemanno, L., Gesteira, A.D., Micheli, F., Mariano, A.C., Gramacho, K.P., Silva, D.D., Meinhardt, L.W., Mazzafera, P., Pereira, G.A., Cascardo, J.D. Involvement of calcium oxalate degradation during programmed cell death in *Theobroma cacao* tissues triggered by the hemibiotrophic fungus *Moniliophthora perniciosa*. *Plant Science*. 173 (2007) 106-117.
- Johnson, E.S., Mora, A., Schnell, R.J. 2007. Field guide efficacy in the identification of reallocated clonally propagated accessions of cacao (*Theobroma cacao* L.). *Genetic Resources and Crop Evolution*. 54:1301-1313.
- Johnson, E., Phillips, W., Bekele, F., Zhang, D., Schnell, R.J. 2007. Field guide to the clones of *Theobroma cacao*. *Proceedings of the International Cocoa Producer's Conference*. San Jose, Costa Rica Oct 9-14. Vol. I: 641-646.
- Zhang, D., Boccara, M., Motilal, L., Butler, D.R., Umaharan, P., Mischke, B.S., Meinhardt, L.W. 2006. "Refractario" cocoa – genetic identity and genetic structure assessed using microsatellite markers. *Conservation Genetics*. 9(2):327-337.

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ENHANCEMENT OF SMALL FRUIT GERMPLASM THROUGH GENOMIC CHARACTERIZATION AND GENETIC IMPROVEMENT WITH EMPHASIS ON DISEASE RESISTANCE – Mark Ehlenfeldt (P), Kimberly Lewers, James Polashock, And Shioh Wang; Beltsville, Maryland

- Ehlenfeldt, M.K., Martin Jr, R.B. 2010. Seed set, berry weight, and yield interactions in highbush blueberry cultivars (*Vaccinium corymbosum* L.) 'Bluecrop' and 'Duke'. *Journal of American Pomological Society*. 64:161-170.
- Ehlenfeldt, M.K., Polashock, J.J., Stretch, A.W., Kramer, M.H. 2010. Mummy Berry Fruit Rot and Shoot Blight Incidence in Blueberry: Prediction, Ranking, and Stability in a Long-term Study. *HortScience*. 45:92-97.
- Wang, S.Y., Chen, C. 2010. Effect of allyl isothiocyanate on antioxidant enzyme activities, flavonoids and fruit quality of blueberry (*Vaccinium corymbosum* L., cv. Duke). *Food Chemistry*. 122(2010)1153-1158.
- Wang, S.Y., Chen, C., Yin, J. 2010. Effect of allyl isothiocyanate on antioxidants and fruit decay of blueberries. *Food Chemistry*. 120:199-204.
- "Wang, S.Y., Millner, P.D. 2009. Effect of Compost Socks System on Antioxidant Capacity, Flavonoid Content, and Fruit Quality of Strawberries. *Journal of Agricultural and Food Chemistry*. 57:9651-9657."
- Cline, W.O., Ballington, J.R., Polashock, J.J. 2009. Blueberry Red Ringspot Observations and Findings in North Carolina. *Meeting Abstract*. 810:305-312.
- Ehlenfeldt, M.K., Finn, C.E. G-435 and ars 96-138, pink-fruited blueberry selections. *HortScience*. 42:172-173.
- Ehlenfeldt, M.K., Rowland, L.J., Ogden, E.L., Vinyard, B.T. 2009. Cold hardiness of southern-adapted blueberry genotypes and the potential for their use in northern adapted blueberry breeding. *Plant Breeding*. 128:393-396.
- Johnson-Cicalese, J., Vorsa, N., Polashock, J.J. 2009. Breeding for fruit rot resistance in *Vaccinium macrocarpon*. In: *Proceedings of the Ninth International Vaccinium Symposium*, July 14-16, 2008, Corvallis, Oregon. p.191-195.
- Polashock, J.J., Caruso, F.L., Oudemans, P.V., Mcmanus, P., Constantelos, C., Crouch, J. 2009. Species identification and variation in the North American cranberry fruit rot complex. In: *Acta Horticulturae International Symposium on Vaccinium Culture Proceedings*, July 13-16, 2008, Corvallis, Oregon. p. 395-399.
- Polashock, J.J., Ehlenfeldt, M.K., Crouch, J.A. 2009. Molecular detection and discrimination of blueberry red ringspot virus strains causing disease in cultivated blueberry and cranberry. *Plant Disease*. 23:720-726.
- Polashock, J.J., Oudemans, P.V., Caruso, F.L., Mcmanus, P., Crouch, J. 2009. Population structure of the North American cranberry fruit rot complex. *Plant Pathology*. 58:1116-1127.
- Vorsa, N., Johnson-Cicalese, J., Polashock, J.J. 2009. A blueberry by cranberry hybrid derived from a *Vaccinium darrowi* x (*V. macrocarpon* x *V. oxycoccus*) intersectional cross. *Proceedings of the Ninth International Vaccinium Symposium*, July 14-16, 2008, Corvallis, Oregon. p.187-189.

- Wang, S.Y., Chen, C., Wang, C.Y. 2009. The influence of light and maturity on fruit quality and flavonoid content of red raspberries. *Food Chemistry*. 112:676-684.
- Ehlenfeldt, M.K., Finn, C.E. G-435 and ars 96-138, pink-fruited blueberry selections. *HortScience*. 42:172-173.
- Wang, S.Y., Bowman, L., Ding, M. 2008. Methyl Jasmonate Enhances Antioxidant Activity, Flavonoid Content and Antiproliferation of Human Cancer Cells in Blackberries (*Rubus* spp.). *Journal of Agricultural and Food Chemistry*. 107:1261-1269.
- Black, B., Frisby, J., Lewers, K.S., Takeda, F., Finn, C.E. 2008. Heat unit model for predicting bloom dates in *Rubus*. *HortScience*. 43(7):2000-2004.
- Polashock, J.J. 2008. New and Emerging Pathogens of Blueberry. New Jersey Annual Vegetable Meeting Proceedings. p. 140.
- Oudemans, P., Polashock, J.J., Vinyard, B. 2008. Fairy Ring Disease of Cranberry: Assessment of Crop Losses and Impact on Cultivar Genotype. *Plant Disease*. 92:616-622.
- Feng, R., Wang, S.Y., Ni, H., Tourkova, I., Shulin, M., Yin, X. 2007. Cyanidin-3-rutinoside, a natural polyphenol antioxidant, selectively kills leukemic cells by induction of oxidative stress. *Oncogene*. 282:13468-13476
- Black, B., Ehlenfeldt, M.K. 2007. Foliar Applications of GA4+7 Reduce Flowering in Highbush Blueberry. *HortScience*, vol. 42, pp. 555-558.
- Polashock, J.J. 2007. A Multifaceted Approach to Cranberry Improvement. Meeting Abstract. p.17.
- Wang, S.Y., Ballington, J.R. 2007. Free radical scavenging capacity and antioxidant enzyme activity in deerberry (*vaccinium stamineum* L.). *Food Science and Technology Today*. 73:1352-1361.
- Wang, S.Y., Bowman, L., Ding, M. 2007. Variations in free radical scavenging capacity and antiproliferative activity among different genotypes of autumn olive (*elaeagnus umbellata* thunb).. *Planta Medica*. 73:451-460
- Wang, S.Y., Chen, C., Wang, C.Y., Chen, P. 2007. Resveratrol Content in Strawberry Fruit is Affected by Preharvest Conditions. *Journal of Agriculture and Food Chemistry*. 55:8269-8274.
- Wang, S.Y., Fordham, I.M. 2007. Differences in Chemical Composition and Antioxidant Capacity among Different Genotypes of Autumn Olive (*Elaeagnus umbellata* Thumb). *Journal of the Science of Food and Agriculture*. 45(4):402-409.
- Feng, R., Wang, S.Y., Ni, H., Tourkova, I., Shulin, M., Yin, X. 2007. Cyanidin-3-rutinoside, a natural polyphenol antioxidant, selectively kills leukemic cells by induction of oxidative stress. *Oncogene*. 282:13468-13476
- Wang, S.Y., Lewers, K.S. 2007. Antioxidant capacity and flavonoid contents in wild strawberries. *Journal American Society Hortscience*. 132(5):629-637.
- Wang, S.Y., Lewers, K.S., Bowman, L., Ding, M. 2007. Antioxidant Activities and Inhibition of Cancer Cell Proliferation in Wild Strawberries. *Journal American Society Hortscience*. 132(5):647-658.
- Polashock, J.J., Oudemans, P. 2006. Fairy ring disease increases host genetic diversity in cultivated cranberry. National American Phytopathology Meetings.

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OPERATING AND DEVELOPING THE GERMPLASM RESOURCES INFORMATION NETWORK FOR THE U.S. GENETIC RESOURCES PROGRAM – Gary Kinard (P); Beltsville, Maryland

- Arnaud, E., S. Dias, M. Mackay, P. Cyr, C. Gardner, P. Bretting, G. Kinard, L. Guarino, and S. Louafi. 2010. Chapter 11: A global portal enabling worldwide access to information on conservation and use of biodiversity for food and agriculture. Pp. 175-185 in L. Maurer and K. Tochtermann (eds.) *Information and Communication Technologies for Biodiversity Conservation and Agriculture*. Shaker Verlag, Aachen, Germany.

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GENOMICS AND PROTEOMICS APPROACHES TO BROADENING RESISTANCE OF SOYBEAN TO PESTS AND PATHOGENS – Benjamin Matthews (P), Bret Cooper, and Mark Tucker; Beltsville, Maryland

- Severin, A.J., Peiffer, G., Xu, W.W., Hyten, D.L., Bucciarelli, B., O'Rourke, J.A., Grant, D.M., Farmer, A.D., May, G.D., Vance, C.P., Shoemaker, R.C., Stupar, R.M., Bolon, Y. 2010. An integrative approach to genomic introgression mapping. *Plant Physiology*. DOI:10.1104/pp.110.158949.
- Tremblay, A., Hosseini, P., Alkharouf, N., Li, S., Matthews, B.F. 2010. Transcriptome Analysis of a Compatible Response by Glycine max to *Phakopsora pachyrhizi* Infection. *Plant Science*. 179:183-193.
- Tucker, M.L., Ping, X., Yang, R. 2010. 1-aminocyclopropane-1-carboxylic acid (ACC) concentration and ACC synthase expression in soybean roots and root tips and soybean cyst nematode (*Heterodera glycines*) colonized root pieces. *Journal of Experimental Botany*. 61(2):463-472.
- Klink, V., Houssein, P., Matsye, P., Alkharouf, N., Matthews, B.F. 2010. An expression analysis of syncytia isolated from the roots of the Glycine max (soybean) genotype PI 88788 undergoing a resistant reaction after infection by *Heterodera glycines* (soybean cyst nematode). *Plant Physiology and Biochemistry*. 48:176-193.

- Klink, V., Matthews, B.F. 2009. Emerging approaches to broaden the resistance of soybean to the soybean cyst nematode. *Plant Physiology*. 151:1017-1022.
- Klink, V., Housseini, P., Matsye, P., Alkharouf, N., Matthews, B.F. 2009. A gene expression analysis of syncytia isolated from the roots of the Glycine max (soybean) genotype PI 548402 (Peking) undergoing a resistant reaction after infection by *Heterodera glycines* (soybean cyst nematode). *Plant Molecular Biology*. 71:525-567.
- Tremblay, A., Li, S., Scheffler, B.E., Matthews, B.F. 2009. Laser capture microdissection (LCM) and expressed sequence tag analysis of uredinia formed by *Phakopsora pachyrhizi*, the causal agent of Asian soybean rust. *Physiological and Molecular Plant Pathology*. 73(6):163-174.

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MOLECULAR GENETIC APPROACHES TO PEST AND PATHOGEN RESISTANCE IN SUGAR BEET, BETA VULGARIS L. – Anna Smigocki (P) and L. David Kuykendall; Beltsville, Maryland

- Ghanem, M.E., Hichri, I., Smigocki, A.C., Albaceta, A., Fauconnier, M., Diatloff, E., Martinez-Andujar, C., Lutts, S., Dodd, I.C., Perez-Alfocea, F. 2011. Root-targeted hormonal biotechnology to improve crop stress tolerance. *Plant Cell Reports*. 30(5):807-823.
- Kuykendall, L.D., Shao, J.Y. 2011. Genes encoding a callose synthase and phytochrome A are adjacent to a MAP3Ka-like gene in *Beta vulgaris* USH20. *International Journal of Plant Genomics*. vol. 2011. Article ID 370548, 8 pages. doi:10.1153/2011/370548.
- Kuykendall, L.D., Burdekin, K.A., Shao, J.Y., Conway, L.B. 2011. A plant gene encodes a 'HSFA9-like' heat shock factor and is part of a cluster of orthologous genes including NPR1, CaMP and CK1 in *Beta vulgaris*, *Populus trichocarpa*, *Solanum lycopersicum* and *Vitis vinifera*. *Advanced Studies in Biology*. vol.3, 2011, no.1, 13-24.
- Ghanem, M.E., Albacete, A., Smigocki, A.C., Frebort, I., Pospisilova, H., Martinez-Andujar, C., Acosta, M., Sanchez-Bravo, J., Lutts, S., Dodd, I.C., Perez-Alfocea, F. 2010. Root-synthesised cytokinins induce salinity tolerance in tomato (*Solanum lycopersicum* L.). *Journal of Experimental Botany*. 62(1):125-140.
- Cingel, A., Vinterhalter, B., Vinterhalter, D., Smigocki, A.C., Ninkovic, S. 2010. Agrobacterium-mediated transformation of two Serbian potato cultivars (*Solanum tuberosum* L. cv. Dragacevka and cv. Jelica). *African Journal of Biotechnology*. 9(30):4644-4650.
- Kuykendall, L.D., Shao, J.Y., Kurushko, T., Burdekin, K.A. 2009. Stress Induced Expression of a *Beta vulgaris* L. Gene for a Chloroplast-Targeted Signal Calmodulin-Binding Protein. *Advanced Studies in Biology*. 1(7):323-332.

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DISCOVERING AND USING DISEASE RESISTANCE GENES IN PHASEOLUS VULGARIS FOR THE CONTROL OF RUSTS – Marcial Pastor Corrales (P); Beltsville, Maryland

- Goncalves-Vidigal, M., Cruz, A., Garcia, A., Vidigal Filho, P., Souza, L., Pastor Corrales, M.A., Mcclean, P., Kami, J., Gepts, P. 2010. Linkage mapping of the Phg-1 and Co-14 genes for resistance to angular leaf spot and anthracnose in the common bean cultivar AND 277. *Theoretical and Applied Genetics*. 122:893-903.
- Pastor Corrales, M.A., Wasonga, C.J., Porch Clay, T.G., Griffiths, P.D. 2010. Targeting gene combinations for broad spectrum rust resistance in heat tolerant snap beans developed for tropical environments. *Journal of the American Society for Horticultural Science*. 135(6):521-532.
- Urrea, C., Steadman, J.R., Pastor Corrales, M.A., Lindgren, D.T., Venegas, J.P., Coyne, D.P. 2009. Registration of Great Northern Common Bean Cultivar NE1-06-12 with Enhanced Disease Resistance to Common Bacterial Blight and Bean Rust. *Journal of Plant Registrations*. 3:219-222.
- Goncalves-Vidigal, M.C., Vidigal Filho, P.S., Pastor Corrales, M.A. 2008. A New Anthracnose Resistance Gene in Andean Common Bean Cultivar Jalo Listras Pretas. *Crop Science*. 49:133-138.
- Brick, M.A., Ogg, J.B., Singh, S.P., Schwartz, H.F., Johnson, J.J., Pastor Corrales, M.A. 2008. Registration of Drought Tolerant, Rust Resistant, High Yielding Pinto Bean Germplasm Line CO46348. *Journal of Plant Registrations*. 2:120-124.
- Miles, M.R., Pastor Corrales, M.A., Hartman, G.L., Frederick, R.D. 2007. Differential response of common bean cultivars to *Phakopsora pachyrhizi*. *Plant Disease*. 91:698-704.
- Pastor Corrales, M.A., Liebenberg, M.M., Sartorato, A., Arraes-Pereira, P.A. 2007. Reaction of common bean cultivars to the Asian soybean rust pathogen, *Phakopsora pachyrhizi*, under field conditions in South Africa and Brazil. *Bean Improvement Cooperative Proceedings*. 50:123-124.

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THE U.S. NATIONAL FUNGUS COLLECTIONS AND DEVELOPMENT OF ON-LINE RESOURCES ABOUT FUNGI – Amy Rossman (P); Beltsville, Maryland

- Braun, U., Farr, D.F., Minnis, A.M. 2009. Nomenclatural notes on some cercosporoid hyphomycetes. *Schlechtendalia*. 19:81-84.
- Minnis, D., Rossman, A.Y., Yun, H. 2008. Proposal to conserve the name *Olivea tectonae* (T.S. Ramakr. & K. Ramakr.) R.L. Mulder against *Olivea tectonae* (Racib.) Thirum (Basidiomycota). *Taxon*. 57:1355-1356.
- Park, J., Park, B., Veeraraghavan, N., Jung, K., Lee, Y., Blair, J.E., Geiser, D., Isard, S., Mansfield, M.A., Nikolaeva, E., Park, S., Kim, S.H., Greene, M., Ivors, K.L., Balci, Y., Peiman, M., Erwin, D.C., Coffey, M.D., Rossman, A.Y., Farr, D.F., Cline, E., Grunwald, N.J., Luster, D.G., Schrandt, J.K., Martin, F.N., Ribeiro, O.K., Makalowska, I., Kang, S. 2008. *Phytophthora* Database: A forensic database supporting the identification and monitoring of *Phytophthora*. *Plant Disease*. 92:966-972.
- Cline, E., Farr, D.F., Rossman, A.Y. 2008. A synopsis of *Phytophthora* with accurate scientific names, host range, and geographic distribution. *Plant Health Progress*. <http://dx.doi.org/10.1094/PHP-2008-0318-01-RS>.
<http://www.plantmanagementnetwork.org/sub/php/review/2008/phytophthora/>
- Cline, E., Vinyard, B.T., Edmonds, R. 2007. Spatial Effects of Retention Trees on Mycorrhizas and Mycorrhizae and Biomass of Douglas-fir Seedlings. *Canadian Journal of Forest Research* 37:430-438.

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DISSECTING COMPLEX TRAITS IN MAIZE AND BIOFUEL GRASSES BY APPLYING GENOMICS, BIOINFORMATICS, AND GENETIC RESOURCES – Edward Buckler IV (P) and Peter Bradbury; Ithaca, New York

- Tian, F., Bradbury, P., Brown, P., Sun, Q., Flint Garcia, S.A., Rocheford, T.R., McMullen, M.D., Holland, J.B., Buckler IV, E.S. 2011. Genome-wide association study of maize identifies genes affecting leaf architecture. *Nature Genetics*. 43 159:162.
- Buckler IV, E.S., Gur, A., Gibon, Y., Sulpice, R., Flint Garcia, S.A., McMullen, M.D., Stitt, M. 2010. Genetic Analysis of Central Carbon Metabolism Unveils an Amino Acid Substitution That Alters Maize NAD-Dependent Isocitrate Dehydrogenase Activity. *PLoS One*. 10.1371/journal.pone.0009991
- Costich, D., Friebe, B., Sheehan, M.J., Casler, M.D., Buckler IV, E.S. 2010. Genome-size variation in switchgrass (*Panicum virgatum*): flow cytometry and cytology reveal rampant aneuploidy. *The Plant Genome*. 3, No 3.
- Zhang, Z., Ersoz, E., Lai, C., Todhunter, R., Tiwari, H.K., Gore, M.A., Bradbury, P., Yu, J., Arnett, D., Ordovas, J.M., Buckler IV, E.S. 2010. Mixed Linear Model Approach Adapted for Genome-Wide Association Studies. *Nature Genetics*. 42:355-360.
- Bernardo, A.N., Bradbury, P., Ma, H., Hu, S., Bowden, R.L., Buckler IV, E.S., Bai, G. 2009. Discovery and mapping of single feature polymorphisms in wheat using affymetrix arrays. *Biomed Central (BMC) Genomics*. 10:251.
- Buckler IV, E.S., Ersoz, E.S., Jianming, Y. 2009. Applications of Linkage Disequilibrium and Association Mapping in Maize. In: A.L. Kriz, B.A. Larkins, editors. *Molecular Genetic Approaches to Maize Improvement*, New York, NY: Springer-Verlag Berlin Heidelberg. p. 173.
- Buckler IV, E.S., Gore, M., Wright, M., Ersoz, E., Bouffard, P., Jarvie, T., Hurwitz, B., Narechania, A., Harkins, T., Grills, G., Ware, D. 2009. Large-scale enrichment and discovery of gene-associated SNPs. *Nature Methods*. 2 (2):121-133.
- Buckler IV, E.S., Gore, M., Zhu, C., Yu, J. 2009. Status and Prospects of Association Mapping in Plants. *The Plant Genome*. 1(1):5-20.
- Buckler IV, E.S., Holland, J.B., McMullen, M.D., Kresovich, S., Acharya, C., Bradbury, P., Brown, P., Browne, C.J., Eller, M.S., Ersoz, E., Flint Garcia, S.A., Garcia, A., Glaubitz, J.C., Goodman, M., Harjes, C., Guill, K.E., Kroon, D., Larsson, S., Lepak, N.K., Li, H., Mitchell, S.E., Pressoir, G., Peiffer, J., Oropeza Rosas, M., Rocheford, T., Romay, C., Romero, S., Salvo, S.A., Sanchez Villeda, H., Sun, Q., Tian, F., Upadaya, N., Ware, D., Yates, H., Yu, J., Zhang, Z. 2009. The Genetic Architecture of Maize Flowering Time. *Science*. 325(5941):714-718.
- Buckler IV, E.S., Myles, S., Peiffer, J., Brown, P., Ersoz, E., Zhang, Z., Costich, D. 2009. Association mapping: critical considerations shift from genotyping to experimental design. *The Plant Cell*. 10.1105/tpc.109.068437.
- Buckler IV, E.S., Tian, F., Stevens, N.M. 2009. Tracking footprints of maize domestication and evidence for a massive selective sweep on chromosome 10. *Proceedings of the National Academy of Sciences*. 106:9979-9986
- Buckler IV, E.S., Yu, J., Zhang, Z., Tabanao, D.A., Gail, P., Kresovich, S., Todhunter, R.J. 2009. On the adequacy of numbers of background markers for relationship estimation and sample sizes for association mapping. *The Plant Genome*. 2(1):63-77.

- Gore, M.A., Chia, J., Elshire, R.J., Sun, Q., Ersoz, E.S., Hurwitz, B.L., Peiffer, J.A., McMullen, M.D., Grills, G.S., Ross-Ibarra, J., Ware, D., Buckler IV, E.S. 2009. A first generation haplotype map of maize. *Science*. 326(5956):1115-1117.
- Zhang, Z., Buckler IV, E.S., Casstevens, T., Bradbury, P. 2009. Software engineering the mixed model for genome-wide association studies on large samples. *Briefings in Bioinformatics*. 10(6):664-675.
- Buckler IV, E.S., Gore, M. 2008. An Arabidopsis haplotype map takes root. *Nature Genetics*. 39:1056-1057.
- Buckler IV, E.S., Hamblin, M.T., Warburton, M.L. 2008. Empirical Comparison of Simple Sequence Repeats and Single Nucleotide Polymorphisms in Assessment of Maize Diversity and Relatedness. *PLoS One*. 2(12):e1367.
- Buckler IV, E.S., Weber, A.L., Briggs, W.H., Rucker, J., Baltazar, B.M., Sanchez-Gonzalez, J.D., Feng, P., Doebley, J.F. 2008. The Genetic Architecture of Complex Traits in Teosinte (*Zea mays ssp. parviglumis*): New Evidence from Association Mapping. *Genetics*. 180(2):1221-1232.
- Stich, B., Moring, J., Piepho, H., Heckenberger, M., Buckler IV, E.S., Melchinger, A.E. 2008. Comparison of Mixed-Model Approaches for Association Mapping. *Genetics*. 178(3):1748-1754.
- Buckler IV, E.S., Harjes, C., Rocheford, T., Bai, L., Brutnell, T., Kandianis, C.B., Sowinski, S.G., Stapleton, A.E., Vallabhaneni, R., Williams, M., Wurtzel, E., Jianbing, Y. 2008. Natural genetic variation in the lycopene epsilon cyclase gene can enhance provitamin A biofortification of maize. *Science*. 319(5861):330-333.
- Buckler IV, E.S., Yu, J., Holland, J.B., McMullen, M.D. 2008. Genome-wide complex trait dissection through nested association mapping. *Genetics*. 178:539-551.
- Blakey, C., Costich, D., Sokolov, V., Islam-Faridi, M. 2007. *Tripsacum* genetics: from observations along a river to molecular genomics. *Maydica*. 52:81-99.
- Bradbury, P., Zhang, Z., Kroon, D., Casstevens, T., Ramdoss, Y., Buckler IV, E.S. 2007. Tassel: software for association mapping of complex traits in diverse samples. *Bioinformatics*. 23:2633-2635.
- Buckler IV, E.S., Ersoz, E., Yu, J. 2007. Applications of linkage disequilibrium and association mapping in crop plants. Book Chapter. In: *Genomics Assisted Crop Improvement Vol. 1: Genomics Approaches and Platforms*, R.K. Varshney and R. Ruberosa, editors. pp 97-119. Publisher: Springer, Dordrecht, The Netherlands
- Buckler IV, E.S., Esch, E., Szymaniak, J., Yates, H., Wojciech, P. 2007. Using Crossover Breakpoints in Recombinant Inbred Lines to Identify Quantitative Trait Loci Controlling the Global Recombination Frequency. *Genetics*. 177:1851-1858.
- Buckler IV, E.S., Stich, B., Yu, J., Maurer, H.P., Melchinger, A.E., Utz, H. 2007. Power to detect higher-order epistatic interactions in a metabolic pathway using nested association mapping and diallel association mapping. *Genetics*. Vol. 176:563-570
- Gore, M., Bradbury, P., Hogers, R., Kirst, M., Verstege, E., Van Oeveren, J., Peleman, J., Buckler IV, E.S., Van Eijk, M. 2007. Evaluation of Target Preparation Methods for Single Feature Polymorphism Detection in Large Complex Plant Genomes. *Crop Science*. 47:S135-S148.
- Sheehan, M., Kennedy, L., Costich, D., Brutnell, T. 2007. Subfunctionalization of PhyB1 and PhyB2 in the control of seedling and mature plant traits in maize. *Plant Journal*. 49:338-353.

1907-21000-030-00D

IDENTIFICATION OF FUNCTIONAL SEQUENCE IN PLANT GENOMES THROUGH BIOINFORMATIC, GENOMIC, AND GENETIC APPROACHES – Doreen Ware (P); Ithaca, New York

- Myles, S., Chia, J., Hurwitz, B., Simon, C.J., Zhong, G., Buckler IV, E.S., Ware, D. 2010. Rapid genomic characterization of the genus *Vitis*. *PLoS One*. 5(1):e8219.
- Brook, Iii, L., Strable, J., Ware, D., Nettleton, D., Scanlon, M.J. 2009. Microdissection of Shoot Meristem Functional Domains. *PLoS Genetics*. 5(5): e1000476. doi:10.1371/journal.pgen.1000476
- Liang, C., Mao, L., Ware, D., Stein, L. 2009. Evidence-based gene predictions in plant genomes. *Genome Research*. 10(2):1912-1923.
- Paterson, A.H., Bowers, J.E., Maher, C.A., Narechania, A., Zhang, L., Ware, D., Messing, J., Rokhsar, D.S. 2009. The *Sorghum bicolor* genome and the diversification of grasses. *Nature*. 457:551-556.
- Schnable, P., Ware, D., Fulton, R.S., Stein, J.C., Wei, F., Pasternak, S., Liang, C., Wing, R., Wilson, R., Zhang, L., Chia, J., Narechania, A. 2009. The B73maize genome: complexity, diversity, dynamics. *Science*. 326(5956):1112-1115.
- Youens-Clark, K., Faga, B., Stein, L., Ware, D. 2009. CMap 1.01: a comparative mapping application for the internet. *Bioinformatics*. 25(22):3040-3042.
- Zhang, L., Chia, J., Kumari, S., Stein, J.C., Liu, Z., Narechania, A., Maher, C.A., Guill, K., McMullen, M.D., Ware, D. 2009. Genome-wide characterization of maize miRNA genes. *PLoS Genetics*. 5(11): e1000716. doi:10.1371/journal.pgen.1000716.
- Zhou, S., Fusheng, W., Pasternak, S., Ware, D., Wing, R., Livny, M., Schwartz, D.C. 2009. A single molecule scaffold for the maize genome. *PLoS Genetics*. 5(11):1-14.

- Canaran, P., Buckler IV, E.S., Glaubitz, J., Stein, L., Sun, Q., Zhao, W., Ware, D. 2009. Panzea: An Update on New Content and Features. *Nucleic Acids Research*. 36:D1041-D1043.
- Wicker, T., Narechania, A., Sabot, F., Stein, J., Giang, V., Graner, A., Ware, D., Stein, N. 2008. Low-pass shotgun sequencing of the barley genome facilitates rapid identification of genes, conserved non-coding sequences and novel repeats. *Biomed Central (BMC) Genomics*. 9:518.
- Avraham, S., Tung, C., Jaiswal, P., Ilic, K., Kellogg, E., Mccouch, S., Pujar, A., Reiser, L., Rhee, S., Sachs, M.M., Schaeffer, M.L., Stein, L., Stevens, P., Vincent, L., Zapata, F., Ware, D. 2008. The Plant Ontology Database: A community resource for plant structure and developmental stages controlled vocabulary and annotations. *Nucleic Acids Research*. 36:D449-D454.
- Canaran, P., Glaubitz, J., Ware, D. 2008. HTML::GMap-A High Level Perl Wrapper Around the Google Maps(TM) API. *Bioinformatics*. 36(1):1041-1043.
- Jaiswal, P., Avraham, S., Buckler IV, E.S., Casstevens, T., Hebbard, C., Liang, C., Ni, J., Ravenscroft, D., Ren, L., Stein, L., Spooner, W., Tecle, I., Thomason, J., Wei Xuehong, Ware, D., Youens-Clark, K., Yap, I., Mccouch, S., Tung, C. 2008. Gramene: a growing plant comparative genomics resource. *Nucleic Acids Research*. 36:D947-D953.
- Kim, H., Hurwitz, B., Yu, Y., Collura, K., Gill, N., Miguel, P.S., Mullikin, J., Nelson, W., Wissotski, M., Braidotti, M., Kudrna, D., Goicoechea, J.L., Stein, L., Ware, D., Jackson, S., Soderlund, C., Wing, R. 2008. The Oryza map alignment project: Construction, alignment and analysis of 12 BAC fingerprint/end sequence framework physical maps that represent the 10 genome types of genus Oryza. *Genome Research*. 9:R45.1-R45.15.
- Lu, C., Jeong, D., Kulkarni, K., Pillay, M., Nobuta, K., German, R., Thatcher, D., Maher, C., Zhang, L., Ware, D., Liu, B., Cao, X., Meyers, B., Green, P. 2008. Genome-wide analysis for discovery of new rice miRNA reveals natural antisense miRNA (nat-miRNAs). *Proceedings of the National Academy of Sciences*. 105(12):4951-4956.
- Canaran, P., Stein, L., Ware, D. 2007. Look-Align: an interactive web-based multiple sequence alignment viewer with polymorphism analysis support. *Bioinformatics*. 22(7):885-886.
- Hass-Jacobus, B., Futrell-Griggs, M., Abernathy, B., Westerman, R., Goicoechea, J., Stein, J., Ware, D. 2007. Integration of hybridization-based markers (overgos) into physical maps for comparative and evolutionary explorations in the genus Oryza and in Sorghum. *Biomed Central (BMC) Genomics*. 7:199.
- Hufford, K., Canaran, P., Ware, D., McMullen, M.D., Gaut, B. 2007. Tissue-specific expression of maize domestication and crop improvement loci. *Plant Physiology*. 144:1642-1653.
- Ilic, K., Kellogg, E., Jaiswal, P., Zapata, F., Stevens, P., Vincent, L., Avraham, S., Reiser, L., Pujar, A., Sachs, M.M., Whitman, N., Mccouch, S., Schaeffer, M.L., Ware, D., Stein, L., Rhee, S. 2007. Plant Structure Ontology, Unified Vocabulary of Anatomy and Morphology of a Flowering Plant. *Plant Physiology*. 143:587-599.
- Jaiswal, P., Ni, J., Yap, I., Ware, D., Spooner, W., Youens-Clark, K., Ren, L., Liang, C., Zhao, W., Ratnapu, K., Faga, B., Canaran, P., Fogleman, M., Hebbard, C., Avraham, S., Schmidt, S., Casstevens, T., Buckler IV, E.S., Stein, L., Mccouch, S. 2007. Gramene: a bird's eye view of cereal genomes. *Nucleic Acids Research*. 34:D717-D723. Available: <http://nar.oxfordjournals.org/>.
- Zhao, W., Canaran, P., Jurkuta, R., Fulton, T., Glaubitz, J., Buckler IV, E.S., Doebley, J., Gaut, B., Goodman, M., Holland, J.B., Kresovich, S., McMullen, M.D., Stein, L., Ware, D. 2007. Panzea: A Database and Resource for Molecular and Functional Diversity in the Maize Genome. *Nucleic Acids Research*. 34:D752-D757.

1907-21000-031-00D

BIOINFORMATIC METHODS AND TOOLS TO PREDICT SMALL GRAIN FIELD PERFORMANCE – Jean Luc Jannink (P) and Peter Bradbury, Ithaca, New York

- Hamblin, M.T., Close, T.J., Bhat, P.R., Chao, S., Abraham, K., Blake, T., Brooks, W.S., Cooper, B., Griffey, C.A., Hayes, P.M., Hole, D.J., Horsley, R.D., Obert, D.E., Smith, K.P., Ullrich, S.E., Muehlbauer, G.J., Jannink, J. 2010. Population structure and linkage disequilibrium in US barley germplasm: implications for association mapping. *Crop Science*. 50:556-566.
- Heffner, E.L., Lorenz, A.J., Jannink, J., Sorrells, M.E. 2010. Plant breeding with genomic selection: potential gain per unit time and cost. *Crop Science*. 50:1681-1690.
- Iwata, H., Jannink, J. 2010. Marker genotype imputation in a low-marker-density panel with a high-marker-density reference panel: accuracy evaluation in barley breeding lines. *Crop Science*. 50:1269-1278.
- Jannink, J., Lorenz, A.J., Iwata, H. 2010. Genomic selection in plant breeding: from theory to practice. *Briefings in Functional Genomics and Proteomics*. 9:166-177.
- Jannink, J. 2010. Dynamics of long-term genomic selection. *Genetics*. 42:35.
- Heffner, E.L., Sorrells, M.E., Jannink, J. 2009. Genomic Selection for Crop Improvement. *Crop Science*. 49:1-12.
- Iwata, H., Ebana, K., Fukuoka, S., Hayashi, T., Jannink, J. 2009. Bayesian multilocus association mapping on ordinal and censored traits and its application to the analysis of genetic variation among Oryza sativa L. germplasms. *Theoretical and Applied Genetics*. 118(5):865-880.

- Iwata, H., Ebana, K., Uga, Y., Hayashi, T., Jannink, J. 2009. Whole genome association mapping of grain shape variation among *Oryza sativa* L. germplasms based on elliptic Fourier analysis. *Theoretical and Applied Genetics*. 114(8):1437-1449.
- Jannink, J., Iwata, H., Bhat, P.R., Chao, S., Wenzl, P., Muehlbauer, G.J. 2009. Marker imputation in barley association studies. *The Plant Genome*. 2:11-22.
- Tinker, N.A., Kilian, A., Rines, H.W., Bjornstad, A., Howarth, C.J., Jannink, J., Anderson, J.M., Rosnagel, B.G., Wight, C.P., Stuthman, D.D., Sorrells, M.E., Scoles, G.J., Eckstein, P.E., Ohm, H.W., Jackson, E.W., Tuveeson, S., Kolb, F.L., Molnar, S.J., Olsson, O., Carson, M.L., Ceplitis, A., Bonman, J.M., Federizzi, L., Langdon, T. 2009. New DArT markers for oat provide enhanced map coverage and global germplasm characterization. *Biomed Central (BMC) Genomics*. 10(39):1471-2164.
- Waugh, R., Muehlbauer, G.J., Jannink, J., Ramsay, L. 2009. Association genetics in barley. *Current Opinion in Plant Biology*. 12(2):218-222.
- Zhong, S., Dekkers, J., Jannink, J. 2009. Association-Based Genomic Selection in Cultivated Barley. *Genetics*. 182:355-364.
- Bodhireddy, P., Jannink, J., Nelson, J. 2009. Selective Advance for Accelerated Development of Recombinant Inbred QTL Mapping Populations. *Crop Science*. 49:1284-1294.
- Gutierrez, L., Nason, J.D., Jannink, J. 2009. Morphological Genetic Diversity of Worldwide Barley and Mega-Targets of Selection. *Crop Science*. 49:483-497.
- Yao, N., Jannink, J., White, P.J., Alavi, S. 2008. Impact of Dry Solids and Bile Acid Concentrations on Bile Acid Binding Capacity of Extruded Oat Cereals. *Journal of Agricultural and Food Chemistry*. 56:8672-8679.
- Jannink, J., Moreau, L., Charcosset, A., Charmet, G. 2008. Overview of QTL detection in plants and tests for synergistic epistatic interactions. *Genetica*. 136:225-236.
- Jannink, J. 2007. Qtl x genetic background interaction: application to predicting progeny value. *Euphytica*. 161:61-69.
- Zhong, S., Jannink, J. 2007. Using QTL results to discriminate among crosses based on their progeny mean and variance. *Genetics*. 177:567-576.

1907-21000-032-00D

DETERMINATION AND CHARACTERIZATION OF UNINTENDED EFFECTS IN GENETICALLY MODIFIED FOOD CROPS – Owen Hoekenga (P), James Giovannoni, and Leon Kochian; Ithaca, New York

- Hoekenga, O., Lung'Aho, M., Tako, E., Kochian, L.V., Glahn, R.P. 2011. Iron biofortification of maize grain. *Plant Genetic Resources: Characterization and Utilization*. 1-3. Available: <http://journals.cambridge.org/>.
- Dubois, P., Olsefski, G.T., Hoekenga, O., Flint Garcia, S.A., Brutnell, T. 2010. Physiological and Genetic Characterization of End-of-Day Far-Red Light Response in maize Seedlings. *Plant Physiology*. 10.1104/pp.110.159830.
- Krill, A.M., Kirst, M., Kochian, L.V., Buckler IV, E.S., Hoekenga, O. 2010. Association and linkage analysis of aluminum tolerance genes in maize. *PLoS One*. 5(4):e9958. doi:10.1371/journal.pone.0009958.

1907-21000-033-00D

GENOMICS APPROACHES FOR IMPROVING NUTRITIONAL QUALITY OF FOOD CROP SPECIES – James Giovannoni (P), Leon Kochian, Yong Yang, Li Li, and Theodore Thannhauser; Ithaca, New York

- Zhou, X., Lu, S., Ling, H., Li, L. 2011. The cauliflower orange gene enhances petiole elongation by suppressing expression of eukaryotic release factor 1. *New Phytologist*. 190:89-100.
- Pan, I., Giovannoni, J.J., Irisch, V. 2010. Functional diversification of AGAMOUS lineage genes in regulating tomato flower and fruit development. *Journal of Experimental Botany*. 61:1795-1806.
- Ramos, S., Rutzke, M., Hayes, R.J., Faquin, V., Guilherme, L.R., Li, L. 2010. Selenium accumulation in lettuce germplasm. *Planta*. 233:649-660.
- Kamenetzky, L., Asis, R., Bassi, S., De Godoy, F., Bermudez, L., Fernie, A., Vrebalov, J., Giovannoni, J.J., Rossi, M., Carrari, F. 2010. Genomic analysis of wild tomato introgressions determining metabolism- and yield-associated traits. *Plant Physiology*. 152:1772-1786.
- Harel-Beja, R., Portnoy, V., Fei, Z., Giovannoni, J.J., Ori, N., Lewinsohn, E., Tadmor, Y., Schaffer, A., Katzir, N. 2010. A genetic map of melon highly enriched with fruit quality QTLs and EST markers, including sugar and carotenoid metabolism genes. *Theoretical and Applied Genetics*. 121:511-533.
- Gonda, I., Bar, E., Portnoy, V., Schaffer, A., Tadmor, Y., Gepstein, S., Giovannoni, J.J., Katzir, N., Lewinsohn, E. 2010. Branched-chain and aromatic amino acid catabolism into aroma volatiles in *Cucumis melo* L. fruit. *Journal of Experimental Botany*. 61:1111-1123.

- Enfissi, E., Barneche, F., Ahmed, I., Lichtle, C., Giovannoni, J.J., Bowler, C., Bramley, P., Fraser, P. 2010. Integrative transcript and metabolite analysis of DE-ETIOLATED1 down-regulated tomato fruit reveals the underlying metabolic and cellular events associated with their nutritionally enhanced chemotype. *The Plant Cell*. 22:1190-1215.
- Elitzur, T., Vrebalov, J., Giovannoni, J.J., Ocampo, E., Goldschmidt, E., Friedman, H. 2010. The Regulation of MADS-box Gene Expression During Ripening of Banana and their Regulatory Interaction with Ethylene. *Journal of Experimental Botany*. 61:1523-1535.
- Chiu, L., Zhou, X., Burke, S., Wu, X., Prior, R.L., Li, L. 2010. The purple cauliflower arises from activation of a MYB transcription factor. *Plant Physiology*. 154:1470-1480.
- Fei, Z., Joung, J., Tang, X., Zheng, Y., Huang, M., Lee, J., Tieman, D., Alba, R., Klee, H., Giovannoni, J.J. 2010. Tomato Functional Genomics Database (TFGD): a comprehensive collection and analysis package for tomato functional genomics. *Nucleic Acids Research*. D1156-1163.
- Waller, J., Akhtar, T., Lara-Nunez, A., Gregory, J., Giovannoni, J.J., Hanson, A. 2010. Developmental and feedforward control of the expression of folate biosynthesis genes in tomato fruit. *Molecular Plant*. 3:66-77.
- Zhou, X., Li, L. 2010. Think outside of the box: selenium volatilization altered by a broccoli gene in the ubiquinone biosynthetic pathway. *Plant Signaling and Behavior*. 5:74-75.
- Joung, J., Corbett, A., Fellman, S., Tieman, D., Klee, H., Giovannoni, J.J., Fei, Z. 2009. Plant MetGenMAP: an integrative analysis system for plant systems biology. *Plant Physiology*. 151:1758-1768.
- Vrebalov, J., Pan, I., Arroyo, A., Chung, M., Poole, M., Rose, J., Seymour, G., Giovannoni, J.J., Irish, V. 2009. Fleshy fruit expansion and ripening are regulated by the tomato SHATTERPROOF gene, TAGL1. *The Plant Cell*. 21:3041:3062.
- Xiangjun, Z., Cooke, P.H., Li, L. 2009. Eukaryotic release factor 1-2 affects Arabidopsis responses to glucose and phytohormones during germination and early seedling development. *Journal of Experimental Botany*. 61:357-367.
- Yuan, Y., Chiu, L., Li, L. 2009. Transcriptional regulation of anthocyanin biosynthesis in red cabbage. *Planta*. 230:1141-1153.
- Stack, S., Royer, S., Shearer, L., Chang, S., Giovannoni, J.J., Westfall, D., Anderson, L. 2009. Role of fluorescence in situ hybridization (FISH) in sequencing the tomato genome. *Cytogenetics and Genome Research*. 124:339-350.

1907-21000-034-00D

GENOMIC APPROACHES TO IMPROVING TRANSPORT AND DETOXIFICATION OF SELECTED MINERAL ELEMENTS IN CROP PLANTS – Leon Kochian (P), Yong Yang, and Theodore Thannhauser; Ithaca, New York

- Hoekenga, O., Mwaniki, A., Glahn, R.P., Kochian, L.V., Lung'Aho, M.G., Szalma, S.J., Hart, J.J., Rutzke, M.A. 2011. Genetic and physiological analysis of iron content and bioavailability in maize kernels. *PLoS One*. 6(6): e20429. doi:10.1371/journal.pone.0020429.
- Maron, L., Guimaraes, C., Pineros, M., Magalhaes, J., Pleiman, J., Mao, C., Kochian, L.V. 2010. Two functionally distinct members of the MATE (multidrug and toxic compound extrusion) family of transporters potentially underlie two major Al tolerance QTL in maize. *Plant Journal*. 61:728-740.
- Shaff, J., Schultz, B., Craft, E.J., Clark, R., Kochian, L.V. 2010. GEOCHEM-EZ: a chemical speciation program with greater power and flexibility. *Plant and Soil Journal*. 330:207-214.
- Shulz, C., Kochian, L.V., Harrison, M. 2010. Genetic variation for root architecture, nutrient uptake and mycorrhizal colonisation in *Medicago truncatula* accessions. *Plant and Soil Journal*. Available: <http://www.springerlink.com/index/73J4KLH3M3607430.pdf>.
- Kuepper, H., Kochian, L.V. 2010. Transcriptional regulation of metal transport genes and mineral nutrition during acclimation to cadmium and zinc in the Cd/Zn hyperaccumulator, *Thlaspi caerulescens* (Ganges population). *New Phytologist*. 185(1):114-129.
- Ligaba, A., Kochian, L.V., Pineros, M. 2009. Phosphorylation at S384 regulates the activity of the TaALMT1 malate transporter that underlies aluminum resistance in wheat. *Plant Journal*. 60:411-423.
- Wang, Y., Mosebach, C.M., Kibbe, A.S., Ryhal, M.K., Jones, A.D., Palmer, J.A., Kochian, L.V. 2009. Generation of Arabidopsis mutants by heterologous expression of a full length cDNA library from tomato fruits. *Plant Molecular Biology Reporter*. 27:454-461.

1910-21000-019-00D

CONSERVATION AND CHARACTERIZATION OF GERMPLASM OF SELECTED VEGETABLE CROPS – Larry Robertson (P), Joanne Labate, and Angelo Baldo; Geneva, New York

- Kopsell, D.A., Carl, S., Deyton, D., Abney, K., Kopsell, D., Robertson, L.D. 2010. Characterization of Nutritionally Important Carotenoids in Welsh Onion Accessions. *HortScience*. 45(3):463-465.
- Griffiths, P.D., Marke, L.F., Robertson, L.D. 2009. Identification of Crucifer Accessions from the NC-7 and NE-9 Plant Introduction Collections that are Resistant to Black Rot (*Xanthomonas campestris* pv. *campestris*) Races 1 and 4. *HortScience* 44:284-288.
- Labate, J.A., Robertson, L.D., Baldo, A.M. 2009. Multilocus sequence data reveal extensive departures from equilibrium in domesticated tomato (*Solanum lycopersicum* L.). *Heredity* 103:257-267.
- Labate, J.A., Robertson, L.D., Wu, F., Tanksley, S.D., Baldo, A.M. 2009. EST, COSII, and arbitrary gene markers give similar estimates of nucleotide diversity in cultivated tomato (*Solanum lycopersicum*). *Journal of Theoretical and Applied Genetics*. 118:1005-1014.
- Labate, J.A., Grandillo, S., Fulton, T., Munos, S., Caicedo, A., Peralta, I., Ji, Y., Chetelat, R., Scott, J.W., Gonzalo, M.J., Francis, D., Yang, W., Van Der Knaap, E., Baldo, A.M., Smith-White, B., Mueller, L., Prince, J., Blanchard, N., Storey, D., Stevens, M., Robbins, M., Wang, J., Liedl, B., Oconnell, M., Stommel, J.R., Aoki, K., Iijima, Y., Slade, A., Hurst, S., Loeffler, D., Steine, M., Vafeados, D., Mcguire, C., Freeman, C., Amen, A., Goodstal, J., Facciotti, D., Van Eck, J., Causse, M. 2007. Tomato. Book Chapter. In: Kole, C. editor. *Genome mapping and molecular breeding in plants. Volume 5 Vegetables*. NY. Springer Publishing Co. p. 1-125.

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CONSERVATION AND UTILIZATION OF THE GENETIC RESOURCES OF APPLES, GRAPES, AND TART CHERRIES – Gan-Yuan Zhong (P) and Angela Baldo; Geneva, New York

- Baldo, A.M., Norelli, J.L., Farrell, R., Bassett, C.L., Aldwinckle, H., Malnoy, M. 2010. Identification of genes differentially expressed during interaction of resistant and susceptible apple cultivars (*Malus x domestica*) with *Erwinia amylovora*. *Biomed Central (BMC) Plant Biology*.
- Al-Turki, S., Shahba, M., Forsline, P.L., Stushnoff, C. 2008. Biodiversity of Total Phenolics, Antioxidant Capacity, and Juice Quality in Apple Cider Taxa. *Journal of Horticulture, Environment and Biotechnology*. 49(6):409-417.
- Su, L., Bukovac, M.J., Forsline, P.L., Van Nocker, S. 2008. Natural Variation in Fruit Abscission-Related Traits in Apple (*Malus*). *Euphytica*.

1910-21000-023-00D

DEVELOPMENT OF PEST, DISEASE RESISTANCE, AND STRESS TOLERANCE IN APPLE ROOTSTOCKS – Gennaro Fazio (P) and Angelo Baldo; Geneva, New York

- Dreesen, R., Vanholme, B., Luyten, K., Van Wynsberghe, L., Fazio, G., Roldan-Ruiz, I., Keulemans, J. 2010. Analysis of *Malus* S-RNase gene diversity based on a comparative study of European wild apple, old and modern apple cultivars. *Molecular Ecology. Molecular Breeding* DOI: 10.1007/s11032-010-9405-5.
- St. Laurent, A., Merwin, I., Fazio, G., Thies, J., Brown, M. 2010. Rootstock genotype succession influences apple replant disease and root-zone microbial community composition in an orchard soil. *Plant and Soil Journal*. DOI:10.1007/s11104-010-0522-2.
- Russo, N., Aldwinckle, H., Robinson, T., Fazio, G. 2008. Budagovsky 9 rootstock: uncovering a novel resistance to fire blight. *Acta Hort*. 793:321-324.
- Russo, N., Robinson, T., Fazio, G., Aldwinckle, H. 2008. Fire Blight Resistance of Budagovsky 9 Apple Rootstock. *Plant Disease*. 92(3):385-391.
- Beers, E., Cockfield, S., Fazio, G. 2007. Biology and management of woolly apple aphid, *Eriosoma lanigerum* (Hausmann), in Washington State. In: *Proceedings of the IOBC*, 30(4):37-42.
- Russo, N.L., Robinson, T.L., Fazio, G., Aldwinckle, H.S. 2007. Field Evaluation of Apple Rootstocks for Orchard Performance and Fire Blight Resistance. *HortScience*. 47:1517-1525.
- Fazio, G., Aldwinckle, H., Mcquinn, R., Robinson, T. 2006. Differential susceptibility to fire blight in commercial and experimental apple rootstock cultivars. *Acta Horticulture Proceedings (ISHS)* 704:527-530.
- Logiudice, N., Aldwinckle, H.S., Robinson, T.L., Fazio, G. 2006. The nature of resistance of the 'b.9' apple rootstock to fire blight. *Acta Hort*. 704:515-520.

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IMPROVING GRAPE ROOTSTOCK AND SCION PEST AND DISEASE RESISTANCE – Peter Cousins (P), Lance Cadle Davidson, Angelo Baldo, and Gan-Yuan Zhong; Geneva, New York

- Cadle Davidson, L.E., Wakefield, L., Seem, R.C., Gadoury, D.M. 2009. Specific isolation of RNA from the grape powdery mildew pathogen *Erysiphe necator*, an epiphytic, obligate parasite. *Journal of Phytopathology*. doi: 10.1111/j.1439-0434.2009.01578.x.
- Cadle Davidson, L.E. 2008. Variation within and between *Vitis* species for foliar resistance to the downy mildew pathogen *Plasmopara viticola*. *Plant Disease*. 92:1577-1584.
- Cadle Davidson, L.E. 2008. Monitoring pathogenesis of natural *Botrytis cinerea* infections in developing grape berries. *American Journal of Enology and Viticulture*. 59:387-395.
- Gee, C., Gadoury, D., Cadle Davidson, L.E. 2008. Ontogenic resistance to *Uncinula necator* varies by genotype and tissue type in a diverse collection of *Vitis* spp. *Plant Disease*. 92:1067-1073.

1910-21220-005-00D

GENETICS AND GENOMICS OF GRAPE GROWTH, DEVELOPMENT, AND QUALITY – Christopher Owens (P), Gan-Yuan Zhong, and Angelo Baldo; Geneva, New York

- Garris, A.J., Cousins, P.S., Ramming, D.W., Baldo, A.M. 2009. Parentage Analysis of Freedom Rootstock. *American Journal of Enology and Viticulture*. Am. J. Enol. Vitic. 60(3):357-361.
- Garris, A.J., Clark, L., Owens, C.L., McKay, S., Luby, J., Matthiason, K., Fennel, A. 2009. Mapping of photoperiod induced growth cessation in the wild grape *Vitis riparia* Michx. using microsatellite markers. *Journal of the American Society for Horticultural Science*. 134:261-272.
- Owens, C.L. 2008. Grapes. in *Breeding Temperate Fruit Crops: Germplasm to Genomics*, Hancock, J.F. (ed.) Kluwer Academic Publishers.
- This, P., Lacombe, T., Cadle Davidson, M., Owens, C.L. 2007. Wine grape (*Vitis vinifera* L.) color associates with allelic variation in the domestication gene *VvmybA1*. *Journal of Theoretical and Applied Genetics*. Vol 114: 723-730.

1931-21000-017-00D

GENETIC IMPROVEMENT OF FRUIT CROPS THROUGH FUNCTIONAL GENOMICS AND BREEDING – Ralph Scorza (P), Richard Bell, Ann Callahan, Zongrang Liu, Christopher Dardick, and Chinnathambi Srinivasan; Kearneysville, West Virginia

- Li, R., Mock, R.G., Fuchs, M., Halbrecht, J., Howell, B., Liu, Z. 2011. Molecular characterization of the partial RNA1 and RNA2 3' untranslated region of tomato ringspot virus isolates from North America. *Canadian Journal of Plant Pathology*. DOI: 10.1080/07060661.2011.536648.
- Manfre, A.J., Glenn, D.M., Nunez, A., Moreau, R.A., Dardick, C.D. 2011. Light quantity and photosystem function mediate host susceptibility to turnip mosaic virus via a salicylic acid-independent mechanism. *Molecular Plant-Microbe Interactions*. 24(3):315-327.
- Srinivasan, C., Liu, Z., Scorza, R. 2011. Ectopic expression of class 1 KNOX genes induce and adventitious shoot regeneration and alter growth and development of tobacco (*Nicotiana tabacum* L) and European plum (*Prunus domestica* L). *Plant Cell Reports*. 30:655-664.
- Dardick, C.D., Callahan, A.M., Chiozzotto, R., Schaffer, R., Piagnani, M., Scorza, R. 2010. Stone formation in peach fruit exhibits spatial coordination of the lignin and flavonoid pathways and similarity to *Arabidopsis* dehiscence. *BioMed Central Biology*. 8:13.
- Jung, K., Cao, P., Seo, Y., Dardick, C.D., Ronald, P. 2010. The rice kinase phylogenomics database: a guide for systematic analysis of the rice kinase super-family. *Trends in Plant Science*. 15(11):595-599.
- Petri, C., Scorza, R. 2010. Factors affecting adventitious regeneration from in vitro leaf explants of 'Improved French' plum, the most important dried plum cultivar in the USA. *Annals of Applied Biology*. 156:79-89.
- Singer, S.D., Liu, Z. 2010. Both the constitutive Cauliflower Mosaic Virus 35S and tissue-specific AGAMOUS enhancers activate transcription autonomously in *Arabidopsis thaliana*. *Plant Molecular Biology*. 74:293-305.
- Yang, Y., Singer, S., Liu, Z. 2010. Evaluation and comparison of insulation efficiency of three enhancer-blocking insulators in plants. *Plant Cell Reports*. DOI: 10.1007/s11240-010-9880-8.
- Yang, Y., Singer, S., Liu, Z. 2010. *Petunia* AGAMOUS enhancer-derived chimeric promoters specify a carpel-, stamen- and petal-specific expression pattern sufficient for engineering male and female sterility in tobacco. *Plant Molecular Biology Reporter*. DOI 10.1007/s11105-010-0215-z.

- Yang, Y., Singer, S., Liu, Z. 2010. Two similar but distinct second intron fragments from tobacco AGAMOUS homologs confer identical floral organ-specific expression sufficient for generating complete sterility in plants. *Planta*. 231:1159-1169.
- Callahan, A.M., Scorza, R., Dardick, C.D. 2009. Characterization of 'Stoneless': A naturally occurring, partially stoneless plum cultivar. *Journal of the American Society for Horticultural Science*. 134:120-125.
- Hily, J., Singer, S.D., Yang, Y., Liu, Z. 2009. A transformation booster sequence (TBS) from *Petunia hybrida* functions as an enhancer-blocking insulator in *Arabidopsis thaliana*. *Plant Cell Reports*. 28:1095-1104.
- Hu, D., Scorza, R. 2009. Analysis of the 'A72' peach tree growth habit and its inheritance in progeny obtained from crosses of 'A72' with columnar peach trees. *Journal of the American Society for Horticultural Science*. 134(2):236-243.
- Singer, S.D., Hily, J., Liu, Z. 2009. A 1-kb bacteriophage lambda fragment functions as an insulator to effectively block enhancer-promoter interactions in *Arabidopsis thaliana*. *Plant Molecular Biology Reporter*. 28:69-76.
- Sun, Q., Sun, H., Bell, R.L., Xin, L. 2009. Effect of polyvinyl alcohol on in vitro rooting capacity of shoots in pear clones (*Pyrus communis* L.) of different ploidy. *Plant Cell Tissue And Organ Culture*. 99:299-304.
- Sun, Q., Sun, H., Li, L., Bell, R.L. 2009. In vitro colchicine-induced polyploid plantlet production and regeneration from leaf explants of the diploid pear (*Pyrus communis* L.) cultivar, 'Fertility'. *Journal of Horticultural Science and Biotechnology*. 84(5):548-552.
- Elkins, R.B., Turner, J.D., Castagnoli, S., Seavert, C.F., Mitcham, E.J., Biasi, W.V., Colonna, A., Bell, R.L. 2008. Evaluation of potential alternative European pear cultivars for U.S. West Coast growers. *Acta Horticulturae*. 800:483-489.
- Hall, J.D., Cobb, J., Iqbal, M., Abidali, M., Liu, Z., Mount, D.W. 2008. UVH6, a plant homolog of the human/yeast TFIIH transcription factor subunit XPD/RAD3, regulates cold-stress genes in *Arabidopsis thaliana*. *Plant Molecular Biology Reporter*. 27:217-228.
- Manfre, A.J., Lahatte, G.A., Climer, C.R., Marcotte, Jr., W.R. 2008. Seed dehydration and the establishment of desiccation tolerance during seed maturation is altered in the *Arabidopsis thaliana* mutant *atem6-1*. *Plant And Cell Physiology*. 50(2):243-253.
- Manfre, A.J., Simon, A.E. 2008. Importance of coat protein and RNA silencing in satellite RNA/virus interactions. *Virology*. 379:161-167.
- Jung, K., Dardick, C.D., Lee, J., Phetsom, J., Xu, X., Seo, Y., Ouyang, S., Ly, E., An, K., Cho, Y., Lee, G., Buell, C., An, G., Ronald, P. 2008. Identification and functional analysis of light-responsive unique or paralogous gene family members in rice using a near genomic gene microarray. *PLoS Genetics*. 4(8):e1000164.
- Jung, K., Dardick, C.D., Phetsom, J., Canlas, P., Seo, Y., Shultz, M., Ouyang, S., Yuan, Q., Frank, B., Ly, E., Yi, Y., Hsia, A., An, K., Chou, H., Rocke, D., Lee, G., Schnable, P., An, G., Buell, R., Ronald, P. 2008. Construction and validation of a 45K rice oligonucleotide array. *PLoS One*. 3(10):e3337.
- Liu, Z., Hily, J. 2008. A simple and sensitive high-throughput GFP screening in woody and herbaceous plants. *Plant Cell Reports*. 28:493-501.
- Lu, R., Lee, G., Shultz, M., Dardick, C.D., Jung, K., Phetsom, J., Jia, Y., Rice, R.H., Goldberg, Z., Schnable, P.S., Ronald, P., Rocke, D.M. 2008. Assessing probe-specific dye and slide biases in two-color microarray data. *BMC Bioinformatics*. 9:314. doi:10.1186/1471-2105-9-314.
- Nagel, A., Schnabel, G., Petri, C., Scorza, R. 2008. Generation and characterization of transgenic plum lines expressing the *Gastrodia* anti-fungal protein. *HortScience*. 43:1514-1521.
- Park, C., Peng, Y., Bart, R., Chen, X., Ruan, D., Canlas, P.E., Dardick, C.D., Ronald, P.C. 2008. Rice Xb15, a protein phosphatase 2C, negatively regulates Xa21-mediated resistance and programmed cell death. *PLoS Biology*. 6(9):e231.
- Peng, Y., Bartley, L.E., Chen, X., Dardick, C.D., Ruan, R., Canlas, P.E., Ronald, P.C. 2008. OsWRKY62 is a negative regulator of basal and Xa21-mediated defense against *Xanthomonas oryzae* pv. *oryzae* in rice. *Molecular Plant*. 1(3):446-458.
- Petri, C., Webb, K.K., Hily, J., Dardick, C.D., Scorza, R. 2008. High transformation efficiency in plum (*Prunus domestica* L.): a new tool for functional genomics studies in *Prunus* spp. *Molecular Breeding*. DOI 10.1007/s11032-008-9200-8.
- Urtubia, C., Devia, J., Castro, A., Zamora, P., Aguirre, C., Tapia, E., Barba, P., Dell'Orto, P., Moynihan, M., Petri, C., Scorza, R., Prieto, H. 2008. Agrobacterium-mediated genetic transformation of *Prunus salicina*. *Plant Cell Reports*. DOI 10.1007/s00299-008-0559-0.
- Zagrai, I., Ravelonandro, M., Scorza, R., Minoiu, N., Zagrai, L. 2008. Bulletin of University of Agricultural Sciences and Veterinary Medicine Cluj-Napoca. University of Agricultural Sciences and Veterinary Medicine Cluj-Napoca. 65(1-2):358:365.
- Zagrai, I., Capote, N., Ravelonandro, M., Cambra, M., Zagrai, L., Scorza, R. 2007. Silencing mechanism of C5 transgenic plums is stable under challenge inoculation with heterologous viruses. *Journal of Plant Pathology*. 90:41-46.
- Kundu, J., Briard, P., Hily, J., Ravelonandro, M., Scorza, R. 2007. Role of the 25-26 nt siRNA in the resistance of transgenic *Prunus domestica* graft inoculated with plum pox virus. *Virus Genes*. 36:215-220.

- Damiano, C., Kondakova, V., Todorovska, E., Kamenova, I., Badjiakov, I., Scorza, R., Gentile, A., Monticelli, S., Atanassov, A. 2007. Improving regeneration and transformation for resistance to sharka in prunus. *Acta Horticulturae*. 738:583-587.
- Dardick, C.D., Chen, J., Richter, T., Shu, O., Ronald, P. 2007. The rice kinase database (rkd): a phylogenomic database for the rice kinome. *Plant Physiology*. 143:579-586.
- Elkins, R.B., Turner, J.D., Castagnoli, S., Seavert, C.F., Mitcham, E.J., Biasi, W.V., Colonna, A., Bell, R.L. 2007. Evaluation of potential alternative varieties for U.S. west coast pear growers. Pear International Symposium, May 22-26, 2007, Consolacao, Portugal.
- Hily, J., Scorza, R., Ravelonandro, M., Damsteegt, V.D., Bassett, C.L., Petri, C., Liu, Z. 2007. Plum pox virus coat protein gene intron-hairpin-RNA (ihpRNA) constructs provide resistance to Plum pox virus in *Nicotiana benthamiana* and *Prunus domestica*. *Journal of the American Society for Horticultural Science*. 132(6):850-858.
- Scorza, R., Hily, J., Callahan, A.M., Malinowski, T., Cambra, M., Capote, N., Zagrai, I., Damsteegt, V.D., Briard, P., Ravelonandro, M. 2007. Dereglulation of plum pox resistant transgenic plum 'Honeysweet'. *Acta Horticulturae*. 738:669-673.
- Dardick, C.D. 2007. Comparative expression profiling of *Nicotiana benthamiana* leaves systemically infected with three fruit tree viruses. *Molecular Plant-Microbe Interactions*. 20(8):1004-1017.
- Cox, K., Layne, D., Scorza, R., Schnabel, G. 2006. Gastrodia anti-fungal protein (GAFP) from the orchid *Gastrodia elata* confers resistance to multiple root disease pathogens in transgenic tobacco. *Planta*. 224:1373-1383.
- Rohila, J.S., Chen, M., Chen, S., Chen, J., Cerny, R., Dardick, C.D., Canlas, P., Xu, X., Gribskov, M., Kanrar, S., Zhu, J., Ronald, P., Fromm, M. 2006. Protein-protein interactions of tandem affinity purification (tap)-tagged protein kinases in rice. *Plant Journal*. 46:1-13.
- Scorza, R. 2006. Dereglulation of plum pox resistant transgenic plum 'Honeysweet'. *Compact Fruit Tree*. 39:29-30.
- Bell, R.L. 2006. Host resistance to pear psylla nymphal feeding in a diverse collection of *pyrus* germplasm. Entomological Society of America Annual Meeting. Paper No. D0574.
- Rohila, J.S., Chen, M., Chen, S., Chen, J., Cerny, R., Dardick, C.D., Canlas, P., Xu, X., Gribskov, M., Kanrar, S., Zhu, J., Ronald, P., Fromm, M. 2006. Protein-protein interactions of tandem affinity purification (tap)-tagged protein kinases in rice. *Plant Journal*. 46:1-13.

1931-21000-020-00D

USING FUNCTIONAL AND APPLIED GENOMICS TO IMPROVE STRESS AND DISEASE RESISTANCE IN FRUIT TREES – Michael Wisniewski (P), John Norelli, Timothy Artlip, and Carole Bassett; Kearneysville, West Virginia

- Wisniewski, M., Norelli, J., Bassett, C., Artlip, T., Macarisin, D. (2011) Ectopic expression of a novel peach (*Prunus persica*) CBF transcription factor in apple (*Malus x domestica*) results in short-day induced dormancy and increased cold hardiness. *Planta* 233:971-983.
- Soria-Guerra, R., Rosales-Mendoza, S., Gasic, K., Wisniewski, M.E., Korban, S.S., Band, M. 2011. Gene expression is highly regulated in early developing fruit of apple. *Plant Molecular Biology Reporter*. DOI 10.1007/s 11105-011-0300-y.
- Norelli, J.L., Malnoy, M., Borejsza-Wysocka, E., Aldwinckle, H.S. 2010. Long-term stability of attacin E expression in transformed apple after 12 years in the field and effect of the expression of this gene on the fruit characteristics. *BioMed Central (BMC) Biotechnology*. doi: 10.1186/1472-6750-10-41.
- Malnoy, M., Borejsza-Wysocka, E., Norelli, J.L., Flaishman, M., Gidoni, D., Aldwinckle, H.S. 2010. Genetic transformation of apple (*Malus x domestica*) without use of a selectable marker gene. *Tree Genetics and Genomes*. 6:423-433.
- Macarisin, D., Wisniewski, M.E., Bassett, C.L., Thannhauser, T.W. 2009. Proteomic analysis of B-aminobutyric acid priming and aba-induction of drought resistance in crabapple (*Malus pumila*): effect on general metabolism, the phenylpropanoid pathway and cell wall enzymes. *Plant Cell and Environment*. 32: 1612-1631.
- Karlson, D., Byard, S., Wisniewski, M.E., Li, J. 2010. Interspecific analysis of xylem freezing responses in *Acer* and *Betula*. *HortScience*. 45:165-168.

3602-21000-006-00D

IDENTIFICATION AND CHARACTERIZATION OF GENES IMPORTANT DURING SEED DEVELOPMENT IN LEGUMES – Karen Hudson (P); West Lafayette, Indiana

- Kim, K.S., Bellendir, S., Hudson, K. A., Hill, C.B., Hartman, G.L., Hyten, D., Hudson, M.E., Diers, B.W. 2010. Fine mapping the soybean aphid resistance gene *Rag1* in soybean. *Theoretical and Applied Genetics*. 20(5):1063-1071.

- Hudson, K.A. 2010. The Circadian Clock-Controlled Transcriptome of Developing Soybean Seeds. *The Plant Genome*. 3(1):1-11.
- Hudson, K.A., Kim, K., Diers, B., Hudson, M. 2009. Microarray-Based Genetic Mapping Using Soybean Near-Isogenic Lines and Generation of SNP Markers in the Rag1 Aphid-Resistance Interval. *The Plant Genome*. 1:89-98.

3602-22000-016-00D

MOLECULAR AND GENETIC MECHANISMS OF HESSIAN FLY RESISTANCE IN SOFT WINTER WHEAT – Richard Shukle (P), Brandon Schemerhorn, and Christopher Williams; West Lafayette, Indiana

- Arrueta, L., Shukle, R.H., Wise, I.L., Mitapalli, O. 2011. Gene Characterization of Two Digestive Serine Proteases in Orange Blossom Wheat Midge (*Sitodiplosis Mosellana*). *The Canadian Entomologist*. 142:6 p. 532-545.
- Behura, S.K., Shukle, R.H., Stuart, J.J. 2010. Assessment of Structural Variation and Molecular Mapping of Insertion Sites of Desmar-like Elements in the Hessian Fly Genome. *Insect Molecular Biology*. 19(4):109.
- Kosma, K., Nemacheck, J.A., Jenks, M., Williams, C.E. 2010. Changes in the Properties of Wheat Leaf Cuticle During Interactions with Hessian Fly. *Plant Journal*. 63:31-43.
- Yu, G., Williams, C.E., Harris, M.O., Cai, X., Mergoum, M., Xu, S.S. 2010. Development and Validation of Molecular Markers Closely LinKed to H32 for Resistance to Hessian Fly in Wheat. *Crop Science*. 50:1325-1332.
- Shukle, R.H., Subramanyam, S., Williams, C.E. 2010. Ultrastructural Changes in the Midgut of Hessian Fly Larvae Feeding on Resistant Wheat. *Journal of Insect Physiology*. 56:754-760.
- Li, H., Sun, L., Muir, W., Xie, J., Schemerhorn, B.J., Pittindrigh, B. 2009. Transcriptional Signatures in Response to Wheat Germ Agglutinin and Starvation in *Drosophila melanogaster* Larval Midgut. *Insect Molecular Biology*. p. 21-31.
- Schemerhorn, B.J., Crane, Y.M., Morton, P.K., Aggarwal, R., Stuart, J.J. 2009. Localization and Characterization of 170 BAC-derived clones and mapping of Ninety-Four Microsatellites in the Hessian Fly. *Journal of Heredity*. Available at: <http://jhered.oxfordjournals.org/cgi/content/abstract/esp045>.
- Williams, C.E., Saltzmann, K.D., Giovanini, M., Ohm, H. 2009. Transcript Profiles of Two Wheat Lipid Transfer Protein-encoding Genes are Altered During Attach by Hessian Fly Larvae. *Plant Physiology and Biochemistry*. 48:54-61.
- Shukle, R.H., Mittapalli, O., Morton, P.K., Chen, M. 2009. Characterization and expression analysis of a gene encoding a secreted lipase-like protein expressed in the salivary glands of larval Hessian fly, *Mayetiola destructor* (Say). *Journal of Insect Physiology*. 55(2):104-111.
- Hong-Mei, L., Buczkowski, G., Schemerhorn, B.J., Murdock, L.L., Pittindrigh, B.R. 2008. Transcriptomic profile of *Drosophila melanogaster* larval midgut and responses to oxidative stress. *Insect Molecular Biology*. 335-339.
- Mittapalli, O., Shukle, R.H. 2008. Molecular characterization and responsive expression of a defender against apoptotic cell death homologue from Hessian fly, *Mayetiola destructor*. *Comparative Biochemistry and Physiology. Part B* 149: 517-523.
- Saltzmann, K.D., Giovanini, M., Zheng, C., Williams, C.E. 2008. Virulent Hessian Fly Larvae Manipulate the Free Amino Acid Content of Host Wheat Plants. *Journal of Chemical Ecology*. 34:1401-1410.
- Subramanyam, S., Smith, D., Clemens, J.C., Webb, M.A., Sardesai, N., Williams, C.E., 2008. Functional Characterization of HFR-1, a high-mannose N-glycan-specific wheat lectin induced by Hessian fly larvae. *Plant Physiology*. 147:1412-1426.
- Hunt, G., Amdam, G.V., Schlupalius, D., Emore, C., Sardesai, N., Williams, C.E., Rueppell, O., Guzman-Novoa, E., Arechavaleta-Velasco, M., Chandra, S. 2007. Behavioral Genomics of Honey Bee foraging and nest defense. *Naturwissenschaften*. Available: <http://dx.doi.org/10.1007/s00114-006-0183-1>.
- Mittapalli, O., Neal, J.J., Shukle, R.H. 2007. Antioxidant defense response in a galling insect. *Proceedings of the National Academy of Sciences*. 104(6):1889-1894.
- Mittapalli, O., Neal, J.J., Shukle, R.H. 2007. Tissue and life state specificity of glutathione s-transferase expression in the Hessian fly, *Mayetiola destructor*: implications for resistance to host allelochemicals. *Journal of Insect Science*. 7(20):1-13.
- Omrakash, M., Sardesai, N., Shukle, R.H. 2007. CDNA cloning and transcriptional expression of a peritorphin-like gene in the Hessian fly, *Mayetiola destructor* (say). *Archives of Insect Biochemistry and Physiology*. 64:19-29.
- Shukle, R.H., Yoshiyama, M., Morton, P.K., Schemerhorn, B.J. 2007. Tissue and developmental expression of a gene from Hessian fly encoding an ABC-active-transporter protein during interactions with wheat. *Journal of Insect Physiology*. 55:146-154.
- Cho, K., Tores, N.L., Subramanyam, S., Deepak, S.A., Sardesai, N., Han, O., Williams, C.E., Ishii, H., Kubo, A., Iwahashi, H. 2006. Protein extraction/solubilization protocol for monocot and dicot plant gel-based proteomics. *Springer Verlag*. 49:413-420.

Giovanini, M.P., Saltzmann, K.D., Putoff, D., Gonzalo, M., Ohm, H.W., Williams, C.E. 2006. A novel wheat gene encoding a putative chitin-binding lectin is associated with resistance against Hessian fly. *Molecular Plant Pathology*. 8:69-82.

3607-21000-012-00D

IMPROVEMENT OF PEST RESISTANCE AND QUALITY TRAITS OF SOYBEAN – Rouf Mian (P) and Margaret Redinbaugh; Wooster, Ohio

- Bai, X., Wei, Z., Orantes, L., Jun, T., Mittapalli, O., Mian, R.M., Michel, A.P. 2010. Combining Next-Generation Sequencing Strategies for Rapid Molecular Resource Development from an Invasive Aphid Species, *Aphis glycines*. *PLoS One*. 5(6): e11370. doi:10.1371/journal.pone.0011370.
- Michel, A.P., Mian, R.M., Olivas, N.D. 2010. Detached Leaf and Whole Plant Assays for Soybean Aphid (*Aphis glycines*) Resistance Show Differential Responses among Resistance Sources and Biotypes. *Journal of Economic Entomology*. 103:949-957.
- Michel, A.P., Zhang, W., Mian, R.M. 2010. Genetic Diversity and Differentiation among Laboratory and Field Populations of the Soybean Aphid, *Aphis glycines*. *Bulletin of Entomological Research*. 27:1-8.
- Sung-Taeg, K., Mian, R.M. 2010. Genetic Map of the Powdery Mildew Resistance Gene in Soybean PI 243540. *Genome*. 53(5):400-405.
- Mian, R.M., Bond, J., Joobeur, T., Mengistu, A., Weibold, W., Grover, S., Wrather, A. 2009. Identification of Soybean Genotypes Resistant to *Cercospora soja* by Field Screening and Molecular Markers. *Plant Disease*. 93:408-411.
- Mian, R.M., Kang, S., Redinbaugh, M.G. 2009. Microsatellite Diversity of Soybean Genotypes Differing in Bean Pod Mottle Virus Leaf Symptom. *Genetic Resources and Crop Evolution*. 89:359-67.
- Michel, A.P., Zhang, W., Jung, J., Kang, S., Mian, R.M. 2009. Cross-Species Amplification and Polymorphism of Microsatellite Loci in the Soybean Aphid, *Aphis glycines*. *Journal of Economic Entomology*. 102:1389-1392.
- Michel, A.P., Zhang, W., Jung, J., Kang, S., Mian, R.M. 2009. Population Genetic Structure of the Soybean Aphid, *Aphis glycines*. *Environmental Entomology*. 38:1301-1311.
- Shannon, J., Jeong-Dong, L., Wrather, J., Sleper, D.A., Mian, R.M., Bond, J.P., Robbins, R.T. 2009. Registration of 'S99-2281' Soybean Germplasm Line with Resistance to Frogeye Leaf Spot and Three Nematode Species. *Journal of Plant Registrations*. 3:94-98.
- Mian, R.M., Bond, J., Joobeur, T., Mengistu, A., Weibold, W., Grover, S., Wrather, A. 2009. Identification of Soybean Genotypes Resistant to *Cercospora soja* by Field Screening and Molecular Markers. *Plant Disease*. 93:408-411.
- Kang, S., Mian, R.M., Hammond, R.B. 2008. Soybean Aphid Resistance in PI 243540 is Controlled by a Single Dominant Gene. *Crop Science*. 48:1744-1748.
- Mian, R.M., Cooper, R.L., Dorrance, A. 2008. Registration of Stout-Rps1k Soybean Germplasm with Phytophthora Stem and Root Rot Resistance. *Journal of Plant Registrations*. 2:255-257.
- Mian, R.M., Dorrance, A.E. 2008. Registration of 'Prohio' Soybean. *Journal of Plant Registrations*. 2:208-210.
- Mian, R.M., Hammond, R.B., St. Martin, S.K. 2008. New Plant Introductions with Resistance to the Soybean Aphid. *Crop Science*. 48:1055-1061.
- Mian, R.M., Kang, S., Beil, S.E. 2008. Genetic Linkage Mapping of the Soybean Aphid Resistance Gene in PI 243540. *Journal of Theoretical and Applied Genetics*. 117:955-962.
- Mian, R.M., Missaoui, A.M., Walker, D.R., Phillips, D.V., Boerma, H.R. 2008. Frogeye Leaf Spot of Soybean: A Review and Proposed Race Designations for Isolates of *Cercospora Sojina* Hara. *Crop Science*. 48:14-24.
- Ortiz-Perez, E., Mian, R.M., Cooper, R.L., Mendiola, T., Tew, J., Horner, H.T., Hanlin, S.J., Palmer, R.G. 2008. Seed-Set Evaluation of Four Male-Sterile, Female-Fertile Soybean Lines Using Alfalfa Leaf-Cutter Bees and Honey Bees as Pollinators. *Journal of Agricultural Science*. 146:461-469.

3611-21000-022-00D

CONSERVATION AND UTILIZATION OF MAIZE GENETIC STOCKS – Martin Sachs (P); Urbana, Illinois

- Sachs, M.M. 2009. Cereal Germplasm Resources. *Plant Physiology*. 149(1):148-151.
- Stinard, P.S., Kermicle, J.L., Sachs, M.M. 2009. The Maize enr System of r1 Haplotype-Specific Aleurone Color Enhancement Factors. *Journal of Heredity*. 100(2):217-228.
- Vartapetian, B.B., Sachs, M.M., Fagerstedt, K.V. 2008. Plant anaerobic stress II. Strategy of avoidance of anaerobiosis and other aspects of plant life under hypoxia and anoxia. *Plant Stress*. 2:1-19.
- Sachs, M.M., Vartapetian, B.B. 2007. Plant anaerobic stress I. Metabolic adaptation to oxygen deficiency. *Plant Stress*. 1(2):123-135.

- Avraham, S., Tung, C., Jaiswal, P., Ilic, K., Kellogg, E.A., McCouch, S.R., Pujar, A., Reiser, L., Rhee, S.Y., Sachs, M.M., Schaeffer, M.L., Stein, L.D., Stevens, P.F., Vincent, L.P., Zapata, F., Ware, D. 2007. The Plant Ontology Database: A Community Resource for Plant Structure and Developmental Stages Controlled Vocabulary and Annotations. *Nucleic Acids Research* 36: Database issue D449-D454
- Ilic, K., Kellogg, E.A., Jaiswal, P., Zapata, F., Stevens, P.F., Vincent, L.P., Avraham, S., Reiser, L., Pujar, A., Sachs, M.M., Whitman, N.T., McCouch, S.R., Schaeffer, M.L., Ware, D., Stein, L.D., Rhee, S.Y. 2006. Plant Structure Ontology. Unified Vocabulary of Anatomy and Morphology of a Flowering Plant. *Plant Physiology* 143:587-599
- Pujar, A., Jaiswal, P., Kellogg, E., Ilic, K., Vincent, L., Avraham, S., Stevens, P., Zapata, F., Reiser, L., Rhee, S., Sachs, M.M., Schaeffer, M.L., Stein, L., Ware, D., McCouch, S. 2006. Whole-plant growth stage ontology for angiosperms and its application in plant biology. *Plant Physiology*. 142:414-428.

3611-21000-023-00D

SOYBEAN GENETIC MANAGEMENT AND UTILIZATION – Steven Clough (P) and Randall Nelson; Urbana, Illinois

- Bilgin, D.D., Zavala, J.A., Zhu, J., Clough, S.J., Ort, D.R., Delucia, E.H. 2010. Biotic Stress Globally Down-Regulates Photosynthesis Genes. *Plant Cell and Environment*. doi: 10.1111/j.1365-3040.2010.02167.x.
- Betzelberger, A.M., Gillespie, K.M., Mcgrath, J.M., Koester, R.P., Nelson, R.L., Ainsworth, E.A. 2010. Effects of Chronic Elevated Ozone Concentration on Antioxidant Capacity, Photosynthesis and Seed Yield of 10 Soybean Cultivars. *Plant Cell and Environment*. 33(9):1569-1581. Available: doi: 10.1111/j.1365-3040.2010.02165.x.
- Mikel, M.A., Diers, B.W., Nelson, R.L., Smith, H.H. 2010. Genetic Diversity and Agronomic Improvement of North American Soybean Germplasm. *Crop Science*. 50:1219-1229.
- Shannon, J.G., Nelson, R.L., Lee, J.D., Wrather, J.A. 2010. Registration of LG04-6863 Soybean Germplasm Line with Diverse Pedigree. *Journal of Plant Registrations*. 4:70-72.
- Tian, Z., Wang, X., Lee, R., Li, Y., Specht, J.E., Nelson, R.L., McClean, P.E., Qiu, I., Ma, J. 2010. Artificial Selection for Determinate Growth Habit in Soybean. In: *Proceedings of the National Academy of Sciences*. PNAS 107(19):8563-8568. Available: www.pnas.org/cgi/doi/10.1073/pnas.1000088107.
- Zhang, M., Wu, Y., Lee, M., Liu, Y., Rong, Y., Santos, T.S., Wu, C., Xie, F., Nelson, R.L., Zhang, H. 2010. Numbers of Genes in the NBS and RLK Families Vary by More than Four-Fold Within a Plant Species and are Regulated by Multiple Factors. *Nucleic Acids Research*. Available: doi:10.1093/nar/gkq524.
- Zhu, J., Patzoldt, W.L., Radwan, O., Tranel, P.J., Clough, S.J. 2009. Effects of Photosystem II Interfering Herbicides Atrazine and Bentazon on the Soybean Transcriptome. *The Plant Genome*. 2:191-205.
- Calla, B., Vuong, T., Radwan, O., Hartman, G.L., Clough, S.J. 2009. Gene Expression Profiling Soybean Stem Tissue Early Response to *Sclerotinia sclerotiorum* and in Silico Mapping in Relation to Resistance Markers. *The Plant Genome*. 2:149-166.
- Hartman, G.L., Haudenschild, S. 2009. Movement of *Phakopsora pachyrhizi* (Soybean Rust) Spores by Non-Conventional Means. *European Journal of Plant Pathology*. 123:225-228.
- Lee, J.D., Shannon, J.G., So, Y.S., Sleper, D.A., Nelson, R.L., Lee, J.H., Choung, M.G. 2009. Environmental Effects on Lutein Content and Relationship of Lutein and Other Seed Components in Soybean. *Plant Breeding*. 128:97-100.
- Libault, M., Farmer, A., Brechenmacher, L., Franck, W.L., Drnevich, J., Langley, R.J., Bilgin, D.D., Radwan, O., Neece, D.J., Clough, S.J., May, G., Stacey, G. 2009. Complete Transcriptome of the Soybean Root Hair Cell, a Single Cell Model, and its Alteration in Response to *Bradyrhizobium japonicum* Infection. *Plant Physiology*. 152: 541-552.
- Bilgin, D.D., Delucia, E.H., Clough, S.J. 2009. A Robust Plant RNA Isolation Method for Affymetrix Genechip® Analysis and Quantitative Real-Time RT-PCR. *Nature Protocols*. 4:333-340.
- Bilgin, D.D., Aldea, M., O'Neill, B.F., Benitez, M., Li, M., Clough, S.J., Delucia, E.H. 2008. Elevated Ozone Alters Soybean-Virus Interaction. *Molecular Plant-Microbe Interactions*. 21(10):1297-1308.
- Brechenmacher, L., Moon-Young, K., Zou, J., Benitez, M., Li, M., Joshi, T., Calla, B., Lee, M., Philip, R., Libault, M., Vodkin, L.O., Xu, D., Lee, S., Clough, S.J., Stacey, G. 2008. Transcription profiling of soybean nodulation by *Bradyrhizobium japonicum*. *Molecular Plant-Microbe Interactions*. 21(5):631-645.
- Libault, M., Thibivilliers, S., Radwan, O., Clough, S.J., Stacey, G. 2008. Identification of four soybean reference genes for gene expression normalization. *The Plant Genome*. 1:44-54.
- Miles, M.R., Morel, W., Ray, J.D., Smith, J.R., Frederick, R.D., Hartman, G.L. 2008. Adult Plant Evaluation of Soybean Accessions for Resistance to *Phakopsora pachyrhizi* in the Field and Greenhouse in Paraguay. *Plant Disease*. 92(1):96-105.
- Li, Y., Zou, J., Li, M., Bilgin, D., Vodkin, L.O., Hartman, G.L., Clough, S.J. 2008. Soybean defense responses to the soybean aphid. *New Phytologist*. 179(1):185-195.
- Young, T., Diers, B., Hartman, G.L. 2008. Identification of QTL for Resistance to *Sclerotinia* Stem Rot (*Sclerotinia sclerotiorum*) in Soybean Plant Introduction 194639. *Crop Science*. 48:2209-2214.

- Wan, J., Zhang, X., Neece, D.J., Ramonell, K.M., Clough, S.J., Kim, S., Stacey, M.G., Stacey, G. 2008. A LysM Receptor-like Kinase Plays a Critical Role in Chitin Signaling and Fungal Resistance in Arabidopsis. *The Plant Cell*. 20:471-481.
- Wan, J., Zhang, X., Neece, D.J., Ramonell, K.M., Clough, S.J., Kim, S., Stacey, M.G., Stacey, G. 2008. A LysM receptor-like kinase plays a critical role in chitin signaling and fungal resistance in Arabidopsis. *The Plant Cell*. 20(2):471-481.
- Wang, W., Dia, V.P., Vasconez, M., Nelson, R.L., De Mejia, E.G. 2008. Analysis of Soybean Protein- Derived Peptides and the Effect of Cultivar, Environmental Conditions, and Processing on Lunasin Concentration in Soybean and Soy Products. *Journal of Association of Official Analytical Chemists International*. 91:936-946.
- Zhu, J., Patzoldt, W.L., Shealy, R.T., Vodkin, L.O., Clough, S.J., Tranel, P.J. 2008. Transcriptome response to glyphosate in sensitive and resistant soybean. *Journal of Agricultural and Food Chemistry*. 56:6355-6363.
- Miles, M.R., Levy, C., Morel W., Mueller, T., Steinlage, T., van Rij, N., Frederick, R.D., Hartman, G.L. 2007. International fungicide efficacy trials for management of soybean rust. *Plant Disease*. 91(11):1450-1458.
- Vodkin, L.O., Thibaud-Nissen, F., Gonzales, D.O., Zabala, G., Clough, S.J., Shealy, R. 2007. An Update on Soybean Functional Genomics and Microarray Resources for Gene Discovery and Crop Improvement. *Petria*. 17(1):43-53.
- Rasolohery, C.A., Berger, M., Lygin, A.V., Lozovaya, V.V., Nelson, R.L., Dayde, J. 2007. Effect of temperature and water availability during late maturation of the soybean seed on germ and cotyledon isoflavone content and composition. *Journal of the Science of Food and Agriculture*. 88:218-228.
- Guzman, P.S., Diers, B.W., St. Martin, S.K., Leroy, A.R., Grau, C.R., Hughes, T.J., Nelson, R.L. 2007. QTL for yield in three backcross-derived populations of soybean. *Crop Science*. 47: 111-122.
- Chen, Y., Gu, C., Mensah, C., Nelson, R.L., Wang, D. 2007. SSR marker diversity of soybean aphid resistance sources in northern America. *Crop Science*. 50:1104-1111.
- Hill, C., Li, Y., Hartman, G.L. 2006. A single dominant gene for resistance to the soybean aphid in the soybean cultivar dowing. *Crop Science*. 46(4):1601-1605.
- Kirakosyan, A., Kaufman, P., Nelson, R.L., Kasperbauer, M.J., Duke, J.A., Seymour, E., Chang, S., Warber, S., Bolling, S. 2006. Isoflavone levels in five soybean (*Glycine max*) genotypes are altered by phytochrome-mediated light treatments. *Journal of Agricultural and Food Chemistry*. 54(1):54-58.
- Zabala, G., Zou, J., Tuteja, J., Gonzalez, D.O., Clough, S.J., Vodkin, L.O. 2006. Transcription changes in the phenylpropanoid pathway of *Glycine max* in response to *Pseudomonas syringae* infection. *Biomed Central (BMC) Genomics*. 6:26.

3611-21000-024-00D

IDENTIFYING AND MANIPULATING DETERMINANTS OF PHOTOSYNTHATE PRODUCTION AND PARTITIONING – Donald Ort (P), Elizabeth Ainsworth, Carl Bernacchi, and Steven Huber; Urbana, Illinois

- Ainsworth, E.A., Bush, D.R. 2011. Carbohydrate Export from the Leaf - A Highly Regulated Process and Target to Enhance Photosynthesis and Productivity. *Plant Physiology*. 155(1):64-69.
- Ort, D.R., Zhu, X., Melis, A. 2011. Optimizing Antenna Size to Maximize Photosynthetic Efficiency. *Plant Physiology*. 155:79-85.
- Sun, J., Zhang, J., Larue, C.T., Huber, S.C. 2011. Decrease in Leaf Sucrose Synthesis Leads to Increased Leaf Starch Turnover and Decreased RuBP-limited Photosynthesis But Not Rubisco-limited Photosynthesis in Arabidopsis Null Mutants of SPSA1. *Plant Cell and Environment*. 34(4):592-604.
- Gillespie, K.M., Rogers, A., Ainsworth, E.A. 2011. Growth at Elevated Ozone or Elevated Carbon Dioxide Concentration Alters Antioxidant Capacity and Response to Acute Oxidative Stress in Soybean (*Glycine max*). *Journal of Experimental Botany*. 62(8):2667-2678.
- Ainsworth, E.A., Ort, D.R. 2010. How Do We Improve Crop Production in a Warming World? *Plant Physiology*. 154:526-530.
- Zhu, X., Long, S.P., Ort, D.R. 2010. Improving Photosynthetic Efficiency for Greater Yield. *Annual Reviews of Plant Biology*. 61:235-261.
- Long, S., Ort, D.R. 2010. More Than Taking the Heat: Crops and Global Change. *Current Opinion in Plant Biology*. 13(3):241-248.
- Oh, M., Wang, X., Wu, X., Zhao, Y., Clouse, S.D., Huber, S.C. 2010. Phosphorylation of Tyr-610 in the Receptor Kinase BAK1 Plays a Role in Brassinosteroid Signaling and Basal Defense Gene Expression. *Proceedings of the National Academy of Sciences*. 107(4):17827-17832.
- Rascher, U., Biskup, B., Leahey, A.D.B., McGrath, J.M., Ainsworth, E.A. 2010. Altered Physiological Function, Not Structure, Drives Increased Radiation-Use Efficiency of Soybean Grown at Elevated CO₂. *Photosynthesis Research*. 105(1):15-25.
- Feng, Z., Pang, J., Kobayashi, K., Zhu, J., Ort, D.R. 2010. Differential Responses in Two Varieties of Winter Wheat to Elevated Ozone Concentration Under Fully Open-air Field Conditions. *Global Change Biology*. doi: 10.1111/j.1365-2486.2010.02184.x.

- Calfapietra, C., Ainsworth, E.A., Beier, C., De Angelis, P., Ellsworth, D.S., Godbold, D.L., Hendrey, G.R., Hickler, T., Hoosbeek, M.R., Karnosky, D.F., King, J., Korner, C., Leakey, A., Lewin, K.F., Liberloo, M., Long, S.P., Lukac, M., Matyssek, R., Miglietta, F., Nagy, J., Norby, R.J., Oren, R., Percy, K.E., Rogers, A., Mugnozza, G.S., Stitt, M., Ceulemans, R. 2010. Challenges in Elevated CO₂ Experiments on Forests. *Trends in Plant Science*. 15(1):5-10.
- Oh, M., Clouse, S.D., Huber, S.C. 2009. Tyrosine Phosphorylation in Brassinosteroid Signaling. *Plant Signaling and Behavior*. 4(12):77-81.
- Rogers, A., Ainsworth, E.A., Leakey, A.D.B. 2009. Will Elevated Carbon Dioxide Concentration Amplify the Benefits of Nitrogen Fixation in Legumes? *Plant Physiology*. 151:1009-1016.

3620-22410-012-00D

MANAGEMENT AND GENETIC CHARACTERIZATION OF AGRICULTURAL AND BIOTECHNOLOGICAL MICROBIAL RESOURCES – David Labeda (P), Todd Ward, Stephen Peterson, Cletus Kurtzman, and Kerry Donnell; Peoria, Illinois

- Gale, L.R., Harrison, S.A., Ward, T.J., O Donnell, K., Milus, E.A., Gale, S.W., Kistler, H.C. 2011. Nivalenol-type populations of *Fusarium graminearum* and *F. asiaticum* are prevalent on wheat in Southern Louisiana. *Phytopathology*. 101(1):124-134.
- Koganti, S., Kuo, T., Kurtzman, C.P., Smith, N., Ju, L. 2011. Production of arabitol from glycerol: strain screening and study of factors affecting production yield. *Applied Microbiology and Biotechnology*. 90(1):257-267.
- Gilbert, J., Clear, R.M., Ward, T.J., Gaba, D., Tekauz, A., Turkington, T.K., Woods, S.M., Nowicki, T., O Donnell, K. 2010. Relative Aggressiveness and Production of 3- or 15-acetyl Deoxynivalenol and Deoxynivalenol by *Fusarium graminearum* in Spring Wheat. *Canadian Journal of Plant Pathology*. 32(2):146-152.
- Kato, A., Rooney, A.P., Furutani, Y., Hirose, S. 2010. Evolution of Trappin Genes in Mammals. *BMC Evolutionary Biology*. 10(31). Available: <http://www.biomedcentral.com/1471-2148/10/31>.
- Von Der Ohe, C., Gauthier, V., Tamburic-Illincic, L., Brule-Babel, A., Fernando, D.W., Clear, R., Ward, T.J., Miedaner, T. 2010. A Comparison of Aggressiveness and Deoxynivalenol Production Between Canadian *Fusarium graminearum* Isolates with 3-Acetyl and 15-Acetyldeoxynivalenol Chemotypes in Field-Grown Spring Wheat. *European Journal of Plant Pathology*. 127(3):407-417.
- Labeda, D.P., Price, N.P., Tan, G.A., Goodfellow, M., Klenk, H. 2010. Emendation of the Genus *Actinokineospora* Hasegawa 1988 and Transfer of *Amycolatopsis fastidiosa* Henssen et al. 1987 as *Actinokineospora fastidiosa* comb. nov. *International Journal of Systematic and Evolutionary Microbiology*. 60(6):1444-1449.
- Park, B., Park, J., Cheong, K., Choi, J., Jung, K., Lee, Y., Ward, T.J., O Donnell, K., Geiser, D.M., Kang, S. 2010. Cyber-infrastructure for *Fusarium* (CiF): Three integrated platforms supporting strain identification, phylogenetics, comparative genomics, and knowledge sharing. *Nucleic Acids Research*. doi:10.1093/nar/gkq1166.
- Connor, N., Sikorski, J., Rooney, A.P., Kopac, S., Koepfel, A.F., Burger, A., Cole, S.G., Perry, E.B., Krizanc, D., Field, N.C., Slaton, M., Cohan, F.M. 2010. Ecology of Speciation in the Genus *Bacillus*. *Applied and Environmental Microbiology*. 76(5):1349-1358.
- Du, Z., Rooney, A.P., Chen, G., Austin, B. 2010. *Corynebacterium marinum* sp. nov. isolated from coastal sediment in Qingdao, China. *International Journal of Systematic and Evolutionary Microbiology*. 60(pt8):1944-1947. doi:10.1099/ijs.0.018523-0.
- Rooney, A.P. 2009. Evolution of Moth Sex Pheromone Desaturases. *Annals of the New York Academy of Sciences*. 1170:506-510.
- Vaishampayan, P., Miyashita, M., Ohnishi, A., Satomi, M., Rooney, A.P., Duc, M.T., Venkateswaran, K. 2009. Description of *Rummeliibacillus stabekisii* gen. nov., sp. nov. and Reclassification of *Bacillus pycnus* Nakamura et al. 2002 as *Rummeliibacillus pycnus* comb. nov. *International Journal of Systematic and Evolutionary Microbiology*. 59(Pt 5):1094-1099.
- Balajee, A., Baddley, J.W., Peterson, S.W., Nickle, D., Varga, J., Boey, A., Lass-Flörl, C., Samson, R.A. 2009. *Aspergillus alabamensis*, a New Clinically Relevant Species in the Section *Terrei*. *Eukaryotic Cell*. 8(5):713-722.
- Nelson, D.M., Glawe, A.J., Labeda, D.P., Cann, I.K., Mackie, R.I. 2009. *Paenibacillus tundrae* sp. nov. and *Paenibacillus xylanexedens* sp. nov., Psychrotolerant, Xylan-Degrading, Bacteria from Alaskan Tundra. *International Journal of Systematic and Evolutionary Microbiology*. 59(pt. 7):1708-1714.
- Rooney, A.P., Price, N.P., Erhardt, C., Swezey, J.L., Bannan, J.D. 2009. Phylogeny and Molecular Taxonomy of the *Bacillus subtilis* species Complex and the Description of *Bacillus subtilis* subsp. *inaquosorum* subsp. nov. *International Journal of Systematic and Evolutionary Microbiology*. 59(Pt 10):2429-2436.

- Balajee, S., Kano, R., Baddley, J.W., Moser, S.A., Marr, K.A., Alexander, B.D., Andes, D., Kontoyiannis, D.P., Perrone, G., Peterson, S.W., Brandt, M.E., Pappas, P.G., Chiller, T. 2009. Molecular Identification of *Aspergillus* Species: Transplant Associated Infection Surveillance Network (TRANSNET). *Journal of Clinical Microbiology*. 47(10):3138-3141.
- Balajee, S.A., Borman, A.M., Brandt, M.E., Cano, J., Cuenca-Estrella, M., Dannaoui, E., Guarro, J., Haase, G., Kibbler, C.C., Meyer, W., O'Donnell, K., Petti, C.A., Rodriguez-Tudela, J.L., Sutton, D., Velegraki, A., Wickes, B.L. 2009. Sequence-Based Identification of *Aspergillus*, *Fusarium*, and *Mucorales* in the Clinical Laboratory: Where Are We and Where Should We Go From Here? *Journal of Clinical Microbiology*. 47(4):877-884.
- Kurtzman, C.P. 2009. Biotechnological Strains of *Komagataella* (*Pichia*) *pastoris* are *Komagataella phaffii* as Determined from Multigene Sequence Analysis. *Journal of Industrial Microbiology and Biotechnology*. 36(11):1435-1438.
- Labeda, D.P., Price, N.P., Donahue, J.M., Williams, N.M., Sells, S.F. 2009. *Streptomyces atriruber* sp. nov. and *Streptomyces silaceus* sp. nov.: New Species of Equine Origin. *International Journal of Systematic and Evolutionary Microbiology*. 59(11):2899-2903.
- Bishop, J.A., Chase, N., Kurtzman, C.P., Merz, W.G. 2008. Production of White Colonies on CHROMagar Candida(TM) by Members of the *Candida glabrata* Clade and Other Species with Overlapping Phenotypic Traits. *Journal of Clinical Microbiology*. 46(10):3498-3500.
- Bishop, J.A., Chase, N., Magill, S.S., Kurtzman, C.P., Fiandaca, M.J., Merz, W.G. 2008. *Candida bracarensis* Detected Among Isolates of *Candida glabrata* by Petide Nucleic Acid Fluorescence in Situ Hybridization: Susceptibility Data and Documentation of Presumed Infection. *Journal of Clinical Microbiology*. 46(2):443-446.
- Kurtzman, C.P. 2008. *Naumovozyma* Kurtzman, nom. nov. Myconet. Available: http://www.fieldmuseum.org/myconet/newnotes.asp#2008_03_05.
- Labeda, D.P., Kroppenstedt, R.M. 2008. Proposal for the new genus *Allokutzneria* gen. nov. within the suborder Pseudonocardineae and transfer of *Kibdelosporangium albatum* Tomita et al. 1993 as *Allokutzneria albata* comb. nov. *International Journal of Systematic and Evolutionary Microbiology*. 58:1472-1475.
- Labeda, D.P., Kroppenstedt, R.M., Euzéby, J.P., Tindall, B.J. 2008. Proposal of *Goodfellowiella* gen. nov. to replace the illegitimate genus name *Goodfellowia* Labeda and Kroppenstedt 2006. *International Journal of Systematic and Evolutionary Microbiology*. 58:1047-1048.
- Tamura, T., Ishida, Y., Otaguro, M., Hatano, K., Labeda, D.P., Price, N.P., Suzuki, K. 2008. Reclassification of *Streptomyces caeruleus* as a Synonym of *Actinoalloteichus cyanogriseus* and Reclassification of *Streptomyces spheroides* and *Streptomyces laceyi* as Later Synonyms of *Streptomyces niveus*. *International Journal of Systematic and Evolutionary Microbiology*. 58(12):2812-2814.
- Gale, L.R., Ward, T.J., Balmas, V., Kistler, H.C. 2007. Population subdivision of *Fusarium graminearum* sensu stricto in the Upper Midwestern United States. *Phytopathology*. 97(11):1434-1439.
- Goodfellow, M., Kumar, Y., Labeda, D.P., Sembiring, L. 2007. The *Streptomyces violaceusniger* clade: a home for streptomycetes with rugose ornamented spores. *Antonie Van Leeuwenhoek*. 92:173-199.
- Kurtzman, C.P. 2007. *Blastobotrys americana* sp. nov., *B. illinoisensis* sp. nov., *B. malaysiensis* sp. nov., *B. muscicola* sp. nov., *B. peoriensis* sp. nov. and *B. raffinofermentans* sp. nov., six anamorphic yeast species. *International Journal of Systematic and Evolutionary Microbiology*. 57:1154-1162.
- Kurtzman, C.P. 2007. New anamorphic yeast species: *Candida infanticola* sp. nov., *Candida polysorbophila* sp. nov., *Candida transvaalensis* sp. nov., and *Trigonopsis californica* sp. nov. *Antonie Van Leeuwenhoek*. 92(2):221-231.
- Kurtzman, C.P., Albertyn, J., Basehoar Powers, E.A. 2007. Multigene phylogenetic analysis of the Lipomycetaceae and the proposed transfer of *Zygozoma* species to *Lipomyces* and *Babjevia anomala* to *Dipodascopsis*. *Federation Of European Microbiological Societies Yeast Research*. 7:1027-1034.
- Labeda, D.P., Kroppenstedt, R.M. 2007. Proposal for *Umezawaea* gen. nov., a new genus of the Actinosynnemataceae related to *Saccharothrix* and transfer of *Saccharothrix tangerinus* Kinoshita et al. 2000 as *Umezawaea tangerina* gen. nov., comb. nov. *International Journal of Systematic and Evolutionary Microbiology*. 57 (12):2758-2761.
- Middelhoven, W.J., Kurtzman, C.P. 2007. Four novel yeasts species from decaying organic matter: *Blastobotrys robertii* sp. nov., *Candida cretensis* sp. nov., *Candida scorzettiae* sp. nov. and *Candida vadensis* sp. nov. *Antonie Van Leeuwenhoek*. 92:233-244.
- Samson, R.A., Hong, S., Peterson, S.W., Frisvad, J.C., Varga, J. 2007. Polyphasic taxonomy of *Aspergillus* section *Fumigati* and its teleomorph *Neosartorya*. *Studies in Mycology*. 59:147-203.
- Xue, B., Rooney, A.P., Kajikawa, M., Okada, N., Roelofs, W.L. 2007. Novel sex pheromone desaturases in the genomes of corn borers generated through gene duplication and retroposon fusion. *Proceedings of the National Academy of Sciences*. 104(11):4467-4472.
- Suh, S., Blackwell, M., Kurtzman, C.P., Lachance, M. 2006. Phylogenetics of *Saccharomycetales*, the ascomycete yeasts. *Mycologia*. 98(6):1008-1019.

3620-42000-041-00D

ENHANCED INSECT RESISTANCE AND CROP MANAGEMENT FOR REDUCTION OF MYCOTOXINS IN MIDWEST CORN
– Patrick Dowd (P), Eric Johnson, And Alejandro Rooney; Peoria, Illinois

- Dowd, P.F., Johnson, E.T., Pinkerton, T.S. 2010. Identification and Properties of Insect Resistance-Associated Maize Anionic Peroxidases. *Phytochemistry*. 71(11-12):1289-1297.
- Johnson, E.T., Berhow, M.A., Dowd, P.F. 2010. Constitutive Expression of the Maize Genes B1 and C1 in Transgenic Hi II Maize Results in Differential Tissue Pigmentation and Generates Resistance to *Helicoverpa zea*. *Journal of Agricultural and Food Chemistry*. 58(4):2403-2409.
- Dowd, P.F., Johnson, E.T. 2009. Differential Resistance of Switchgrass *Panicum virgatum* L. Lines to Fall Armyworms *Spodoptera frugiperda* (J. E. Smith). *Genetic Resources and Crop Evolution*. Available: <http://www.springerlink.com/content/4561455q7288w1l2/>.
- Dowd, P.F., Johnson, E.T. 2009. Field Incidence of Mycotoxins in Commercial Popcorn and Potential Environmental Influences. *Mycotoxin Research*. 26(1):15-22.
- Johnson, E.T., Berhow, M.A., Dowd, P.F. 2008. Colored and White Sectors from Star-Patterned *Petunia* Flowers Display Differential Resistance to Corn Earworm and Cabbage Looper Larvae. *Journal of Chemical Ecology*. 34:757-765.
- Dowd, P.F., Johnson, E.T., Pinkerton, T.S. 2007. Oral toxicity of beta-N-acetyl hexosaminidase to insects. *Journal of Agricultural and Food Chemistry*. 55(9):3421-3428.
- Johnson, E.T., Berhow, M.A., Dowd, P.F. 2007. Expression of a maize Myb transcription factor driven by a putative silk-specific promoter significantly enhances resistance to *Helicoverpa zea* in transgenic maize. *Journal of Agricultural and Food Chemistry*. 55(8):2998-3003.

3622-21000-030-00D

GENOMIC CHARACTERIZATION AND MANIPULATION OF VALUE-ADDED TRAITS FROM RYE FOR UTILIZATION IN WHEAT IMPROVEMENT – Perry Gustafson (P); Columbia, Missouri

- Emon, R.M., Islam, M.M., Begum, S.N., Jahiruddin, M., Islam, M.A., Hassan, M.M., Haque, M.S., Musket, T., Nguyen, H., Gustafson, J.P. 2010. Molecular Marker Based Characterization And Genetic Diversity Of Wheat Genotypes In Relation To Boron Efficiency. *Euphytica*. 70(4):339-348.
- Gustafson, J.P., Borlaug, N., Raven, P. 2010. World Food Supply and Biodiversity. *World Agriculture*. 1(2):38-41.
- Bento, M., Gustafson, J.P., Viegas, W., Silva, M. 2010. Genomes Behave as Social Entities: Alien Chromatin Minorities Evolve Through Specificities Reduction. *Theoretical and Applied Genetics*. 121(3):489-497. 10.1007/s00122-010-1325-6.
- Shi, B., Gustafson, J.P., Button, J., Miyazaki, J., Pallotta, M., Gustafson, N., Zhou, H., Langridge, P., Collins, N. 2009. Physical Analysis of the Complex Rye (*Secale cereale* L.) Alt4 Aluminium (Aluminum) Tolerance Locus Using a Whole-Genome BAC Library of Rye cv. Blanco. *Theoretical and Applied Genetics*. 119:695-704.
- Benito, M., Pereira, H.S., Rocheta, M., Gustafson, J.P., Viegas, W., Silva, M. 2008. Polyploidization as a Retraction Force in Plant Genome Evolution: Sequence Rearrangements in Triticale. *PLoS One*. Available at: <http://www.plosone.org/doi/pone.0001402>.
- Collins, N., Shirley, N.J., Saeed, M., Pallotta, M., Gustafson, J.P. 2008. An ALMT1 gene cluster controlling aluminium (aluminum) tolerance at the Alt4 locus of rye (*Secale cereale* L.). *Genetics*. 179(1):669-682.
- Gustafson, J.P., Ma, X., Korzun, V., Snape, J.W. 2008. A Consensus Map of Rye Integrating Mapping Data from Five Mapping Populations. *Theoretical and Applied Genetics*. 118:793-800.
- Ma, X., Gustafson, J.P. 2008. Allopolyploidization Accommodated Genomic Sequence Changes in Triticale. *Annals Of Botany*. 101(6):825-832.
- Chappell, A.S., Bilyeu, K.D. 2007. The low linolenic acid soybean line PI 361088B contains a novel GmFAD3A mutation. *Crop Science*. 47(4):1705.
- Ramalingam, J., Pathan, M.S., Feril, O., Miftahudin, Ross, K., Ma, X., Mahmoud, A.A., Layton, J., Rodriguez-Milla, M.A., Chikmawati, T., Valliyodan, B., Skinner, R., Matthews, D.E., Gustafson, J.P., Nguyen, H.T. 2006. Structural and functional analyses of the wheat genomes based on expressed sequence tags (ests) related to abiotic stresses. *Genome*. 49:1324-1340.

3622-21000-032-00D

MODIFICATION OF SOYBEAN SEED COMPOSITION FOR FOOD, FEED, AND OTHER INDUSTRIAL USES – Jan Miernyk (P), Perry Gustafson, Jason Gillman, Melvin Oliver, Kristin Bilyeu, and Hari Krishnan; Columbia, Missouri

- Miernyk, J.A., Hajduch, M. 2010. Seed proteomics. *Journal of Proteomics*. 74(4):389-400.

3622-21000-033-00D

GENETIC ENHANCEMENT OF SOYBEAN SEED VALUE BY BIOTECHNOLOGY – Ivan Baxter (P), Douglas Allen, and Melvin Oliver; Columbia, Missouri

Thiele, I., Huduke, D.R., Steeb, B., Fankam, G., Allen, D.K., Bazzani, S., Charusanti, P., Chen, F., Fleming, R.M., Hsiung, C.A., Dekeersmaecker, S.C., Liao, Y., Marchal, K., Mo, M.L., Ozdemir, E., Raghunathan, A., Reed, J.L., Shin, S., Sigurbjornsdottir, S., Steinmann, J., Sudarsan, S., Swainston, N., Thijs, I.M., Zengler, K., Palsson, B.O., Adkins, J.N., Dirk, B. 2011. A community effort towards a knowledge-base and mathematical model of the human pathogen *Salmonella Typhimurium* LT2. *BMC Systems Biology*. 5:8.

3622-21000-034-00D

GENETIC MECHANISMS AND MOLECULAR GENETIC RESOURCES FOR MAIZE – Michael McMullen (P), Mary Schaeffer, Melvin Oliver, and Sherry Flint Garcia; Columbia, Missouri

Schaeffer, M.L., Harper, E.C., Gardiner, J.M., Andorf, C.M., Campbell, D.A., Cannon, E.K., Sen, T.Z., Lawrence, C.J. 2011. MaizeGDB: Curation and outreach go hand-in-hand. *Database: The Journal of Biological Databases and Curation*. 2011. Article bar022. Available: <http://database.oxfordjournals.org/content/2011/bar022.full?keytype=ref&ijkey=TWrbetpkJmuYDzz>

Green, J.M., Harnsomburana, J., Schaeffer, M.L., Lawrence, C.J., Shyu, C. 2011. Multi-Source and Ontology-Based Retrieval Engine for Maize Mutant Phenotypes. *Database: The Journal of Biological Databases and Curation*. 2011:Article bar012. Available: <http://database.oxfordjournals.org/content/2011/bar012>.

Oliver, M.J., Jain, R., Balbuena, T.S., Agrawal, G.K., Gasulla, F., Thelen, J.J. 2010. Proteome analysis of leaves of the desiccation-tolerant grass, *sporobolus stapfianus*, in response to desiccation. *Phytochemistry*. 72(2011):1273-1284.

Zhao, Q., Weber, A., McMullen, M.D., Guill, K.E., Doebley, J. 2010. MADS-box Genes in Maize: Frequent Targets of Selection During Domestication. *Genetics Research*. 93(2011):65-75.

Bottoms, C., Flint Garcia, S.A., McMullen, M.D. 2010. IView: Introgression Library Visualization and Query Tool. *BMC Bioinformatics*. 11(6):S28.

Koster, K.L., Balsamo, R.A., Catherine, E., Oliver, M.J. 2010. Desiccation sensitivity and tolerance in the moss *Physcomitrella patens*: Assessing limits and damage. *Plant Growth Regulation*. 62:293-302.

Oliver, M.J., Murdock, A.G., Mishler, B.D., Kuehl, J.V., Boore, J.L., Mandoli, D.F., Everett, K.D., Wolf, P.G., Duffy, A.M., Karol, K.G. 2010. Chloroplast Genome Sequence of the Moss *Tortula ruralis*: Gene Content and Structural Arrangement Relative to Other Green Plant Chloroplast Genomes. *Biomed Central (BMC) Genomics*. 11:143.

Wei, F., Zhang, J., Zhou, S., He, R., Schaeffer, M.L., Collura, K., Kudrna, D., Faga, B., Wissotski, M., Golser, W., Fulton, L., Courtney, W., Fronick, C., Fulton, B., Coe, E., Schwartz, D.C., Ware, D., Clifton, S., Wilson, R.K., Wing, R.A. 2009. The Physical and Genetic Framework of the B73 Maize Genome. *PLoS Genetics*. 5(11):e1000715.

Flint Garcia, S.A., Buckler Iv, E.S., Tiffin, P., Ersoz, E., Springer, N.M. 2009. Heterosis is Prevalent for Multiple Traits in Diverse Maize Germplasm. *PLoS One*. 4:e7433.

Butron, A., Chen, Y., Rottinghaus, G., McMullen, M.D. 2009. Genetic Variation at Bx 1 Controls DIMBOA Content in Maize. *Theoretical and Applied Genetics*. 120:721-734.

Weber, A.L., Zhao, Q., McMullen, M.D., Doebley, J. 2009. Using Association Mapping in Teosinte (*Zea Mays* ssp *Parviglumis*) to Investigate the Function of Selection-Candidate Genes. *PLoS One*. 4:e8227.

3622-21000-035-00D

FUNCTIONAL GENOMICS FOR IDENTIFICATION OF SOYBEAN QUALITY TRAIT GENES – Yong Qiang An (P) and Melvin Oliver; Columbia, Missouri

An, Y., Meagher, R.B. 2010. The Strong Expression and Conserved Regulation of ACT2 in *Arabidopsis* and *Physcomitrella Patens*. *Plant Molecular Biology Reporter*. 28:481-490.

Zhang, M., Zhang, Y., Liu, L., Yu, L., Tsang, S., Tan, J., Yao, W., Kang, M., An, Y., Fan, X. Gene Expression Browser: Large-Scale and Cross-Experiment Microarray Data Management, Search & Visualization. *BMC Bioinformatics*. 11(433)1471-2105.

3625-21000-050-00D

GERMPLASM ENHANCEMENT OF MAIZE PROJECT (GEM) – Michael Blanco (P) and Candice Gardner; Ames, Iowa

- Jiang, H., Campbell, M., Blanco, M.H., Jane, J. 2010. Characterization of Maize Amylose-extender (ae) Mutant Starches. Part II: Structures and Properties of Starch Residues Remaining After Enzymatic Hydrolysis at Boiling-water Temperature. *Carbohydrate Polymers*. 80(1):1-12.
- Jiang, H., Horner, H.T., Pepper, T.M., Blanco, M.H., Campbell, M., Jane, J. 2010. Formation of Elongated Starch Granules in High-amylose Maize. *Carbohydrate Polymers*. 80(2):534-539.
- Jiang, H., Lio, J., Blanco, M.H., Campbell, M., Jane, J. 2010. Resistant-starch Formation in High-amylose Maize Starch During Kernel Development. *Journal of Agriculture and Food Chemistry*. 58:8043-8047.
- Srichuwong, S., Gutsea, J., Blanco, M.H., Duvick, S.A., Gardner, C.A., Jane, J. 2010. Characterization of Corn Grains for Ethanol Production. *Journal of ASTM International*. 7(2):1-10.
- Li, L., Jiang, H., Campbell, M., Blanco, M.H., Jane, J. 2008. Characterization of Maize Amylose-Extender (ae) Mutant Starches. Part I: Relationship Between Resistant Starch Contents and Molecular Structures. *Carbohydrate Polymers*. 74(3):396-404.

3625-21000-051-00D

THE MAIZE GENETICS AND GENOMICS DATABASE – Carolyn Lawrence (P), Craig Abel, and Taner Sen; Ames, Iowa

- Yennamalli, R.M., Rader, A.J., Wolt, J.D., Sen, T.Z. 2011. Thermostability in endoglucanases is fold-specific. *BMC Structural Biology*. 11(10):1-15.
- Andorf, C.M., Lawrence, C.J., Harper, E.C., Schaeffer, M.L., Campbell, D.A., Sen, T.Z. 2010. The Locus Lookup Tool at MaizeGDB: Identification of Genomic Regions in Maize by Integrating Sequence Information with Physical and Genetic Maps. *Bioinformatics*. 26(3):434-436.
- Bastow, R., Beynon, J., Estelle, M., Friesner, J., Grotewold, E., Lavagi, I., Lindsey, K., Meyers, B., Provart, N., Benfey, P., Birney, E., Braun, P., Brendel, V., Buell, R., Caccamo, M., Carrington, J., Cherry, M., Ecker, J., Eppig, J., Forster, M., Gutierrez, R., Hilson, P., Huala, E., Katari, M., Kersey, P., Kudla, J., Ma, H., Matsui, M., Matthews, K., May, S., Mayer, K., Millar, A., Millar, H., Mjolsness, E., Mockler, T., Nikolau, B., Nordborg, M., Rawlings, C., Schofield, P., Schoof, H., Schroeder, J., Sen, T.Z., Stanzione, D., Town, C., Toyoda, T., Vision, T., Walsh, S., Wang, X., Ware, D., Weckwerth, W., Yang, W. 2010. An international bioinformatics infrastructure to underpin the Arabidopsis community. *The Plant Cell*. 22(8):2530-2536.
- Sen, T.Z., Harper, E.C., Schaeffer, M.L., Andorf, C.M., Seigfried, T.E., Campbell, D.A., Lawrence, C.J. 2010. Choosing a Genome Browser for a Model Organism Database (MOD): Surveying the Maize Community. *Database: The Journal of Biological Databases and Curation*. doi: 10.1093/database/baq007. p. 1.
- Sen, T.Z., Andorf, C.M., Schaeffer, M.L., Harper, E.C., Sparks, M., Duvick, J., Brendel, V., Cannon, E., Campbell, D.A., Lawrence, C.J. 2009. MaizeGDB Becomes Sequence-centric. *Database: The Journal of Biological Databases and Curation*. doi: 10.1093/database/bap020. p. 1.
- Yi, G., Luth, D., Goodman, T.D., Lawrence, C.J., Becraft, P.W. 2009. High-throughput Linkage Analysis of Mutator Insertion Sites in Maize. *Plant Journal*. 58(5):883-892.
- Kurkcuoglu, O., Doruker, P., Sen, T.Z., Kloczkowski, A., Jernigan, R.L. 2008. The Ribosome Shape Directs mRNA Translocation through Entrance and Exit Dynamics. *Physical Biology*. 5(4):40005.
- Lawrence, C.J., Harper, E.C., Schaeffer, M.L., Sen, T.Z., Seigfried, T.E., Campbell, D.A. 2008. MaizeGDB: The Maize Model Organism Database for Basic, Translational, and Applied Research. *International Journal of Plant Genomics*. 2008:496957.
- Lushbough, C., Bergman, M.K., Lawrence, C.J., Jennewein, D., Brendel, V. 2008. BioExtract Server - An Integrated Workflow-enabling System to Access and Analyze Heterogeneous, Distributed Biomolecular Data. *IEEE/ACM Transactions on Computational Biology and Bioinformatics*. <http://doi.ieeecomputersociety.org/10.1109/TCBB.2008.98>.
- Kurkcuoglu, O., Doruker, P., Sen, T.Z., Kloczkowski, A., Jernigan, R.L. 2008. The Ribosome Shape Directs mRNA Translocation through Entrance and Exit Dynamics. *Physical Biology*. 5(4):40005.
- Duvick, J., Fu, A., Muppirala, U., Sabharwal, M., Wilkerson, M.D., Lawrence, C.J., Lushbough, C., Brendel, V. 2007. PlantGDB: A Resource for Comparative Plant Genomics. *Nucleic Acids Research*. 36:D959-965.
- Lawrence, C.J., Schaeffer, M.L., Seigfried, T.E., Campbell, D.A., Harper, L.C. 2007. MaizeGDB's New Data Types, Resources, and Activities. *Nucleic Acids Research*. 35:D895-D900.
- Lawrence, C.J., Walbot, V. 2007. Translational Genomics for Bioenergy Production from Fuelstock Grasses: Maize as the Model Species. *The Plant Cell*. 19(7):2091-2094.
- Lawrence, C.J., Walbot, V. 2007. Reply: Specific Reasons to Favor Maize in the U.S. *The Plant Cell*. 19(10):2973.

- Lushbough, C., Bergman, M.K., Lawrence, C.J., Jennewein, D., Brendel, V. 2007. Creating Bioinformatic Workflows within the BioExtract Server. In: BIOCOMP 07 Proceedings. The 2007 International Conference on Bioinformatics and Computational Biology, June 25-28, 2007, Las Vegas, NV. p. 18.
- Lawrence, C.J., Seigfried, T.E., Bass, H.W., Anderson, L.K. 2006. Predicting chromosomal locations of genetically mapped loci in maize using the Morgan2McClintock translator. *Genetics*. 172(3):2007-2009.

3625-21000-052-00D

CURATION AND DEVELOPMENT OF THE SOYBEAN BREEDER'S TOOLBOX AND ITS INTEGRATION WITH OTHER PLANT GENOME DATABASES – Randy Shoemaker (P), Rex Nelson, Craig Abel, David Grant, and Steven Cannon; Ames, Iowa

- Cannon, S.B., Ilut, D., Farmer, A.D., Maki, S.L., May, G.D., Singer, S.R., Doyle, J.J. 2010. Nodule Evolution Did Not Depend on Early Polyploidy in the Legumes. *PLoS One*. 5(7):e11630.doi:10.1371/journal.pone.0011630.
- Du, J., Tian, S., Hans, C., Jackson, S., Cannon, S.B., Shoemaker, R.C., Ma, J. 2010. Evolutionary Conservation, Diversity and Specificity of LTR Retrotransposons in Flowering Plants: New Insights from Genome-Wide Analysis and Multi-Specific Comparison. *Plant Journal*. 10. Available: <http://onlinelibrary.wiley.com/doi/10.1111/j.1365-313X.2010.04263>.
- Findley, S.D., Cannon, S.B., Varala, K., Du, J., Ma, J., Hudson, M.E., Birchler, J., Stacey, G. 2010. A Fluorescence In Situ Hybridization System for Karyotyping Soybean. *Genetics*. 185:727-744.
- Peto, M.F., Grant, D.M., Shoemaker, R.C., Cannon, S.B. 2010. Applying Small-Scale DNA Signatures as an Aid in Assembling Soybean Chromosome Sequences. *Advances in Bioinformatics*. doi:10.1155/2010/976792.
- Schmutz, J., Cannon, S.B., Schlueter, J., Ma, J., Hyten, D.L., Song, Q., Mitros, T., Nelson, W., May, G.D., Gill, N., Peto, M.F., Shu, S., Goodstein, D., Thelen, J.J., Cheng, J., Sakurai, T., Umezawa, T., Shinozaki, K., Du, J., Bhattacharyya, M., Sandhu, D., Grant, D.M., Joshi, T., Libault, M., Zhang, X., Hguyen, H., Valliyodan, B., Xu, D., Futrell-Griggs, M., Abernathy, B., Hellsten, U., Berry, K., Grimwood, J., Yu, Y., Wing, R.A., Cregan, P.B., Stacey, G., Specht, J., Rokhsar, D., Shoemaker, R.C., Jackson, S. 2010. Genome Sequence of the Paleopolyploid Soybean (*Glycine max* (L.) Merr.). *Nature*. 463:178-183.
- Cannon, S.B., May, G.D., Jackson, S.A. 2009. Update on Comparative Genomics of Legumes. *Plant Physiology*. 151:970-977.
- Grant, D.M., Nelson, R., Cannon, S.B., Shoemaker, R.C. 2009. SoyBase, The USDA-ARS Soybean Genetics and Genomics Database. *Nucleic Acids Research*. Doi: 10.1093/nar/gkp798.
- Nelson, R., Avraham, S., Shoemaker, R.C., May, G., Ware, D., Gessler, D.D. 2009. Applications and Methods Utilizing the Simple Semantic Web Architecture and Protocol (SSWAP) for Bioinformatics Resource Discovery and Disparate Data and Service Integration. *BioMed Central (BMC) BioData Mining*. 10:309.
- David, P., Chen, N., Pedrosa-Harand, A., Seignac, M., Thareau, V., Cannon, S.B., Debouck, D., Langin, T., Geffroy, V. 2009. A Nomadic Subtelomeric Disease Resistance Gene Cluster in Common Bean. *Plant Physiology*. 151:1048-1065.
- Gessler, D.D., Schiltz, G.S., May, G.D., Avraham, S., Town, C.D., Grant, D.M., Nelson, R. 2009. SSWAP: A Simple Semantic Web Architecture and Protocol for Semantic Web Services. *BMC Bioinformatics*. 10:309doi:10.1186/1471-2105-10-309.
- Grant, D.M., Nelson, R., Cannon, S.B., Shoemaker, R.C. 2009. SoyBase, The USDA-ARS Soybean Genetics and Genomics Database. *Nucleic Acids Research*. Doi: 10.1093/nar/gkp798.
- Nelson, R., Avraham, S., Shoemaker, R.C., May, G., Ware, D., Gessler, D.D. 2009. Applications and Methods Utilizing the Simple Semantic Web Architecture and Protocol (SSWAP) for Bioinformatics Resource Discovery and Disparate Data and Service Integration. *BioMed Central (BMC) BioData Mining*. 10:309.
- Zhang, X., Cannon, S.B., Stacey, G. 2009. Evolutionary Genomics of LysM Genes in Land Plants. *BMC Evolutionary Biology*. 9:183.
- Singer, S.R., Maki, S.L., Farmer, A.D., Ilut, D., May, G.D., Cannon, S.B., Doyle, J.J. 2009. Beyond the Papilionoids – What can We Learn from Chamaecrista? *Plant Physiology*. 151:1041-1047.
- Bertioli, D., Moretzsohn, M., Madsen, L.H., Sandal, N., Leal-Bertioli, S., Guimaraes, P., Hougaard, B.K., Fredslund, J., Schauser, L., Nielsen, A.M., Sato, S., Tabata, S., Cannon, S.B. 2009. An analysis of synteny of *Arachis* with *Lotus* and *Medicago* sheds new light on the structure, stability and evolution of legume genomes. *BMC Genomics*. 10:45.
- Wawrzynski, A., Ashfield, T., Chen, N.W., Mammadov, J., Nguyen, A., Podicheti, R., Cannon, S.B., Thareau, V., Ameline-Torregrosa, C., Cannon, E., Chacko, B., Couloux, A., Dalwani, A., Denny, R., Deshpande, S., Egan, A., Glover, N., Howell, S., Ilut, D., Lai, H., Martin Del Campo, S., Metcalf, M., O'Bleness, M., Pfeil, B., Ratnaparkhe, M., Roe, B., Maroof, M., Samain, S., Sanders, I., Segurens, B., Seignac, M., Sherman-Broyles, S., Tucker, D.M., Yi, J., Doyle, J.J., Geffroy, V., Roe, B., Maroof, M., Young, N., Innes, R.W. 2008. Replication of Nonautonomous Retroelements in Soybean Appears to be Both Recent and Common. *Plant Physiology*. 148:1760-1771.

- Ameline-Torregrosa, C., Wang, B., O'Bleness, M., Deshpande, S., Zhu, H., Roe, B., Young, N.D., Cannon, S.B. 2008. Identification and characterization of NBS-LRR encoded genes in the model plant *medicago truncatula*. *Plant Physiology*. 146:5-21.
- Ameline-Torregrosa, C., Cazaux, M., Danesh, D., Chardon, F., Cannon, S.B., Esquerre-Tugaye, M., Dumas, B., Young, N.D., Samac, D.A., Huguet, T., Jacquet, C. 2008. Genetic dissection of resistance to anthracnose and powdery mildew in *Medicago truncatula*. *Molecular Plant-Microbe Interactions*. 21(1):61-69.
- Cannon, S.B. 2008. Legume comparative genomics. In: Stacey, G. editor. *Genetics and Genomics of Soybean*. Series: Plant Genetics and Genomics: Crops and Models 2. New York, NY: Springer-Verlag. p. 35-54.
- Grant, D.M., Nelson, R., Graham, M.A., Shoemaker, R.C. 2008. Bioinformatic resources for soybean genetic and genomic research. In: Stacey, Gary, editor. *Genetics and Genomics of Soybean*. New York, New York. Springer. p. 141-162.
- Shoemaker, R.C., Grant, D.M., Olson, T., Warren, W.C., Wing, R., Yu, Y., Kim, H., Cregan, P.B., Joseph, B., Futrell-Griggs, M., Nelson, W., Davito, J., Walker, J., Wallis, J., Kremitski, C., Scheer, D., Clifton, S., Graves, T., Nguyen, H., Wu, X., Luo, M., Dvorak, J., Nelson, R., Cannon, S.B., Thomkins, J., Schmutz, J., Stacey, G., Jackson, S. 2008. Microsatellite Discovery from BAC End Sequences and Genetic Mapping to Anchor the Soybean Physical and Genetic Maps. *Genome*. 51:294-302.
- Febrer, M., Cheung, F., Town, C.D., Cannon, S.B., Young, N.D., Abberton, M., Jenkins, G., Milbourne, D. 2007. Construction, characterization and preliminary BAC-end sequencing analysis of a bacterial artificial chromosome library of white clover (*Trifolium repens* L.). *Genome*. 50:412-421.
- Zhang, X., Wu, X., Findley, S., Wan, J., Libault, M., Nguyen, H.T., Cannon, S.B. 2007. Molecular evolution of lysin motif-type receptor-like kinases in plants. *Plant Physiology*. 144:623-636.
- Cannon, S.B., Sterck, L., Rombauts, S., Sato, S., Cheung, F., Gouzy, J., Wang, X., Mudge, J., Vasdewani, J., Humphray, S.J., Nicholson, C., Spannagl, M., Schiex, T., Mayer, K., Schoof, H., Rogers, J., Quetier, F., Oldroyd, G., Debelle, F., Cook, D.R., Roe, B., Town, C.D., Tabata, S., Van De Peer, Y., Young, N.D. 2006. Legume genome evolution viewed through the *medicago truncatula* and *lotus japonicus* genomes. *Proceedings of the National Academy of Sciences*. 103(40):14959-14964.
- Leebens-Mack, J., Vision, T., Brenner, E., Bowers, J.E., Cannon, S.B., Clement, M.J., Cunningham, C.W., Depamphilis, C., Desalle, R., Doyle, J.J., Eisen, J.A., Gu, X., Harshman, J., Kellogg, E., Koonin, E.V., Philippe, H., Pires, J., Qiu, Y., Rhee, S.Y., Sjolander, K., Soltis, D.E., Stevens, P., Stevenson, D.W., Warnow, T., Zmasek, C., Soltis, P.S. 2006. Taking the first steps towards a standard for reporting on phylogenies: minimal information about a phylogenetic analysis (miapa). *OMICS: A Journal of Integrative Biology*. 10(20):231-237.
- Nelson, R., Shoemaker, R.C. 2006. Identification and Analysis of Gene Families from the Duplicated Genome of Soybean Using EST Sequences. *Biomed Central (BMC) Genomics*. 7:204.
- Young, N.D., Shoemaker, R.C. 2006. Genome studies and molecular genetics: exploring the structure function and evolution of legume genomes. *Current Opinion in Plant Biology*. 9:1-16.

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PLANT GENETIC RESOURCE AND INFORMATION MANAGEMENT – Candice Gardner (P), Charles Block, and Mark Widrechner; Ames, Iowa

- Kovach, D.A., Widrechner, M.P., Brenner, D.M. 2010. Variation in Seed Dormancy in *Echinochloa* and the Development of a Standard Protocol for Germination Testing. *Seed Science and Technology*. 38:559-571.
- Widrechner, M.P., Reitsma, K.R., Clark, L.D., Kirkbride, J.H. 2010. Length and Rapid Elongation of Pedicels of the Female Flowers of *Cucumis anguria* L. *Cucurbit Genetics Cooperative Report*. 31/32:36-40 and back cover.
- Arnaud, E., S. Dias, M. Mackay, P. Cyr, C. Gardner, P. Bretting, G. Kinard, L. Guarino, and S. Louafi. 2010. Chapter 11: A global portal enabling worldwide access to information on conservation and use of biodiversity for food and agriculture. Pp. 175-185 in L. Maurer and K. Tochtermann (eds.) *Information and Communication Technologies for Biodiversity Conservation and Agriculture*. Shaker Verlag, Aachen, Germany.
- Qu, L., Widrechner, M.P., Rigby, S.M. 2010. Analysis of Breeding Systems, Ploidy, and the Role of Hexaploids in Three *Hypericum perforatum* L. Populations. *Industrial Crops and Products*. 32(1):1-6.
- Widrechner, M.P. 2009. Ornamental Plants and the US National Plant Germplasm System: Conserving, Evaluating, Seeking, and Sharing. *International Plant Propagators Proceedings*. 58:213-217.
- Brindley, M.A., Widrechner, M.P., McCoy, J., Murphy, P., Hauck, C., Rizshsky, L., Nikolau, B., Maury, W. 2009. Inhibition of lentivirus replication by aqueous extracts of *Prunella vulgaris*. *Virology Journal*. 6:8.
- Birt, D.F., Widrechner, M.P., Hammer, K., Hillwig, M.L., Wei, J., Kraus, G.A., Murphy, P.A., McCoy, J.H., Wurtele, E.S., Neighbors, J.D., Wiemer, D.F., Maury, W.J., Price, J.P. 2009. *Hypericum* in Infection: Identification of Anti-viral and Anti-inflammatory Constituents. *Pharmaceutical Biology*. 47(8):774-782.
- Huang, N., Hauck, C., Yum, M., Rizshsky, L., Widrechner, M.P., McCoy, J., Murphy, P.A., Dixon, P.M., Nikolau, B.J., Birt, D.F. 2009. Rosmarinic Acid in *Prunella vulgaris* Ethanol Extract Inhibits LPS-induced Prostaglandin

- E2 and Nitric Oxide in RAW264.7 Mouse Macrophages. *Journal of Agricultural and Food Chemistry*. p. 10579-10589.
- Maury, W., Price, J., Brindley, M., Oh, C., Neighbors, J., Wiemer, D., Wills, N., Carpenter, S., Hauck, C., Murphy, P., Widrechner, M.P., Delate, K., Kumar, G., Kraus, G.A., Rizshsky, L., Nikolau, B. 2009. Identification of Light-independent Inhibition of Human Immunodeficiency Virus-1 Infection through Bioguided Fractionation of *Hypericum perforatum*. *Virology Journal*. 6:101. doi:10.1186/1743-422X-6-101. Available: <http://www.virologyj.com/content/pdf/1743-422X-6-101.pdf>.
- Widrechner, M.P., Thompson, J.R., Kapler, E.J., Kordecki, K., Dixon, P.M., Gates, G. 2009. A Test of Four Models to Predict the Risk of Naturalization of Non-native Woody Plants in the Chicago Region. *Journal of Environmental Horticulture*. 27(4):241-250.
- Wu, L., Dixon, P.M., Nikolau, B.J., Kraus, G.A., Widrechner, M.P., Wurtele, E. 2009. Metabolic Profiling of Echinacea Genotypes and a Test of Alternative Taxonomic Treatments. *Planta Medica*. 75:178-183.
- Birt, D.F., Widrechner, M.P., Lalone, C.A., Wu, L., Bae, J., Solco, A.K., Kraus, G.A., Murphy, P.A., Wurtele, E.S., Leng, Q., Hebert, S.C., Maury, W.J., Price, J.P. 2008. Echinacea in Infection. *American Journal of Clinical Nutrition*. 87(suppl.):488S-492S.
- Block, C.C., Shepherd, L.M. 2008. Long-term survival and seed transmission of *Acidovorax avenae* ssp. *citrulli* in melon and watermelon seed. *Plant Health Progress*. Available: <http://www.plantmanagementnetwork.org/sub/php/brief/2008/melon/>.
- Lopez, P.A., Widrechner, M.P., Simon, P.W., Rai, S., Boylston, T.D., Isbell, T., Bailey, T.B., Gardner, C.A., Wilson, L.A. 2008. Assessing phenotypic, biochemical, and molecular diversity in coriander (*Coriandrum sativum* L.) germplasm. *Genetic Resources and Crop Evolution*. 55:247-275.
- Widrechner, M.P., Kirkbride, J.H., Ghebretinsae, A.G., Reitsma, K.R. 2008. *Cucumis zambianus* (Cucurbitaceae): A New Species from Northwestern Zambia. *Systematic Botany*. 33:732-738.
- Cruz, V.V., Rife, C.L., Nason, J.D., Brummer, E.C., Gardner, C.A. 2008. Measuring the effectiveness of isolation of *Brassica napus* L. accessions during caged germplasm regeneration. *Plant Genetic Resources Newsletter*. 154:14-19.
- Flagel, L.E., Rapp, R.A., Grover, C.E., Widrechner, M.P., Hawkins, J., Grafenberg, J.L., Álvarez Fernández, I., Young Chung, G., Wendel, J.F. 2008. Phylogenetic, morphological, and chemotaxonomic incongruence in the North American endemic genus *Echinacea* (Moench). *American Journal of Botany*. 95:756-765.
- Cruz, V.V., Luhman, R., Rife, C.L., Shoemaker, R.C., Marek, L.F., Brummer, E., Gardner, C.A. 2007. Characterization of flowering time and SSR marker analysis of spring and winter type *Brassica napus* L. germplasm. *Euphytica*. 153:43-57.
- Haidet, M., Byrne, M., McCoy, J.H. 2007. Safeguarding the Seeds of Native Plants. *Herbal Gram*. 75:30-37.
- Lebeda, A., Stepankova, J., Krskova, M., Widrechner, M.P. 2007. Resistance in *Cucumis melo* germplasm to *Pseudoperonospora cubensis* pathotypes. *Advances in Downy Mildew Research*. 3:157-167.
- Widrechner, M.P. 2007. Old and New Trends Influencing the Introduction of New Nursery Crops. In: Janick, J. and Whipkey, A., editors. *Issues in New Crops and New Uses*. 2006 AAIC Annual Meeting and Sixth New Crops Symposium, October 14-18, 2006, San Diego, California. p. 237-245.
- Widrechner, M.P. 2007. While they were asleep: Do seeds after-ripen in cold storage? Experiences with *Calendula*. *International Plant Propagators Proceedings*. 56:377-382.
- Lalone, C.A., Hammer, K., Wu, L., Bae, J., Leyva, N., Liu, Y., Solco, A., Kraus, G.A., Murphy, P.A., Wurtele, E.S., Kim, O., Seo, K.I., Widrechner, M.P., Birt, D.F. 2007. Echinacea Species and Alkamides Inhibit Prostaglandin E2 Production in RAW264.7 Mouse Macrophage Cells. *Journal of Agricultural and Food Chemistry*. 55:7314-7322.
- Percifield, R., Hawkins, J., McCoy, J.H., Widrechner, M.P., Wendel, J. 2007. Genetic Diversity in *Hypericum* and AFLP Markers for Species-specific Identification of *H. perforatum* L. *Planta Medica*. 73:1614-1621. doi: 10.1055/s-2007-993749.
- Cruz, V.V., Nason, J.D., Luhman, R., Marek, L.F., Shoemaker, R.C., Brummer, E., Gardner, C.A. 2006. Analysis of bulked and redundant accessions of *Brassica* germplasm using assignment tests of microsatellite markers. *Euphytica*. 152:339-349.
- Hillwig, M., Wurtele, E., Hammer, K., Birt, D., Maury, W., Price, J., McCoy, J.H. 2006. Understanding bioactivity and metabolite diversity in St. John's Wort and related species [abstract]. *The Second Scientific Meeting of the Metabolomics Society*. 151:106.
- McCoy, J.H. 2006. Preservation of medicinal and aromatic crops [abstract]. *Association for the Advancement of Industrial Crops Conference*. p. 5.
- McCoy, J.H., Davis, J.M., Camper, N.D., Kahn, I., Bharathi, A. 2006. Influence of rhizome propagule size on yields and triterpene glycoside concentrations of black cohosh (*Actaea Racemosa* L.). *HortScience*. 42(1):61-64.
- Ortiz-Perez, E., Horner, H.T., Hanlin, S.J., Palmer, R.G. 2006. Insect-mediated seed-set evaluation of 21 soybean lines segregating for male sterility at 10 different loci. *Euphytica*. 152:351-360.
- Senchina, D.S., Wu, L., Flinn, G.N., Konopka, D.N., McCoy, J.H., Widrechner, M.P., Kohut, M.L. 2006. Year-and-a-half old, dried Echinacea spp. roots retain cytokine-modulating capabilities in an in vitro human older adult model of influenza vaccination. *Planta Medica*. 72:1207-1215.

3625-21000-054-00D

IDENTIFICATION OF PLANT GENETIC MECHANISMS THAT INFLUENCE INSECT-POLLINATOR PREFERENCE – Reid Palmer (P) and Craig Abel; Ames, Iowa

- Frasch, R.M., Weigand, C., Perez, P., Palmer, R.G., Sandhu, D. 2011. Molecular mapping of two environmentally sensitive male-sterile mutants in soybean. *Journal of Heredity*. 102:11-16.
- Yang, N., Moon, J., Lee, Y., Lee, S., Kim, H., Hwang, C., Back, K., Palmer, R.G., Jeong, S. 2010. Genetic and sequence analysis of genes controlling natural variation of seed-coat and flower colors in soybean. *Journal of Heredity*. 101(6):757-768.
- Xu, M., Battacharyya, M.K., Palmer, R.G. 2010. Excision of an Active CACTA-Like Transposable Element from DFR2 Causes Variegated Flowers in Soybean [*Glycine max* (L.) Merr.]. *Genetics*. 184:53-63.
- Cervantes-Martinez, I., Xu, M., Ortiz-Perez, E., Kato, K.K., Horner, H.T., Palmer, R.G. 2009. The Male Sterility Locus ms3 is Present in a Fertility Controlling Gene Cluster in Soybean. *Journal of Heredity*. 10:565-570.
- Healy, R.A., Palmer, R.G., Horner, H.T. 2009. Multicellular Secretory Trichome Development on Soybean and Related *Glycine* Gynoecia. *International Journal of Plant Science*. 170(4):444-456.
- Palmer, R.G., Perez, P.T., Ortiz-Perez, E., Maalouf, F., Suso, M.J. 2009. The Role of Crop-Pollinator Relationships in Breeding for Pollinator-Friendly Legumes: From a Breeding Perspective. *Euphytica*. 170:35-52.
- Palmer, R.G., Doyle, J.J. 2009. Anthony H.D. Brown: Conservation Geneticist. *Plant Breeding Reviews*. 31:1-20.
- Perez, P.T., Cianzio, S.R., Palmer, R.G. 2009. Heterotic Patterns of Soybean Lines from 3-Way, 4-Way and 5-way Crosses, and Backcross Populations. *Journal of Crop Production*. 23:95-118.
- Perez, P.T., Cianzio, S.R., Palmer, R.G. 2009. Evaluation of Soybean [*Glycine max* (L.) Merr.] F1 Hybrids. *Journal of Crop Improvement*. 23:1-18.
- Palmer, R.G., Perez, P.T. 2008. Qualitative Genetics - Examples from Soybean and Other Crops. In: *Proceedings of the Conventional and Molecular Breeding of Field and Vegetable Crops International Conference*, November 24-27, 2008, Novi Sad, Serbia. p. 1-6.
- Palmer, R.G., Min, X. 2008. Positioning Three Qualitative Trait Loci on Soybean Molecular Linkage Group E. *Journal of Heredity*. 99:674-678.
- Palmer, R.G., Sandhu, D., Curran, K., Bhattacharyya, M.K. 2008. Molecular Mapping of 36 Soybean Male-Sterile, Female-Sterile Mutants. *Theoretical and Applied Genetics*. 117:711-719.
- Palmer, R.G., Zhang, L., Huang, Z., Xu, M. 2008. Allelism and Molecular Mapping of Soybean Necrotic Root Mutants. *Genome*. 51:243-250.
- Ortiz-Perez, E., Wiley, H., Horner, H.T., Davis, W.H., Palmer, R.G. 2007. Insect-mediated cross-pollination in soybean [*Glycine max* (L.) Merrill]: ii. phenotypic recurrent selection. *Euphytica* 152:1573-1595.

3625-21000-055-00D

GENETIC ANALYSIS OF SELECTION RESPONSE IN MAIZE POPULATIONS – Marvin Scott (P), Craig Abel, and Jode Edwards; Ames, Iowa

- Gutierrez-Rojas, A., Betran, J., Scott, M.P., Atta, H., Menz, M. 2010. Quantitative Trait Loci for Endosperm Modification and Amino Acid Contents in Quality Protein Maize. *Crop Science*. 50:870-879.
- Edwards, J.W. 2010. Testcross Response to Four Cycles of Half-sib and S2 Recurrent Selection in the BS13 Maize (*Zea mays* L.) Population. *Crop Science*. 50:1840-1847.
- Rohlfing, K.A., Pollak, L.M., White, P.J. 2010. Exotic Corn Lines with Increased Starch and Impact on Starch Thermal Characteristics. *Cereal Chemistry*. 87:190-193.
- Scott, M.P., Peterson, J.M., Hallauer, A.R. 2010. Evaluation of Combining Ability and Grain Quality of Quality Protein Maize Derived from U.S. Public Inbred Lines. *Maydica*. 54:449-456.
- Paraman, I., Moeller, L., Scott, M.P., Wang, K., Glatz, C., Johnson, L. 2010. Utilizing protein-lean co-products from corn containing recombinant pharmaceutical proteins for ethanol production. *Journal of Agricultural and Food Chemistry*. 58(19):10419-10425.
- Hasjim, J., Srichuwong, S., Scott, M.P., Jane, J. 2009. Kernel Composition, Starch Structure, and Enzyme Digestibility of Opaque-2 Maize and Quality Protein Maize. *Journal of Agricultural and Food Chemistry*. 57:2049-2055.
- So, Y., Edwards, J.W. 2009. A Comparison of Mixed-Model Analyses of the Iowa Crop Performance Test for Corn. *Crop Science*. 49:1593-1601.
- Sheperd, C.T., Scott, M.P. 2009. Construction and Evaluation of a Maize Chimeric Promoter with Activity in Kernel Endosperm and Embryo. *Biotech and Applied Biochemistry*. 52:233-243.
- Wardyn, B.M., Edwards, J.W., Lamkey, K.R. 2009. Predicted Gains from Inbred-Progeny Selection Is Inferior to Half-sib Selection for Two Maize Populations. *Crop Science*. 49:443-450.

- Scott, M.P., Blanco, M.H. 2009. Evaluation of the Methionine Content of Maize (*Zea mays* L.) Germplasm in the Germplasm Enhancement of Maize Program. *Plant Genetic Resources*. 7:237-243.
- Edwards, J.W. 2008. Predicted Genetic Gain and Inbreeding Depression with General Inbreeding Levels in Selection Candidates and Offspring. *Crop Science*. 48:2086-2096.
- Gutierrez-Rojas, A., Scott, M.P., Leyva, O.R., Menz, M., Betran, J. 2008. Phenotypic Characterization of Quality Protein Maize (QPM) Endosperm Modification and Amino Acid Contents in a Segregating Recombinant Inbred Population. *Crop Science*. 48:1714-1722.
- Haney, L.J., Lamkey, K.R., Kirkpatrick, K., Coors, J.G., Lorenz, A.J., Raman, D.R., Anex, R.P., Scott, M.P. 2008. Development of a Fluorescence-based Method for Monitoring Glucose Catabolism and its Potential use in a Biomass Hydrolysis Assay. *Biotechnology for Biofuels*. 1:17.
- Lorenz, A., Scott, M.P., Lamkey, K.R. 2008. Genetic variance and breeding potential of phytate and inorganic phosphorus in a maize population. *Crop Science*. 48:79-84.
- Scott, M.P., Darrigues, A., Stahly, T., Lamkey, K. 2008. Recurrent Selection to Control Grain Methionine Content and Improve Nutritional Value of Maize. *Crop Science*. 48:1705-1713.
- Shepard, C.T., Vignaux, N., Peterson, J.M., Johnson, L.R., Scott, M.P. 2008. Green Fluorescent Protein as a Tissue Marker in Transgenic Maize Seed. *Cereal Chemistry*. 85:188-195.
- Sheperd, C.T., Vignaux, N., Peterson, J.M., Scott, M.P., Johnson, L.R. 2008. Dry-milling and Fractionation of Transgenic Maize Seed Tissues with Green Fluorescent Protein as a Tissue Marker. *Cereal Chemistry*. 85:196-201.
- Williams, C.L., Liebman, M., Edwards, J.W., James, D.E., Singer, J.W., Arritt, R.W., Herzmann, D. 2008. Predicting spatial variation of crop yield across a landscape using aggregated environmental data. *Crop Science*. 48:1545-1559.
- Chernyshova, A.A., White, P.J., Scott, M.P., Jannink, J. 2007. Selection for nutritional function and agronomic performance in oat. *Crop Science*. 47:2330-2339.
- Wardyn, B.M., Edwards, J.W., Lamkey, K.R. 2007. The genetic structure of a maize synthetic: the role of dominance. *Crop Science*. 47:467-476.
- Lorenz, A., Scott, M.P., Lamkey, K.R. 2007. Quantitative determination of phytate and available phosphorus for maize breeding. *Crop Science*. 47:600-604.

3625-21000-056-00D

BREEDING HIGH-QUALITY CORN FOR LOW-INPUT AND ORGANIC FARMING SYSTEMS –Craig Abel; Ames, Iowa

- Ali, F., Scott, M.P., Bakht, J., Chen, Y., Lubberstedt, T. 2010. Identification of Novel Brown-Midrib (bm) Genes in Maize by Tests of Allelism. *Plant Breeding*. doi:10.1111/j.1439-0523.2010.01791.x.
- Taboada-Gaytan, O., Pollak, L.M., Johnson, L., Fox, S.R., Montgomery, K.T. 2010. Variation among Physical, Compositional, and Wet-Milling Characteristics of the F1 Generation of Corn Hybrids of Introgressed Exotic and Adapted Inbred Lines. *Cereal Chemistry*. 87:175-181.
- Carena, M., Pollak, L.M., Salhuana, W., Denuc, M. 2009. Development of Unique and Novel Lines for Early-Maturing Hybrids: Moving GEM Germplasm Northward and Eastward. *Euphytica*. 170:87-97.
- Taboada-Gaytan, O., Pollak, L.M., Johnson, L., Fox, S. 2009. Wet-milling Characteristics of Ten Lines from the Germplasm Enhancement of Maize Project Compared to Five Corn Belt Lines. *Cereal Chemistry*. 86:204-209.
- De Geus, Y.N., Goggi, S., Pollak, L.M. 2008. Seed Quality of High Protein Corn Lines in Organic and Conventional Farming Systems. *Agronomy for Sustainable Development*. 28(4):541-550.
- Goggi, S., Caragea, P., Pollak, L.M., McAndrews, G., Devries, M., Montgomery, K. 2008. Seed quality assurance in maize breeding programs: tests to explain variations in corn inbreds and populations. *Agronomy Journal*. 100:337-343.
- Goggi, A., Pollak, L.M., Golden, J.C., Devries, M., McAndrews, G., Montgomery, K. 2007. Impact of early seed quality selection on maize inbreds and hybrids. *Maydica*. 52:223-233.

3625-21000-057-00D

FUNCTIONAL GENOMICS OF CEREAL DISEASE DEFENSE – Roger Wise (P), Craig Abel, and Nicholas Lauter; Ames, Iowa

- Moscou, M.J., Lauter, N.C., Steffenson, B., Wise, R.P. 2011. Quantitative and qualitative stem rust resistance factors in barley are associated with transcriptional suppression of defense regulons. *PLoS Genetics*. 7(7):DOI: 10.1371/journal.pgen.1002208.
- Moscou, M.J., Lauter, N.C., Caldo, R.A., Nettleton, D., Wise, R.P. 2011. Quantitative and temporal definition of the Mla transcriptional regulon during barley-powdery mildew interactions. *Molecular Plant-Microbe Interactions*. 24(6):694-705.

- Crossett, A., Lauter, N.C., Love, T.M. 2010. An Empirical Method for Establishing Positional Confidence Intervals Tailored for Composite Interval Mapping of QTL. *PLoS One*. 5:e9039.
- Abebe, T., Melmaiee, K., Berg, V., Wise, R.P. 2009. Drought Response in the Spikes of Barley: Gene Expression in the Lemma, Palea, Awn, and Seed. *Functional and Integrative Genomics*. 10(2):191-205.
- Zhou, S., Wei, F., Nguyen, J., Bechner, M., Potamouis, K., Goldstein, S., Pape, L., Mehan, M.R., Churas, C., Pasternak, S., Forrest, D.K., Wise, R.P., Ware, D., Wing, R., Waterman, M., Livny, M., Schwartz, D.C. 2009. A Single Molecule Scaffold for the Maize Genome. *PLoS Genetics*. 5(11):1-14.
- Kronmiller, B.A., Wise, R.P. 2009. Computational Finishing of Large Sequence Contigs Reveals Interspersed Nested Repeats and Gene Islands in the rf1-associated Region of Maize. *Plant Physiology*. doi: 10.1104. p. 109.143370. 151(2):483-495.
- Wang, K., Frame, B., Xu, X., Moeller, L., Lamkey, K., Wise, R.P. 2009. Strategies for the Production of Maize-derived Pharmaceuticals using Cytoplasmic Male Sterile Lines: in vitro Tissue Culture/Transformation and Field Breeding Approaches. *Maydica*. 54(1):199-210.

3625-21220-004-00D

FUNCTIONAL AND STRUCTURAL GENETIC ANALYSIS OF SOYBEAN AND OTHER LEGUMES – Randy Shoemaker (P), Craig Abel, and Michelle Graham; Ames, Iowa

- Du, J., Grant, D.M., Tian, Z., Nelson, R., Zhu, L., Shoemaker, R.C., Ma, J. 2010. SoyTEdb: A Comprehensive Database of Transposable Elements in the Soybean Genome. *Biomed Central (BMC) Genomics*. 11:113.
- Du, J., Tian, Z., Bowen, N., Schmutz, J., Shoemaker, R.C., Ma, J. 2010. Bifurcation and Enhancement of Autonomous-Non-Autonomous Retrotransposon Partnership through LTR Swapping in Soybean. *The Plant Cell*. 22:48-61.
- Joseph, B., Schlueter, J., Du, J., Graham, M.A., Ma, J., Shoemaker, R.C. 2009. Retrotransposons Within Syntenic Regions Between Soybean and Medicago Truncatula and Their Contribution to Local Genome Evolution. *The Plant Genome*. 2(3):211-223.
- Meyer, J.D., Silva, D.C., Yang, C., Van De Mortel, M., Pedley, K.F., Hill, J.H., Shoemaker, R.C., Abdelnoor, R., Whitham, S.A., Graham, M.A. 2009. Identification and Analyses of Candidate Genes for Rpp4 Mediated Resistance to Asian Soybean Rust in Soybean (Glycine max). *Plant Physiology*. 150:295-307.
- O'Rourke, J.A., Nelson, R., Grant, D.M., Schmutz, J., Grimwald, J., Cannon, S.B., Vance, C.P., Graham, M.A., Shoemaker, R.C. 2009. Integrating Microarray Analysis and the Soybean Genome to Understand the Soybean's Iron Deficiency Response. *Biomed Central(BMC)Genomics*. 10:376. Available: <http://www.biomedcentral.com/1471-2164/10/376>.
- Narayanan, N.N., Tasma, I.M., Grant, D.M., Shoemaker, R.C., Bhattacharyya, M.K. 2009. Identification of Candidate Signaling Genes Including Regulators of Chromosome Condensation 1 Protein Family Differentially Expressed in the Soybean - Phytophthora Sojae Interaction. *Journal of Theoretical and Applied Genetics*. 118(3):399-412.
- Innes, R.W., Ameline-Torregrosa, C., Ashfield, T., Cannon, E., Cannon, S.B., Chacko, B., Chen, N.W., Couloux, A., Dalvani, A., Denny, R., Deshpade, S., Doyle, J.J., Egan, A., Geffroy, V., Glover, N., Hans, C.S., Howell, S., Ilut, D., Jackson, S., Lai, H., Mammadov, J., Martin Del Campo, S., Metcalf, M., Nguyen, A., O'Brien, M., Pfeil, B., Podicheti, R., Ratnaparkhe, M., Roe, B., Saghai, M.A., Samain, S., Sanders, I., Segurens, B., Sevigna, M., Sherman-Broyles, S., Thareau, V., Tucker, D., Walling, J., Wawrzynski, A., Yi, J., Young, N.D. 2008. Differential Accumulation of Retroelements and Diversification of NB-LRR Disease Resistance Genes in Duplicated Regions Following Polyploidy in the Ancestor of Soybean. *Plant Physiology*. 148:1740-1759.
- Mathieu, M., Winters, E., Kong, F., Wan, J., Want, S., Eckert, H., Donovan, C., Somers, D., Wang, K., Nguyen, H., Shoemaker, R.C., Stacey, G., Clemente, T. 2008. Establishment of a Soybean (Glycine max L. Merr.) Transposon-based Mutagenesis Repository. *Planta*. 229:279-289.
- Graham, M.A., Silverstein, K.A., Vandenbosch, K.A. 2008. Defensin-like genes: genomic perspectives on a diverse superfamily in plants. *The Plant Genome/Crop Science*. 48:S3-S11.
- Van, K., Kim, D., Mei Cai, C., Young Kim, M., Shin, J., Graham, M.A., Shoemaker, R.C., Choi, B., Yang, T., Lee, S. 2008. Sequence Level Analysis of Recently Duplicated Regions in the Soybean [Glycine max (L.) Merr.] Genome. *DNA Research*. 15:93-102.
- Shoemaker, R.C., Schlueter, J.A., Jackson, S. 2008. Soybean genome structure and organization. In: Stacey, G. editor. *Genetics and Genomics of Soybean*. Series: Plant Genetics and Genomics: Crops and Models 2. New York, NY: Springer-Verlag. p. 125-140.
- O'Rourke, J., Graham, M.A., Vodkin, L., Gonzalez, D., Cianzio, S., Shoemaker, R.C. 2007. Recovering from iron deficiency chlorosis in near isogenic soybeans: a microarray study. *Plant Physiology and Biochemistry*. 45:287-292.
- O'Rourke, J.A., Charlson, D.V., Gonzalez, D.O., Vodkin, L.O., Graham, M.A., Cianzio, S.R., Grusak, M.A., Shoemaker, R.C. 2007. Microarray analysis of iron deficiency chlorosis in near-isogenic soybean lines. *Biomed Central (BMC) Genomics*. 8:476.

- O'Rourke, J.A., Graham, M.A., Cianzio, S.R., Shoemaker, R.C. 2007. Recovering from iron deficiency chlorosis in near isogenic soybeans: a microarray study [abstract]. Iron Nutrition and Interactions in Plants Symposium. Paper No. S6-P-21.
- Schlueter, J., Lin, J., Schlueter, S., Vasylenko-Sanders, I., Deshpandem, S., Yi, J., O'Bleness, M., Roe, B., Nelson, R., Scheffler, B.E., Jackson, S., Shoemaker, R.C. 2007. Gene duplication and paleopolyploidy in soybean and the implications for whole genome sequencing. *Biomed Central (BMC) Genomics*. 8:330.
- Silverstein, K.A., Moskal, Jr, W.A., Wu, H.C., Underwood, B.A., Graham, M.A., Town, C.D., Vandenbosch, K.A. 2007. Small cysteine-rich peptides resembling antimicrobial peptides have been under-predicted in plants. *Plant Journal*. 51:262-280.
- Van De Mortel, M., Recknor, J.C., Graham, M.A., Nettleton, D., Dittman, J., Nelson, R., Godoy, C.V., Abdelnoor, R.V., Almeida, A.M., Baum, T.J., Whitham, S.A. 2007. Biphasic responses of resistant and susceptible soybean plants to soybean rust, *Phakopsora pachyrhizi*. *Molecular Plant-Microbe Interactions*. 20:887-899.
- Schlueter, J.A., Sanders, I.F., Deshpande, S., Yi, J., Seigfried, M., Roe, B.A., Schlueter, S.D., Scheffler, B.E., Shoemaker, R.C. 2006. The fad2 gene family of soybean: insights into the structural and functional divergence of a paleopolyploid genome. *The Plant Genome*. 1:14-26.
- Graham, M.A., Ramirez, M., Valdes-Lopez, O., Lara, M., Tesfaye, M., Vance, C.P., Hernandez, G. 2006. Identification of candidate phosphorus stress induced genes in *Phaseolus vulgaris* L. through clustering analysis across several plant species. *Functional Plant Biology*. 33(8):789-797.
- Jackson, S.A., Rokhsar, D., Stacy, G., Shoemaker, R.C., Schmutz, J., Grimwood, J. 2006. Toward reference sequence of the soybean genome: a multi-agency effort. *The Plant Genome*. 1:55-61.

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BREEDING SELECTION AND MOLECULAR CHARACTERIZATION FOR IMPROVED SUGAR BEET GERMLASM – Jon McGrath (P), Renfu Lu, and Linda Hanson; East Lansing, Michigan

- Hanson, L.E. 2010. Interaction of *Rhizoctonia solani* and *Rhizopus stolonifer* Causing Root Rot of Sugar Beet. *Plant Disease*. 94(5):504-509.
- Hanson, L.E., Duckert, T.M., Goodwill, T.R., McGrath, J.M. 2010. Beta PIs From the USDA-ARS NPGS Evaluated for Resistance to *Cercospora beticola*, 2009. Plant Disease Management Reports. Available: <http://www.plantmanagementnetwork.org/pub/trial/PDMR/reports/2010/FC005.pdf>.
- El-Zohairy, El-Awady, A., Eissa, H.F., El-Khishin, D.A., Nassar, A., McGrath, J.M. 2009. Differential Expression of Salt Stress-related Genes in Wild *Beta vulgaris*. *Egyptian Journal of Genetics and Cytology*. 38:187-206.
- Hanson, L.E., Duckert, T.M., McGrath, J.M. 2009. Beta PIs from the USDA-ARS NPGS evaluated for resistance to *Cercospora beticola*, 2008. Plant Disease Management Reports. The American Phytopathological Society. 3:V017. DOI: 10.1094/PDMR01. Available: <http://www.plantmanagementnetwork.org/pub/trial/PDMR/reports/2009/V017.pdf>.
- Saccomani, M., Stevanato, P., Trebbi, D., McGrath, J.M., Biancardi, E. 2009. Molecular and Morpho-Physiological Characterization of Sea, Ruderal and Cultivated Beets. *Euphytica*. 169(1):19-29.
- Trebbi, D., McGrath, J.M. 2009. Functional differentiation of the sugar beet root system as indicator of developmental phase change. *Physiologia Plantarum*. 135(1):84-97.
- Hanson, L.E., Duckert, T.M., McGrath, J.M. 2009. Beta PIs from the USDA-ARS NPGS evaluated for resistance to *Cercospora beticola*, 2008. Plant Disease Management Reports. The American Phytopathological Society. 3:V017. DOI: 10.1094/PDMR01. Available: <http://www.plantmanagementnetwork.org/pub/trial/PDMR/reports/2009/V017.pdf>.
- Nagendran, S., Hammerschmidt, R., McGrath, J.M. 2009. Identification of Sugar Beet Germplasm EL51 as a Source of Resistance to Post-Emergence *Rhizoctonia* Damping-Off. *European Journal of Plant Pathology*. 123:461-471.
- Crowell, E.F., McGrath, J.M., Douches, D. 2008. Accumulation of vitamin E in potato (*Solanum tuberosum*) tubers. *Transgenic Research*. 17(2):205-217.
- McGrath, J.M., Elawady, A., El-Khishin, D., Naegel, R.P., Carr, K.M., De Los Reyes, B. 2008. Sugar Beet Germination: Phenotypic Selection and Molecular Profiling to Identify Genes Involved in Abiotic Stress Response. *Acta Horticulturae*. 782:35-50.
- McGrath, J.M., Trebbi, D., Fenwick, A.L., Panella, L.W., Schulz, B., Laurent, V., Barnes, S., Murray, S.C. 2007. An open-source first-generation molecular genetic map from a sugar x table beet cross and its extension to physical mapping. *The Plant Genome*. 47(1):S27-S44.
- Dale, T.M., McGrath, J.M., Renner, K.A. 2006. Response of sugarbeet (*Beta vulgaris*) varieties and populations to postemergence herbicides. *Journal of Sugarbeet Research*. p.119-126.
- McGrath, J.M. 2006. Registration of EL53 sugarbeet germplasm with smooth-root and moderate resistance to *rhizoctonia* crown and root rot. *Crop Science*. 46(5):2334-2335.

3640-21000-025-00D

GENOMICS, GERMLASM EVALUATION, AND GENETIC IMPROVEMENT OF OATS – St. Paul, Minnesota

- Rines, H.W., Phillips, R.L., Kynast, R.G., Okagaki, R.J., Galatowitsch, M.W., Huettl, P.A., Stec, A.O., Jacobs, M.S., Suresh, J., Porter, H.L., Walch, M.D., Cabral, C.B. 2009. Addition of Individual Chromosomes of Maize Inbreds B73 and Mo17 to Oat Cultivars Starter and Sun II: Maize Chromosome Retention, Transmission, and Plant Phenotype. *Theoretical and Applied Genetics*. 119(7):1255-1264.
- Okagaki, R.J., Jacobs, M.S., Stec, A.O., Kynast, R.G., Buescher, E., Rines, H.W., Vales, M., Riera-Lizarazu, O., Schneerman, M., Doyle, G., Friedman, K.L., Staub, R.W., Weber, D.F., Kamps, T.L., Amarillo, I.F., Chase, C.D., Bass, H.W., Phillips, R.L. 2008. Maize centromere mapping: A comparison of physical and genetic strategies. *Journal of Heredity*. 99:85-93.
- Rines, H.W., Porter, H.L., Carson, M.L., Ochocki, G.E. 2007. Introgression of crown rust resistance from diploid oat *Avena strigosa* into hexaploid cultivated oat *A. sativa* by two methods: direct crosses and through an initial 2x+4x synthetic hexaploid. *Euphytica*. 158:67-79.

3640-21000-027-00D

IMPROVING RUST AND FHB RESISTANCE IN HARD RED SPRING WHEAT THROUGH GENETICS AND GENOMICS – David Garvin (P) and Carroll Vance; St. Paul, Minnesota

- Garvin, D.F., Hareland, G.A., Gregoire, B.R., Finley, J.W. 2011. Impact of wheat grain selenium content variation on milling and bread baking. *Cereal Chemistry*. 88(2). 195-200.
- Garvin, D.F., McKenzie, N., Vogel, J.P., Mockler, T.C., Blankenheim, Z., Wright, J., Huo, N., Cheema, J.J., Dicks, J., Hayden, D.M., Gu, Y.Q., Tobias, C.M., Chang, J.H., Chu, A., Trick, M., Michael, T.P., Bevan, M.W., Snape, J.W. 2010. An SSR-Based Genetic Linkage Map of the Model Grass *Brachypodium distachyon*. *Genome*. 53(1):1-13.
- Garvin, D.F., Stack, R.W., Hansen, J.M. 2009. Quantitative Trait Locus Mapping of Increased Fusarium Head Blight Susceptibility Associated with a Wild Emmer Wheat Chromosome. *Phytopathology*. 99(4):447-452.
- Bolton, M.D., Kolmer, J.A., Garvin, D.F. 2008. Wheat Leaf Rust Caused by *Puccinia triticina*. *Molecular Plant Pathology*. 9(5):563-575.
- Bolton, M.D., Kolmer, J.A., Xu, W., Garvin, D.F. 2008. Lr34-Mediated Leaf Rust Resistance in Wheat: Transcript Profiling Reveals a High Energetic Demand Supported by Transient Recruitment of Multiple Metabolic Pathways. *Molecular Plant-Microbe Interactions*. 21(12):1515-1527.
- Garvin, D.F., Gu, Y.Q., Hasterok, R., Hazen, S.P., Jenkins, G., Mockler, T.C., Mur, L.A., Vogel, J.P. 2008. Development of genetic and genomic research resources for *Brachypodium distachyon*, a new model system for grass crop research. *The Plant Genome*. 48:S69-S84.
- Bilgic, H., Cho, S., Garvin, D.F., Muehlbauer, G.J. 2007. Mapping barley genes to chromosome arms by transcript profiling of wheat-barley ditelosomic chromosome addition lines. *Genome*. 50:898-906.
- Garvin, D.F. 2007. *Brachypodium*: a new monocot model plant system emerges. *Journal of the Science of Food and Agriculture*. 87:1177-1179.
- Van Eck, J., Conlin, B., Garvin, D.F., Mason, H., Navarre, D.A., Brown, C.R. 2007. Enhancing beta-carotene content in potato by RNAi-mediated silencing of the b-carotene hydroxylase gene. *American Journal of Potato Research*. 84:331-342.

3640-21000-028-00D

FUNCTIONAL GENOMICS FOR IMPROVING NUTRIENTS AND QUALITY IN ALFALFA AND SOYBEAN – Carroll Vance (P); St. Paul, Minnesota

- Yamagishi, M., Zhou, K., Osaki, M., Miller, S.S., Vance, C.P. 2011. Real-time RT-PCR profiling of transcription factors including 34 MYBs and signaling components in white lupin reveals their P status dependent and organ-specific expression. *Plant and Soil Journal*. 342(1-2):481-493.
- Bolon, Y.E., Haun, W.J., Xu, W.W., Grant, D.M., Stacey, M.G., Nelson, R., Gerhardt, D.J., Jeddeloh, J.A., Stacey, G., Muehlbauer, G.J., Orf, J.H., Naeve, S.L., Stupar, R.M., Vance, C.P. 2011. Phenotypic and genomic analyses of a fast neutron mutant population resource in soybean. *Plant Physiology*. 156(1):240-253.
- Woody, J.L., Severin, A., Bolon, Y.E., Joseph, B., Diers, B.W., Farmer, A.T., Weeks, N.T., Muehlbauer, G.J., Nelson, R., Grant, D.M., Specht, J.E., Graham, M.A., Cannon, S.B., May, G.D., Vance, C.P., Shoemaker, R.C. 2011. Gene expression patterns are correlated with genomic and genic structure in soybean. *Genome*. 54(1):10-18.
- Vance, C.P. 2010. Quantitative trait loci, epigenetics, sugars, and microRNAs: quaternaries in phosphate acquisition and use. *Plant Physiology*. 154(2):582-588.

- Valdes-Lopez, O., Yang, S.H., Aparicio-Fabre, R., Graham, P.H., Reyes, J., Vance, C.P., Hernandez, G. 2010. MicroRNA Expression Profile in Common Bean (*Phaseolus vulgaris*) under Nutrient Deficiency Stresses and Manganese Toxicity. *New Phytologist*. 187(3):805-818.
- Wang, B.L., Tang, X.Y., Cheng, L.Y., Zhang, A.Z., Zhang, W.H., Zhang, F.S., Liu, J.Q., Cao, Y., Allan, D.L., Vance, C.P., Shen, J.B. 2010. Nitric Oxide is Involved in Phosphorus Deficiency-Induced Cluster-Root Development and Citrate Exudation in White Lupin. *New Phytologist*. 187(4):1112-1123.
- Severin, A.J., Woody, J.L., Bolon, Y.E., Joseph, B., Diers, B.W., Farmer, A.D., Muehlbauer, G.J., Nelson, R., Grant, D.M., Specht, J.E., Graham, M.A., Cannon, S.B., May, G.D., Vance, C.P., Shoemaker, R.C. 2010. RNA-Seq Atlas of Glycine max: A Guide to the Soybean Transcriptome. *Biomed Central (BMC) Plant Biology*. 10:610.
- Valdes-Lopez, O., Yang, S.H., Graham, P.H., Reyes, J.L., Vance, C.P., Hernandez, G. 2010. Identification of Nutritional Stress-Responsive miRNAs in *Phaseolus vulgaris* [abstract]. XVIII Plant and Animal Genome Conference Proceedings, January 9-13, 2010, San Diego, California. Abstract W367.
- Liu, J., Allan, D.L., Vance, C.P. 2010. Systemic Signaling and Local Sensing of Phosphate in Common Bean: Cross-Talk Between Photosynthate and MicroRNA399. *Molecular Plant*. 3(2):428-437.
- Bolon, Y.E., Joseph, B., Cannon, S.B., Graham, M.A., Diers, B.W., Farmer, A.D., May, G.D., Muehlbauer, G.J., Specht, J.E., Tu, Z., Weeks, N.T., Xu, W.W., Shoemaker, R.C., Vance, C.P. 2010. Complementary Genetic and Genomic Approaches Help Characterize the Linkage Group I Seed Protein QTL in Soybean. *Biomed Central (BMC) Plant Biology*. 10:41. Available: <http://www.biomedcentral.com/1471-2229/10/41>.
- Sbabou, L., Bucciarelli, B., Miller, S.S., Liu, J., Brhada, F., Filali-Maltouf, A., Allan, D., Vance, C.P. 2010. Molecular Analysis of SCARECROW Genes Expressed in White Lupin Cluster Roots. *Journal of Experimental Botany*. 61(5):1351-1363.
- Hernandez, G., Valdes-Lopez, O., Ramirez, M., Goffard, N., Weiller, G., Aparicio-Fabre, R., Fuentes, S., Erban, A., Kopka, J., Udvardi, M.K., Vance, C.P. 2009. Global changes in the transcript and metabolic profiles during symbiotic nitrogen fixation in phosphorus-stressed common bean plants. *Plant Physiology*. 151(3):1221-1238.
- Kuppusamy, K.T., Ivashuta, S., Bucciarelli, B., Vance, C.P., Gantt, S.J., VandenBosch, K.A. 2009. Knockdown of Cell Division Cycle16 Reveals an Inverse Relationship between Lateral Root and Nodule Numbers and a Link to Auxin in *Medicago truncatula*. *Plant Physiology*. 151(3):1155-1166.

3655-21000-048-00D

ALLIUM, CUCUMIS, AND DAUCUS GERMLASM ENHANCEMENT, GENETICS, AND BIOCHEMISTRY – Philipp Simon (P), Yiqun Weng, and Michael Havey; Madison, Wisconsin

- Miao, H., Zhang, S., Wang, X., Zhang, Z., Li, M., Mu, S., Cheng, Z., Zhang, R., Huang, S., Xie, B., Fang, Z., Zhang, Z., Weng, Y., Gu, X. 2011. A Linkage Map of Cultivated Cucumber (*Cucumis Sativus* L.) with 248 Microsatellite Marker Loci and Seven Genes for Horticulturally Important Traits. *Euphytica*. Available: <http://www.springerlink.com/content/l8u2603qg4657g53/fulltext.html>.
- Weng, Y., Perumal, A., Rudd, J.C., Burd, J.D. 2010. Biotypic Diversity in Greenbug (Hemiptera: Aphididae): Microsatellite Marker-Based Regional Divergence and Host-Adapted Differentiation. *Journal of Economic Entomology*. 103(4):1454-1463.
- Weng, Y., Staub, J.E., Johnson, S., Huang, S. 2010. An Extended Intervarietal Microsatellite Linkage Map of Cucumber, *Cucumis Sativus* L. *HortScience*. 45:882-886.
- Cavagnari, P.F., Weng, Y., Senalik, D.A., Yang, L., Simon, P.W., Harkins, T.T., Kodira, C.D., Huang, S. 2010. Genome-wide Characterization of Simple Sequence Repeats in Cucumber (*Cucumis Sativus* L.). *Biomed Central (BMC) Genomics*. Available: <http://www.biomedcentral.com/1471-2164/11/569>.
- Cuevas, H.E., Song, H., Staub, J.E., Simon, P.W. 2010. Inheritance of Beta-Carotene-Associated Flesh Color in Cucumber (*Cucumis Sativus* L.) Fruit. *Euphytica*. 171(3):301-311.
- Cuevas, H.E., Staub, J.E., Simon, P.W. 2010. Inheritance of the Quantity of β -carotene and Fruit Maturity of Melon (*Cucumis Melon* L.). *Euphytica*. 171(3):301-311.
- Garcia Lampasona, S., Burba, J.L., Simon, P.W. 2010. Molecular Markers: Are They Really Useful to Detect Genetic Variability in Local Garlic Collections? *The Americas Journal of Plant Science and Biotechnology*. 4(S1):104-112.
- Jakse, M., Hirscheegger, P., Bohanec, B., Havey, M.J. 2010. Evaluation of Gynogenic Responsiveness and Pollen Viability of Selfed Doubled Haploid Onion Lines and Chromosome Doubling via Somatic Regeneration. *Journal of the American Society for Horticultural Science*. 135(1):67-73.
- Kang, H., Yang, Y., Zhang, Z., Zhang, S., Mao, Z., Cheng, G., Gu, X., Huang, S., Weng, Y., Xie, B. 2010. Fine Genetic Mapping Localizes Cucumber Scab Resistance Gene *Ccu* into an R Gene Cluster. *Journal of Theoretical and Applied Genetics*. Available: <http://www.springerlink.com/content/lqu520225154x357/>.
- Melgar, S., De Biologia, E., Havey, M.J. 2010. The Dominant Ms Allele in Onion Shows Reduced Penetrance. *Journal of the American Society for Horticultural Science*. 135:49-52.

- Simkova, H., Safar, J., Kubalakova, M., Suchankova, P., Cihalikova, J., Robert-Quatre, H., Azhaguvel, P., Weng, Y., Peng, J.H., Lapitan, N.L., Ma, Y., You, F.M., Luo, M., Bartos, J., Dolezel, J. 2010. BAC Libraries from Wheat Chromosome 7D – Efficient Tool for Positional Cloning of Aphid Resistance Genes. *Journal of Biomedicine and Biotechnology*. Available: <http://www.hindawi.com/journals/jbb/2011/302543.html>.
- Vittal, R., Stewart, B.A., Weng, Y., Ghosh, N. 2010. Genetic Diversity among Sorghum Bicolor L. Moench Benotypes as Revealed by Prolamines and SSR Markers. *Journal of Biotechnology*. 2:101-111.
- Weng, Y. 2010. Low Genetic Diversity Among USDA Cucumis Metuliferus Collections Revealed by Cucumber Genomic Microsatellite Markers. *HortScience*. 45(2):214-219.
- Bartoszewski, G., Gawronski, P., Szklarczyk, M., Verbakel, H., Havey, M.J. 2009. A One-Megabase Physical Map Provides Insights on Gene Organization in the Enormous Mitochondrial Genome of Cucumber. *Genome*. 52:299-307.
- Havey, M.J., Raines, S., Henson, C.A. 2009. Genetic Analyses of Carbohydrate Accumulation in Onion. *Journal of the American Society for Horticultural Science*. 134:618-623.
- Huang, S., Li, R., Zhang, Z., Li, L., Gu, X., Fan, W., Lucas, W., Wang, X., Xie, B., Ni, P., Ren, Y., Zhu, H., Li, J., Lin, K., Jin, W., Fei, Z., Li, G., Staub, J.E., Kilian, A., Van Der Vossen, E.A., Wu, Y., Guo, J., He, J., Jia, Z., Ren, Y., Tian, G., Lu, Y., Ruan, J., Qian, W., Wang, M., Huang, Q., Li, B., Xuan, Z., Cao, J., Wu, Z., Zhang, J., Cai, Q., Bai, Y., Zhao, B., Han, Y., Li, Y., Li, X., Wang, S., Shi, Q., Liu, S., Cho, W.K., Kim, J., Xu, Y., Heller-Uszynska, K., Miao, H., Cheng, Z., Zhang, S., Wu, J., Yang, Y., Kang, H., Li, M., Liang, H., Ren, X., Shi, Z., Wen, M., Jian, M., Yang, H., Zhang, G., Yang, Z., Chen, R., Liu, S., Li, J., Ma, L., Liu, H., Zhou, Y., Zhao, J., Fang, X., Li, G., Fang, L., Li, Y., Liu, D., Zheng, H., Zhang, Y., Qin, N., Li, Z., Yang, G., Yang, S., Bolund, L., Kristiansen, K., Zheng, H., Li, S., Zhang, X., Yang, H., Wang, J., Sun, R., Zhang, B., Jiang, S., Wang, J., Du, Y., Li, S. 2009. The Genome of the Cucumber, *Cucumis Sativus* L. *Nature Genetics*. 41(12):1275-1281.
- Ortiz, R., Simon, P.W., Jansky, S.H., Stelly, D. 2009. Ploidy Manipulation of the Gametophyte, Endosperm, and Sporophyte in Nature and for Crop Improvement – A Tribute to Prof. Stanley J. Peloquin (1921-2008). *Annals Of Botany*. 104(5):795-807.
- Simon, P.W., Cavagnaro, P.F., Chung, S., Szklarczyk, M., Grzebelus, D., Senalik, D.A., Atkins, A.E. 2009. Characterization of a deep-coverage carrot (*Daucus carota* L.) BAC library and initial analysis of BAC-end sequences. *Molecular Genetics and Genomics*. 281(3):273-288.
- Simon, P.W., Cavagnaro, P.F., Senalik, D.A. 2009. SplinkBES - A Splinkerette-Based Method for Generating Long End Sequences From Large Insert DNA Libraries. *Biotechniques*. 47:681-690.
- Sun, T., Simon, P.W., Tanumihardjo, S.A. 2009. Antioxidants and Antioxidant Capacity of Biofortified Carrots (*Daucus Carota*, L.) of Various Colors. *Journal of Agricultural and Food Chemistry*. 57(10):4142-4147.
- Weng, Y., Lu, H., Rudd, J.C., Burd, J.D. 2009. Molecular Mapping of Greenbug Resistance Genes Gb2 and Gb6 in 1AL.1RS Wheat-Rye Translocations. *Plant Breeding*. Available: <http://www3.interscience.wiley.com/cgi-bin/fulltext/123224851/HTMLSTART>.
- Simon, P.W., Pollak, L.M., Clevidence, B.A., Holden, J.M., Haytowitz, D.B. 2009. Plant Breeding for Human Nutritional Quality. *Plant Breeding Reviews*. 31:325-415.
- Just, B.J., Santos, C.F., Yandell, B.S., Simon, P.W. 2009. Major QTL for Carrot Color are Associated with Carotenoid Biosynthetic Genes and Interact Epistatically in a Domesticated x Wild Carrot Cross. *Theoretical and Applied Genetics*. 119(7):1155-1169.
- Mills, J.P., Simon, P.W., Tanumihardjo, S.A. 2008. Biofortified Carrot Intake Enhances Liver Antioxidant Capacity and Vitamin A Status in Mongolian Gerbils. *Journal of Nutrition*. 138(9):1692-1698.
- Mccallum, J., Thomson, S., Pither-Joyce, M., Kenel, F., Clarke, A., Havey, M.J. 2008. Genetic Diversity Analysis and Single-nucleotide Polymorphism Marker Development in Cultivated Bulb Onion Based on Expressed Sequence Tag–Simple Sequence Repeat Markers. *Journal of the American Society for Horticultural Science*. 133:810-818.
- Iovene, M., Wielgus, S.M., Simon, P.W., Buell, C.R., Jiang, J. 2008. Chromatin structure and physical mapping of chromosome 6 of potato and comparative analyses with tomato. *Genetics*. 180(3):1307-1317.
- Ipek, M., Ipek, A., Simon, P.W. 2008. Genetic Characterization of *Allium Tuncelianum*: An Endemic Edible *Allium* Species With Garlic Odor. *Scientia Horticultureae*. 115(4):409-415.
- Ipek, M., Ipek, A., Simon, P.W. 2008. Molecular Characterization of Kastamonu Garlic: An Economically Important Garlic Clone in Turkey. *Scientia Horticultureae*. 115(2):203-208.
- Jakse, J., Meyer, J., Suzuki, G., Mccallum, J., Cheung, F., Town, C.D., Havey, M.J. 2008. Pilot Sequencing of Onion Genomic DNA Reveals Fragments of Transposable Elements, Low Gene Densities, and Significant Gene Enrichment After Methyl Filtration. *Molecular Genetics and Genomics*. 280(4):287-292.
- Camadro, E.L., Cauhepe, M.A., Simon, P.W. 2008. Compatibility Relations Between the Edible Carrot *Daucus Carota* and *D. Pusiillus*, a Related Wild Species from the Argentinian Pampas. *Euphytica*. 159(1-2):103-109.
- Al-Faifi, S., Meyer, J.D., Garcia-Mas, J., Monforte, A.J., Havey, M.J. 2008. Exploiting Synteny in *Cucumis* for Mapping of Psm, A Unique Locus Controlling Paternal Mitochondrial Sorting. *Theoretical and Applied Genetics*. 117(4):523-529.
- Behera, T., Singh, A.K., Staub, J.E. 2008. Genetic Diversity Analysis of Indian Bitter Gourd (*Momordica Charantia* L.) Allows for the Development of Crop Improvement Strategies. *Scientia Horticultureae*. 115:209-217.

- Gaikwad, A.B., Singh, A.K., Behera, T.K., Chandel, D., Staub, J.E. 2008. AFLP Analysis Provides Strategies for Improvement of *Momordica Charantia* L. (Bitter Gourd). *HortScience*. 43(1):127-133.
- Grzebelus, D., Simon, P.W. 2008. Diversity of DcMaster-like elements of the PIF/Harbinger superfamily in the carrot genome. *Genetica*. Available: <http://www.ncbi.nlm.nih.gov/pubmed/18535910>.
- Iovene, M., Grzebelus, E., Carputo, D., Jiang, J., Simon, P.W. 2008. Major Cytogenetic Landmarks and Karyotype analysis in Carrot and Other Apiaceae. *American Journal of Botany*. 95:793-804.
- Khar, A., Jakse, J., Havey, M.J. 2008. Segregations for Onion-Bulb Colors Reveal that Red is Controlled by at Least Three Loci. *Journal of the American Society for Horticultural Science*. 133:42-47.
- Luan, F., Dellanay, I., Staub, J.E. 2008. Melon (*Cucumis melo* L.) Diversity Analyses Provide Strategies for Germplasm Curation, Genetic Improvement and Evidentiary Support of Domestication Patterns. *Euphytica*. 164(2):445-461.
- Meyer, J.D., Deleu, W., Garcia-Mas, J., Havey, M.J. 2008. Construction of a Fosmid Library of Cucumber (*Cucumis sativus*) and Comparative Analyses of the eIF4E and eIF(iso)4E Regions from Cucumber and Melon (*Cucumis melo*). *Molecular Genetics and Genomics*. 279(5):473-480.
- Peloquin, S.J., Boiteux, L.S., Simon, P.W., Jansky, S.H. 2008. A chromosome-specific estimate of transmission of heterozygosity by 2n gametes in potato. *Journal of Heredity*. 99(2):177-181.
- Mills, J.P., Simon, P.W., Tanumihardjo, S.A. 2007. β -Carotene from Red Carrot Maintains Vitamin A Status, but Lycopene Bioavailability Is Lower Relative to Tomato Paste in Mongolian Gerbils. *Journal of Nutrition*. 137:1395-1400.
- Ipek, M., Ipek, A., Senalik, D.A., Simon, P.W. 2007. Characterization of an Unusual Cytoplasmic Chimera Detected in Bolting Garlic (*Allium sativum* L.) Clones. *Journal of the American Society for Horticultural Science*. 132:664-669.
- Grzebelus, D., Jagosz, B., Simon, P.W. 2007. The DcMaster Transposon Display maps polymorphic insertion sites in the carrot (*Daucus carota* L.) genome. *Gene*. 390:67-74.
- Zalapa, J., Staub, J.E., McCreight, J.D. 2007. Mapping and QTL analysis of plant architecture and fruit yield in melon. *Theoretical and Applied Genetics*. 114:1185-1201.
- Camadro, E.L., Cauhepe, M.A., Simon, P.W. 2007. Geographical Distribution of Wild *Daucus* Species in the Natural Grasslands of the Argentinian Pampas. *Genetic Resources and Crop Evolution*. 54(4):855-863.
- Cavagnaro, P.F., Camargo, A., Galmarini, C.R., Simon, P.W. 2007. Effect of cooking on garlic (*Allium sativum*) antiplatelet activity and thiosulfinates content. *Journal of Agricultural and Food Chemistry*. 55:1280-1288.
- Chen, J., Ren, G., Luo, X., Staub, J.E., Jahn, M. 2006. Inheritance of aspartate aminotransferase (AAT) in cucumis species as revealed by interspecific hybridization. *Canadian Journal of Botany*. 84:1503-1507.
- Chung, S., Gordon, V., Staub, J.E. 2006. Sequencing of cucumber (*Cucumis sativus* L.) chloroplast genomes identifies putative candidate genes for chilling tolerance. *Genome*. 50:215-225.
- Mccallum, J., Pither-Joyce, M., Shaw, M., Kenel, F., Davis, S., Butler, R., Scheffer, J., Jakse, J., Havey, M.J. 2006. Genetic Mapping of Sulfur Assimilation Genes Reveals a QTL for Onion Bulb Pungency. *Journal of Theoretical and Applied Genetics*. 114:815-822.
- Porter Dosti, M., Mills, J.P., Simon, P.W., Tanumihardjo, S.A. 2006. Bioavailability of beta-carotene (betaC) from purple carrots is the same as typical orange carrots while high-betaC carrots increase betaC stores in Mongolian gerbils (*Meriones unguiculatus*). *British Journal of Nutrition*. 96(2):260-269.
- Zhuang, F., Chen, J., Staub, J.E., Qian, C. 2006. Taxonomic relationships of a rare cucumis species (*C. hystrix* chakr.) and its interspecific hybrid with cucumber. *HortScience*. 41:571-574.

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POTATO GENETICS, CYTOGENETICS, DISEASE RESISTANCE, AND PRE-BREEDING UTILIZING WILD AND CULTIVATED SPECIES – Shelley Jansky (P), Philipp Simon, Michael Havey, Paul Bethke, and Dennis Halterman; Madison, Wisconsin

- Chen, Y., Halterman, D.A. 2011. Phenotypic characterization of potato late blight resistance mediated by the broad-spectrum resistance gene RB. *Phytopathology*. 101(2):263-270.
- Chung, Y.S., Jansky, S.H., Spooner, D.M. 2011. A Test of Taxonomic and Biogeographic Predictivity: Resistance to Soft Rot in Wild Relatives of Cultivated Potato. *Phytopathology*. 101(2):205-212.
- Jansky, S.H., Hamernik, A.J., Bethke, P.C. 2011. Germplasm Release: Tetraploid Clones with Resistance to Cold-induced Sweetening. *American Journal of Potato Research*. Available: <http://www.springerlink.com/content/120949/?Content+Status=Accepted>.
- Halterman, D.A., Chen, Y., Sopee, J., Berduo, J., Sanchez, A. 2010. Competitive Interaction Between *Phytophthora Infestans* Effectors Leads to Increased Aggressiveness on Plants Containing Broad-Spectrum Late Blight Resistance. *PLoS One*. *PLoS ONE* 5(5): e10536. Available: <http://www.plosone.org/article/info:doi%2F10.1371%2Fjournal.pone.0010536>.
- Shahbazi, H., Aminian, H., Sahebani, N., Halterman, D.A. 2010. Biochemical Evaluation of Resistance Responses of Potato to Different Isolates of *Alternaria Solani*. *Phytopathology*. 100(5):454-459.

- Simon, R., Conghua, X., Clausen, A., Jansky, S.H., Halterman, D.A., Connor, T., Symon, D., Spooner, D.M. 2010. Wild and Cultivated Potato (*Solanum* sect. *Petota*) Escaped and Persistent Outside of its Natural Range. *Journal of Invasive Plant Science and Management*. 3:286-293.
- Bhaskar, P.B., Wu, L., Bethke, P.C., Vaillancourt, B., Jansky, S.H., Busse, J.S., Buell, R., Hamernik, A.J., Jiang, J. 2010. Suppression of the Vacuolar Invertase Gene Prevents Cold-induced Sweetening in Potato. *Plant Physiology*. Available: <http://www.plantphysiol.org/cgi/reprint/pp.110.162545v1?maxtoshow=&hits=10&RESULTFORMAT=&author1=bethke&andorexacttitle=&andorexacttitleabs=&andorexactfulltext=&andsearchid=1&FIRSTINDEX=0&sortspec=relevance&resourcetype=HWCIT&eaf>
- Bethke, P.C., Busse, J.S. 2010. Vine-Kill Treatment and Harvest Date Have Persistent Effects on Tuber Physiology After Harvest. *American Journal of Potato Research*. 87(3):299-309.
- Jansky, S.H. 2010. Potato Flavor. *American Journal of Potato Research*. 87(2):209-217.
- Jansky, S.H., Miller, J.C. 2010. Evaluation of Verticillium Wilt Resistance in Russet Norkotah and Six Strain Selections. *American Journal of Potato Research*. Available: <http://www.springerlink.com/content/j1w51303k0576246/fulltext.pdf>.
- Mccann, L.C., Bethke, P.C., Simon, P.W. 2010. Extensive Variation in Fried Chip Color and Tuber Composition in Cold-Stored Tubers of Wild Potato (*Solanum*) Germplasm. *Journal of Agriculture and Food Chemistry*. 58(4):2368-2376.
- Rodriguez, F., Ghislain, M., Clausen, A., Jansky, S.H., Spooner, D.M. 2010. Hybrid Origins of Cultivated Potatoes. *Journal of Theoretical and Applied Genetics*. 121:1187-1198.
- Bethke, P.C., Sabba, R.P., Bussan, A.J. 2009. Tuber Water and Pressure Potentials Decrease and Sucrose Contents Increase in Response to Moderate Drought and Heat Stress. *American Journal of Potato Research*. 86(6):519-532.
- Halterman, D.A., Liu, Z. 2009. Analysis Of Proteins Differentially Accumulated During Potato Late Blight Resistance Mediated by the RB Resistance Gene. *Physiological and Molecular Plant Pathology*. 74(2):151-160.
- Hamernik, A.J., Hanneman Jr, R.E., Jansky, S.H. 2009. Introgression of Wild Species Germplasm with Resistance to Cold Sweetening into the Cultivated Potato. *Crop Science*. 49:529-542.
- Jansky, S.H. 2009. Verticillium Wilt Resistance in U.S. Potato Breeding Programs. *American Journal of Potato Research*. 86(6):504-512.
- Jansky, S.H., Hamernik, A.J. 2009. The Use of *Solanum verrucosum* as a Bridge Species. *Genetic Resources and Crop Evolution*. 56(8):1107-1116.
- Bae, J., Tai, G., Jansky, S.H. 2009. Selection for Verticillium Wilt Resistance in Potato Breeding Populations Derived from Four Cross Types. *Plant Breeding*. 128(3):290-294.
- Halterman, D.A., Zhenyu, L. 2009. Different Genetic Mechanisms Control Foliar and Tuber Resistance to *Phytophthora infestans* in Wild Potato *Solanum verrucosum*. *American Journal of Potato Research*. 86(6):476-480.
- Jansky, S.H., Liping, J., Kaiyun, X., Xie, C., Spooner, D.M. 2009. Potato Production and Breeding in China. *American Journal of Potato Research*. 52(1):57-65.
- Rosenthal, S., Jansky, S.H. 2008. Effect of production site and storage on antioxidant levels in potato (*Solanum tuberosum* L.) tubers. *Journal of Food Composition and Analysis*. 88(12):2087-2092.
- Bae, J., Jansky, S.H., Rouse, D.I. 2008. The potential for early generation selection to identify potato clones with resistance to Verticillium wilt. *Euphytica*. 164(2):385-393.
- Bae, J., Halterman, D.A., Jansky, S.H. 2008. Development of a Molecular Marker Associated with Verticillium Wilt Resistance in Diploid Interspecific Potato Hybrids. *Molecular Breeding*. 22(1):61-69.
- Bethke, P.C., Busse, J.S. 2008. Validation of a simple, colorimetric, microplate assay using amplex red for the determination of glucose and sucrose in potato tubers and other vegetables. *American Journal of Potato Research*. 85(6):414-421.
- Bethke, P.C., Jansky, S.H. 2008. The Effects of Boiling and Leaching on the Content of Potassium and Other Minerals in Potatoes. *Journal of Food Science*. 73(5):H80-H85.
- Halterman, D.A., Colton, L., Weigus, S., Jiang, J. 2008. Rate-Limiting Late Blight Resistance Conferred by the RB Gene in *Solanum tuberosum* Transgenic Lines Does Not Impact Yield. *Plant Disease*. 92:339-343.
- Jansky, S.H. 2008. Genotypic and environmental contributions to baked potato flavor. *American Journal of Potato Research*. 84(1):95-96.
- Jansky, S.H., Simon, R., Spooner, D.M. 2008. A Test of Taxonomic Predictivity: Resistance to Early Blight in Wild Relatives of Cultivated Potato. *Phytopathology*. 98(6):680-687.
- Atallah, Z.K., Bae, J.J., Jansky, S.H., Rouse, D.I., Stevenson, W.R. 2007. Multiplex real-time quantitative PCR methodology to assist in the breeding of potato lines with resistance to Verticillium wilt. *Phytopathology*. 97(7):865-872.
- Bae, J., Atallah, Z.K., Jansky, S.H., Rouse, D.I., Stevenson, W.R. 2007. Colonization dynamics and spatial progression of *Verticillium dahliae* in individual stems of two potato cultivars with differing responses to potato early dying. *Plant Disease*. 91(9):1137-1141.

- Bethke, P.C., Libourel, I., Aoyama, N., Chung, Y., Still, D.W., Jones, R.L. 2007. The Arabidopsis thaliana aleurone layer responds to nitric oxide, gibberellin, and abscisic acid and is sufficient and necessary for seed dormancy. *Plant Physiology*. 143:1173-1188.
- Pfaff, K., Jansky, S.H. 2007. Effect of time and storage temperature on survival of *Verticillium dahliae* microsclerotia in dried potato stem tissue. *American Journal of Potato Research*. 84:271-273.
- Speransky, A.S., Krinitsina, A.A., Poltronieri, P., Fasano, P., Cimaglia, F., Santino, A., Bogacheva, A.M., Shevelev, A.B., Halterman, D.A., Valueva, T.A. 2007. Kunitz-type protease inhibitors group B from *Solanum palustre*. *Plant Biotechnology Journal*. 2(11):1417-1424.
- Halterman, D.A., Liu, Z. 2006. Identification and Characterization of RB Orthologous Genes from the Late Blight Resistant Wild Potato Species *Solanum verrucosum*. *Physiological and Molecular Plant Pathology*. 69:230-239.

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SYSTEMATICS, GENETIC DIVERSITY ASSESSMENT, AND ACQUISITION OF POTATOES, CARROTS, AND THEIR RELATED WILD RELATIVES – David Spooner (P) And Philipp Simon; Madison, Wisconsin

- Ovchinnikova, A., Krilova, E., Gavrilenko, T., Smekalova, T., Dorofeev, V., Zhuk, M., Knapp, S., Spooner, D.M. 2011. Taxonomy of Cultivated Potatoes (*Solanum* Section *Petota*: *Solanaceae*). *Botanical Journal of the Linnean Society*. 165:107-155.
- Ames, M., Spooner, D.M. 2010. Phylogeny of *Solanum* Series *Piurana* and Putatively Related Species in *Solanum* Section *Petota* Based on Five Conserved Ortholog Sequences. *Taxon*. 59(4):1091-1101.
- Cai, D., Zheng, X., Teng, Y., Spooner, D.M. 2010. Genetic Relationships Among *Pyrus pyrifolia* Cultivars from Southeastern China and Japan. *Acta Horticulturae*. 859:89-92.
- Gavrilenko, T., Antonova, O., Ovchinnikova, A., Novikova, L., Krilova, E., Mironenko, N., Pendinen, G., Smekalova, T., Islamshina, A., Shvachko, N., Kiru, S., Kostina, L., Afanasenko, O., Spooner, D.M. 2010. A Microsatellite and Morphological Assessment of the Russian National Potato Collection. *Genetic Resources and Crop Evolution*. 57:1151-1164.
- Luo, Q., Iovene, M., Spooner, D.M., Buell, C.R., Jiang, J. 2010. Evolution of Chromosome 6 of *Solanum* Species Revealed by Comparative Fluorescence in Situ Hybridization Mapping. *Chromosoma*. 119:435-442.
- Mione, T., Spooner, D.M. 2010. A New Species and Key to the *Jaltomata* (*Solanaceae*) of Mexico. *Novon*. 20(2):186-189.
- Spooner, D.M., Jansky, S.H., Gavrilenko, T., Ovchinnikova, A., Krilova, E., Simon, R. 2010. Ecogeography of Ploidy Variation in Cultivated Potato (*Solanum* Sect. *Petota*). *American Journal of Botany*. 97(12):2049-2060.
- Ghislain, M., Nunez, J., Del Rosario Herrera, M., Bonierbale, M., Spooner, D.M. 2009. The Single Andigenum Origin of Neo-Tuberosum Potato Materials is not Supported by Microsatellite and Plastid Marker Analyses. *Theoretical and Applied Genetics*. 118(5):963-969.
- Rodriguez, F., Ghislain, M., Spooner, D.M. 2009. Genomic Origins of the Cultivated Potato Species: GBSSI Sequencing Data. *The Plant Genome*. 48(S1):S27-S36.
- Rodriguez, F., Spooner, D.M. 2009. Nitrate Reductase Phylogeny of Potato (*Solanum* sect. *Petota*) Genomes with Emphasis on the Origins of the Polyploid Species. *Systematic Botany*. 34:207-219.
- Rodriguez, F., Tanksley, S., Wu, F., Spooner, D.M. 2009. Do potatoes and tomatoes have a single evolutionary history, and what proportion of the genome supports this history. *BMC Evolutionary Biology*. 9:191.
- Spooner, D.M., Clausen, A., Peralta, I. 2009. Taxonomic Treatment of *Solanum* Section *Petota* (Wild Potatoes) in *Catálogo de Plantas Vasculares del Cono Sur* (Argentina, Chile, Paraguay, Uruguay, y sur del Brasil). *Systematic Botany Monographs*. 107:3011-3053.
- Spooner, D.M., Jansky, S.H., Simon, R. 2009. Tests of Taxonomic and Biogeographic Predictivity: Resistance to Multiple Disease and Insect Pests in Wild Relatives of Cultivated Potato. *Crop Science*. 49:1367-1376.
- Ghislain, M., Nunez, J., Del Rosario Herrera, M., Pignataro, J., Guzman, F., Spooner, D.M. 2009. Robust and Highly Informative Microsatellite-Based Genetic Identity Kit for Potato. *Molecular Breeding*. 23:377-388.
- Spooner, D.M. 2009. DNA Barcoding Will Frequently Fail in Complicated Groups: An Example in Wild Potatoes. *American Journal of Botany*. 96:1177-1189.
- Alvarez, N., Peralta, I., Salas, A., Spooner, D.M. 2008. A morphometric study of species boundaries of the wild potato *Solanum brevicaulis* complex: replicated field trials in Argentina and Peru. *Plant Systematics and Evolution*. 274:37-45.
- Ames, M., Salas, A., Spooner, D.M. 2008. A morphometric study of species boundaries of the wild potato *Solanum* series *Piurana* (*solanaceae*) and putatively related species from seven other series in *Solanum* sect. *Petota*. *Systematic Botany*. 33:183-192.
- Ames, M., Spooner, D.M. 2008. DNA from herbarium specimens settle a long-held controversy about origins of the European potato. *American Journal of Botany*. 95:252-257.

- Fajardo, D., Salas, A., Spooner, D.M. 2008. A Replicated Morphological Study in Peru of *Solanum* Series Conicibaccata. *Systematic Botany*. 33(1):183-192.
- Jimenez, J., Brenes, A., Fajardo, D., Salas, A., Spooner, D.M. 2008. The Use and Limits of AFLP Data in the Taxonomy of Polyploid Wild Potato Species in *Solanum* Series Conicibaccata. *Conservation Genetics*. 9(2):381-387.
- Pendinen, G., Gavrilenko, T., Jiang, J., Spooner, D.M. 2008. Evidence of Allopolyploid Speciation of Wild Tetraploid Mexican Species of *Solanum* Series Longipedicellata Obtained by In Situ Hybridization. *Genome*. 51(9):714-720.
- Spooner, D.M., Rodriguez, F., Polgar, Z., Ballard, H., Jansky, S.H. 2008. Genomic origins of potato polyploids: GBSSI gene sequencing data. *The Plant Genome*. 48(S1):27-36.
- Spooner, D.M., Salas, A., Ames, M. 2008. Revision of the *Solanum* medians complex (*Solanum* section Petota). *Systematic Botany*. 33(3):579-588.
- Ames, M., Salas, A., Spooner, D.M. 2007. The discovery and taxonomic implications of a novel 41bp chloroplast DNA deletion in wild potatoes. *Plant Systematics and Evolution*. 268:159-175.
- Fajardo, D., Castillo, R., Salas, A., Spooner, D.M. 2007. A morphometric study of species boundaries of *Solanum* series Conicibaccata: a replicated field trial in Andean Peru. *Systematic Botany*. 33:183-192.
- Ghislain, M., Nunez, J., Trujillo, G., Herrera, M., Guzman, F., Spooner, D.M. 2007. Extensive SSR genotyping of potato landraces supports a drastic revaluation of their gene pool structure and classification. *Proceedings of the National Academy of Sciences*. 104:19398-19403.
- Knapp, S., Spooner, D.M., Leon, B. 2007. Solanaceae endémicas del Perú. *Revista Peruana de Biología*. 13:612-643.
- Rios, D., Ghislain, M., Rodriguez, F., Spooner, D.M. 2007. What is the origin of the European potato? Evidence from Canary Island landraces. *Crop Science*. 47:1271-1280.
- Spooner, D.M., Fajardo, D., Bryan, G. 2007. Species limits of *Solanum berthaultii* Hawkes and *S. tarijense* the implications for species boundaries in *Solanum* sect. Petota. *Taxon*. 56:987-999.
- Spooner, D.M., Jansky, S.H., Bussan, A.J. 2007. Experiences of the local organizing committee for a large scientific meeting. *Acta Horticulturae*. 745:513-532.

3655-21000-051-00D

CONSERVATION AND UTILIZATION OF POTATO GENETIC RESOURCES – John Bamberg (P), David Spooner, and Shelley Jansky; Sturgeon Bay, Wisconsin

- Bamberg, J.B. 2010. Tuber Dormancy Lasting Eight Years in the Wild Potato *Solanum Jamesii*. *American Journal of Potato Research*. 87(2):226-228.
- Bamberg, J.B., Del Rio, A.H., Fernandez, C.J., Salas, A., Vega, S.E., Zorilla, C., Roca, W., Tay, D. 2010. Comparison of "Remote" Versus "Easy" In Situ Collection Locations for USA Wild *Solanum* (Potato) Germplasm. *American Journal of Potato Research*. 87:277-284.
- Bamberg, J.B., Kiszonas, A.M. 2010. Survey of Tuber pH Variation in Potato (*Solanum*) Species. *American Journal of Potato Research*. 87(2):167-176.
- Nzaramba, M.N., Reddivari, L., Bamberg, J.B., Miller, J.C. 2009. Anti-Proliferative Activity and Cytotoxicity of *Solanum Jamesii* Tuber Extracts in Human Colon and Prostate Cancer Cells In Vitro. *Journal of Agriculture and Food Chemistry*. 57:8308-8315.
- Bamberg, J.B. Crazy sepal: a new floral sepallata-like mutant in the wild potato *Solanum microdontum* bitter. *American Journal of Potato Research*. 83:433-435.
- Bamberg, J.B., Del Rio, A.H. 2008. The canon of potato science. . . 50 topics in potato science that every potato scientist should know: 1) Genetic diversity and gene banks. *Potato Research*. 50:207-210.
- Hale, A.L., Reddivari, L., Bamberg, J.B., Miller, J.C. 2008. Interspecific variability for antioxidant activity and phenolic content among *solanum* species. *American Journal of Potato Research*. 85(5):332-341.
- Nzaramba, M., Ndambe, M., Bamberg, J.B., Miller, Jr., J.C. 2007. Effect of propagule type and growing environment on antioxidant activity and total phenolic content in potato germplasm. *American Journal of Potato Research*. 84:323-330.

3655-21000-052-00D

PHYSIOLOGICAL, BIOCHEMICAL AND GENETIC REGULATION OF CARBOHYDRATE METABOLISM IN CEREAL TISSUES – Cynthia Henson (P); Madison, Wisconsin

- Vinje, M.A., Willis, D.K., Henson, C.A., Duke, S.H. 2011. Differential expression of two β -amylase genes (*Bmy1* and *Bmy2*) in developing and mature barley grain. *Planta*. 233(5):1001-1010.

- Duke, S.H., Henson, C.A. 2011. Tracking the progress of congress mashing with osmolyte concentrations and malt extract values in North American barley cultivars and relationships between wort osmolyte concentrations, malt extract values and ASBC measures of malt quality. *Journal of American Society of Brewing Chemists*. 69(1):28-38.
- Vinje, M.A., Willis, D.K., Duke, S.H., Henson, C.A. 2010. Differential RNA expression of bmy1 during seed development and the association with beta-amylase accumulation, activity, and total protein. *Plant Physiology and Biochemistry*. 49:39-45.
- Filichkin, T.P., Vinje, M.A., Budde, A.D., Corey, A.E., Duke, S.H., Gallagher, L., Helgesson, J., Henson, C.A., Obert, D.E., Ohm, J.B., Petrie, S.E., Ross, A.S., Hayes, P.M. 2010. Phenotypic Variation for Diastatic power, β -Amylase Activity, and β -Amylase Thermostability vs. Allelic Variation at the Bmy1 Locus in a Sample of North American Barley Germplasm. *Crop Science*. 50:826-834.
- Duke, S.H., Henson, C.A. 2010. A Comparison of Barley Malt Osmolyte Concentrations and Standard Malt Quality Measurements as Indicators of Barley Malt Amylolytic Enzyme Activities. *Journal of American Society of Brewing Chemists*. 67(4):206-216.
- Vinje, M.A., Duke, S.H., Henson, C.A. 2010. Utilization of Different Bmy1 Intron III Alleles for Predicting β -Amylase Activity and Thermostability in Wild and Cultivated Barley. *Plant Molecular Biology Reporter*. 28:491-501.

3655-21000-053-00D

METABOLISM AND ANALYSIS OF CEREAL PHYTOCHEMICALS – Mitchell Wise (P) and Cynthia Henson; Madison, Wisconsin

- Wise, M.L. 2011. Effect of chemical systemic acquired resistance elicitors on avenanthramide biosynthesis in oat (*Avena sativa*). *Journal of Agriculture and Food Chemistry*. Available: <http://pubs.acs.org/doi/abs/10.1021/jf2008869>.
- Koenig, R.T., Dickman, J.R., Wise, M.L., Ji, L. 2011. Avenanthramides are bioavailable and accumulate in hepatic, cardiac, and skeletal muscle tissue following oral gavage in rats. *Journal of Agricultural and Food Chemistry*. <http://pubs.acs.org/doi/abs/10.1021/jf2002427>.
- Cai, S., Huang, C., Ji, B., Wise, M.L., Zhang, D., Yang, P. 2011. In vitro antioxidant activity and inhibitory effect, on oleic acid-induced hepatic steatosis, of fractions and subfractions from oat (*Avena sativa* L.) ethanol extract. *Journal of Food Science*. 124(3):900-905.

3655-21000-054-00D

GENETIC EXCHANGE AND GENE FLOW RISKS FROM PLANTS IN AGRICULTURE – Johanne Brunet (P) and Philipp Simon; Madison, Wisconsin

- Brunet, J., Stewart, C.M. 2010. The Impact of Distinct Insect Species on Pollination and Gene Flow in Alfalfa. *Psyche*. Available: <http://www.hindawi.com/journals/psyche/2010/201858.html>.
- Zalapa, J.E., Brunet, J. 2010. The Extent of Hybridization and Its Impact on the Genetic Diversity and Population Structure of an Invasive Tree, *Ulmus pumila* (Ulmaceae). *Evolutionary Applications*. 3(2):157-168.
- Brunet, J., Holmquist, K.G. 2009. The Influence of Distinct Insect Pollinators on Female and Male Reproductive Success in the Rocky Mountain Columbine. *Molecular Ecology*. 18(17):3745-3758.

5302-21220-005-00D

IMPROVEMENT OF PRUNUS AND VITIS SCIONS FOR FRUIT QUALITY AND PEST RESISTANCE – Craig Ledbetter (P) and David Ramming; Parlier, California

- Ramming, D.W., Gabler, F., Smilanick, J.L., Cadle Davidson, M., Barba, P., Consolieg, N.H., Mahanil, S., Cadle Davidson, L.E. 2011. A single dominant locus Ren4 confers non-race-specific penetration resistance to grapevine powdery mildew. *Phytopathology*. 101(4):502-508.
- Ramming, D.W. 2010. Greenhouse Screening of Grape Rootstock Populations to Determine Inheritance of Resistance to Phylloxera. *American Journal of Enology and Viticulture*. 61(2):234-239.
- Ledbetter, C.A. 2010. 'Bolaroja' and 'Primarosa': Two New Mid Season Apricots for the Fresh Market. *HortScience*. 45(3): 441-442.
- Ledbetter, C.A. 2010. Register of New Fruit and Nut Varieties List 45. *HortScience*. 45(5):716-756.
- Ledbetter, C.A., Sisterson, M.S. 2010. Carpological Variability of Almond (*Prunus dulcis* [Mill.] D.A. Webb cv Nonpareil) in a Single Orchard During Seven Consecutive Harvests. *HortScience*. 45(12):1788-1792.

- Ledbetter, C.A. 2010. Almond breeding and evaluation activities in Central California: Past and Future. *Options Mediterraneennes*. 94:255-260.
- Ledbetter, C.A. 2010. Apricot Breeding in North America: Current Status and Future Prospects. *Acta Horticulturae*. 862:85-91.
- Ledbetter, C.A. 2009. Using Central Asian germplasm to Improve Fruit Quality and Enhance Diversity in California Adapted Apricots. *Acta Horticulturae*. 814:77-80.
- Ramming, D.W. 2009. Water Loss from Fresh Berries of Raisin Cultivars under Controlled Drying Conditions. *American Journal of Enology and Viticulture*. 60:2:208-214.
- Ramming, D.W. 2008. 'Thomcord' grape. *HortScience* 43(3):945-946.
- Ledbetter, C.A. 2008. Register of New Fruit and Nut Varieties List 44. *HortScience* 43(5):1321-1343.
- Ledbetter, C.A. 2008. Shell Cracking Strength in Almond (*Prunus dulcis* [Mill.] D.A. Webb.) and its implication in uses as a value-added product. *Bioresource Technology*. 99(13): 5567-5573.
- Ledbetter, C.A., Sisterson, M.S. 2007. Advanced Generation Peach-Almond Hybrids as Seedling Rootstocks for Almond: First Year Growth and Potential Pollenizers for Hybrid Seed Production. *Euphytica*. 160(2): 259-266.
- Vizzotto, M., Cisneros-Zevallos, L., Byrne, D., Okie, W.R., Ramming, D.W. 2007. Large Variation Found in the Phytochemical and Antioxidant Activity of Peach and Plum Germplasm. *Journal of the American Society for Horticultural Science*. 132(3):334-340.
- Ledbetter, C.A., Sisterson, M.S. 2007. Advanced Generation Peach-Almond Hybrids as Seedling Rootstocks for Almond: First Year Growth and Potential Pollenizers for Hybrid Seed Production. *Euphytica*. 160(2): 259-266.
- Ramming, D.W. 2007. The USDA/ARS Grape Breeding Program at Parlier, CA. *Proceedings. First Annual National Viticulture Research Conference*, July 18-20, 2007, Davis, California. p. 73-74.

5305-21000-013-00D

SUGARBEET GERMLASM ENHANCEMENT, BREEDING, AND GENETICS – Kelley Richardson (P) and Frank Martin; Salinas, California

- Kongprakhon, P., Cuesta-Marcos, A., Hayes, P.M., Richardson, K.L., Sirithunya, P., Sato, K., Steffenson, B., and Toojinda, T. 2009. Validation of rice blast resistance genes in barley using a QTL mapping population and near-isolines. *Breeding Science*. 59: 341-349.
- Lewellen, R.T. 2007. Registration of CN927-202, CN926-11-3-22, and CN921-306 Sugarbeet Cyst Nematode Resistant Sugarbeet Lines. *Crop Science*. p. 167-169

5305-21220-004-00D

EVALUATION, ENHANCEMENT, GENETICS AND BREEDING OF LETTUCE, SPINACH, AND MELON – James McCreight (P), Bei quan Mou, Ivan Simko, and Ryan Hayes; Salinas, California

- Mou, B., Ryder, E.J. 2010. MU06-857, a Green Leaf Lettuce Breeding Line with Resistance to Leafminer and Lettuce Mosaic Virus. *HortScience*. 45: 666-667.
- Simko, I., Pechenick, D.A., Mchale, L.K., Truco, M.J., Ochoa, O.E., Michelmore, R.W., Scheffler, B.E. 2010. Development of molecular markers for marker-assisted selection in lettuce (*Lactuca sativa*). *Acta Horticulture Proceedings*, Year 2010, volume 859, pages 401-408.
- Simko, I., Hayes, R.J., Subbarao, K.V., Sideman, R.G. 2010. SM09A and SM09B: Romaine Lettuce Breeding Lines Resistant to Dieback and with Improved Shelf-life.. *HortScience*, Year 2010, volume 45, pages 670-672.
- Simko, I., Lincare, J.M. 2010. Combining partially ranked data in plant breeding and biology: II. Analysis with Rasch model.. *Communications in Biometry and Crop Science (CBCS)*. 2010, Vol 5, Pages 56-65.
- Simko, I., Pechenick, D.A. 2010. Combining partially ranked data in plant breeding and biology: I. Rank aggregating methods.. *Communications in Biometry and Crop Science (CBCS)*, Year 2010, volume 5, pages 41-55.
- Hayes, R.J., Wu, B., Pryor, B.M., Subbarao, K.V. Assessment of resistance in lettuce (*Lactuca sativa* L.) to mycelial and ascospore infection by *S.sclerotinia minor* Jagger and *S. sclerotiorum* (Lib.) de Bary. *HortScience* 45:333-341.
- Jenni, S., Hayes, R.J. 2010. Genetic variation, genotype x environment interaction, and selection for tipburn resistance in lettuce in multi-environments.. *Euphytica*, Feb 2010, Vol 171 (3), pages 427-439.
- Martin, F.N. (2009) *Pythium Genetics*. In: *Oomycete Genetics and Genomics: Diversity, Plant and Animal Interactions, and Toolbox*, Kurt Lamour and Sophien Kamoun eds. John Wiley & Sons, Hoboken, NJ. 574 pp.
- Sedlarova, M., Lebeda, A., Miksikova, P., Duchoslav, M., Sedlakova, B., McCreight, J.D. 2009. Histological Aspects of Cucumis melo PI 313970 Resistance to *Podosphaera xanthii* and *Golovinomyces cichoracearum*. *Journal of Plant Diseases and Protection* 116: 169-176.

- Simko, I., Pechenick, D.A., Mchale, L.K., Truco, M.J., Ochoa, O.E., Micheltmore, R.W., Scheffler, B.E. 2009. Association mapping and marker-assisted selection of the lettuce dieback resistance gene Tvr1.. Biomed Central (BMC) Plant Biology, Year 2009, volume 9, article number 135.
- Simko, I. 2009. Development of EST-SSR Markers for the Study of Population Structure in Lettuce (*Lactuca sativa* L.). Journal of Heredity 180: 256-262.
- Rauscher, G.M., Mayton, H., Bonierbale, M., Simko, I., Smart, C., Grunwald, N.J., Fry, W. Effect of Isolate, Environment and a Defeated R-gene (RPI-ber) on Quantitative Resistance of Potato to Late Blight. American Phytopathological Society Annual Meeting, Minneapolis, Minnesota, July 26-July 30. Phytopathology 98:S130.
- Paris, M.K., Zalapa, J.E., McCreight, J.D., Staub, J.E. 2008. Genetic Dissection of Fruit Quality Components in Melon (*Cucumis melo* L.) using a RIL Population Derived from Exotic x Elite US Western Shipping Germplasm. Molecular Breeding 22:405-419.
- Hayes, R.J., Liu, Y. 2008. Genetic Variation for Shelf-life of Salad-cut Lettuce in Modified-atmosphere Environments. Journal American Society Horticultural Science. 133(2):228-233.
- Mou, B. Evaluation of Oxalate Concentration in the U.S. Spinach Germplasm Collection. HortScience 43(6):1690-1693.2008.
- Mou, B., Koike, S.T., Du Toit, L.J. Screening for Resistance to Leaf Spot Diseases of Spinach.. HortScience 43(6):1706-1710.2008.
- Simko, I., Hu, J. Populations structure in cultivated lettuce (*Lactuca sativa* L.) Journal of the American Society for Horticultural Science, Vol. 133, Pages 61-68, 2008.
- Simko, I., Van Den Berg, J.H., Vreugdenhil, D., Ewing, E.E. 2008. Mapping loci for chlorosis associated with chlorophyll b deficiency in potato. Euphytica Vol 162, Pages 99-107, 2008.
- Hayes, R.J., Vallad, G., Qin, Q., Grube, R., Subbarao, K. 2007. Variation for Resistance to Verticillium Wilt in Lettuce (*Lactuca Sativa* L.). Plant Disease. 91(4):439-445.
- Mou, B. 2007. Leafminer-resistant spinach germplasm 03-04-63. HortScience. Vol. 42(7):1717-1718.
- Mou, B. 2007. Leafminer-resistant spinach germplasm 03-04-9. HortScience. 42:699-700.
- Mou, B., Hayes, R.J., Ryder, E.J. 2007. Crisphead breeding lines with resistances to corky root and lettuce mosaic. HortScience. 42:701-703.
- Rancher, G., Smart, C.D., Simko, I., Bonierbale, M., Mayton, H., Greenland, A., Fry, W.E. 2006. Characterization and mapping of rpi-ber, a novel potato late blight resistance gene from *solanum berthaultii*. Theoretical and Applied Genetics. 112(4):674-687.
- Vallad, G., Qin, Q., Hayes, R.J., Subbarao, K.V. 2006. The development and impact of verticillium wilt on post-harvest quality in commercial lettuce fields naturally infested with *V. dahliae*. Phytopathology.
- Vallad, G.E., Qin, Q., Grube, R., Hayes, R.J., Subbarao, K.V. 2006. Characterization of race-specific interactions among isolates of verticillium dahliae pathogenic on lettuce. Phytopathology. 96(12):1380-1387. DOI: 10.1094/PHYTO-96-1380.

5306-21000-017-00D

GENETIC DISSECTION OF SEEDLING COLD TOLERANCE IN RICE (*ORYZA SATIVA* L.) – Thomas Tai (P); Davis, California

- Kim, S.I. and Tai, T.H. 2011. Evaluation of seedling cold tolerance in rice cultivars: a comparison of visual ratings and quantitative indicators of physiological changes. Euphytica, 178:437-447
- Kim, S.I., Andaya, V.C., and Tai, T.H. 2011. Cold sensitivity in rice (*Oryza sativa* L.) is strongly correlated with a naturally occurring I99V mutation in the multifunctional glutathione transferase isoenzyme GSTZ2. Biochem J 435: 373-380
- Kim, S., Tai, T. 2010. Genetic analysis of two OsLpa1-like genes in Arabidopsis reveals that only one is required for wild-type seed phytic acid levels. Planta. 232:1241-1250.

5306-21000-018-00D

MANAGEMENT OF GENETIC RESOURCES FOR VITIS, PRUNUS, JUGLANS, FICUS, OLEA, PISTACIA, PUNICA, DIOSPYROS, ACTINIDIA, AND MORUS – John Preece (P) and Mallikarjuna Aradhya; Davis, California

- Aradhya, M.K., Woeste, K., Velasco, D. 2010. Genetic Diversity, Structure and Differentiation in Cultivated Walnut (*Juglans regia* L.). Acta Horticulturae. 861: 127-132.
- Gunn, B.F., Aradhya, M.K., Salick, J.M., Miller, A.J., Yongping, Y., Lin, L. 2010. Genetic variation in walnuts (*Juglans regia* and *J. sigillata*; Juglandaceae): Species distinctions, human impacts, and the conservation of agrobiodiversity in Yunnan, China1. American Journal of Botany. American Journal of Botany 97:660-671.

- Ibrahimovi, Z., Mcgranahan, G.H., Leslie, C.A., Aradhya, M.K. 2010. Genetic diversity in walnut (*Juglans regia*) from the caucasus nation of Azerbaijan. *Acta Horticulturae*. 861:163170.
- Soleri, D., Koehmstedt, A.M., Aradhya, M.K., Polito, V., Pinney, K. 2010. Comparing the historic olive trees (*Olea europaea* L.) of Santa Cruz with contemporaneous trees in the Santa Barbara, CA area: a case study of diversity and structure in an introduced agricultural species conserved in situ. *Genetic Resources and Crop Evolution*. *Genet. Resour. Crop Evol.* 57:973-984.
- Stover, E.W., Aradhya, M.K., Dangls, G., Prins, B.H., Cousins, P.S. 2009. GRAPE GENETIC RESOURCES AND RESEARCH AT THE DAVIS CALIFORNIA NATIONAL CLONAL GERMPLASM REPOSITORY. *Acta Horticulture Proceedings*. *Acta Hort(ISHS)*827:193-196.
- Bautista, J.C., Gerald, D., Judy, Y., Bruce, R., Stover, E.W. 2008. Use of Genetic Markers to Assess Pedigrees of Grape Cultivars and Breeding Program Selections. *American Journal of Enology and Viticulture*. 59:248-254.
- Stover, E.W. 2007. Mediterranean fruits: ancient history and modern promise. *HortScience*.
- Stover, E.W., Aradhya, M.K., Crisosto, C., Ferguson, L. 2007. The fig: overview of an ancient fruit. *HortScience*. Volume 42: pages 1083-1087
- Stover, E.W., Aradhya, M.K., Prins, B.H. 2007. Table Grape Genetic Resources at the Davis California National Clonal Germplasm Repository. *HortScience* 42:833-834.
- Stover, E.W., Mercure, E.W. 2007. The pomegranate: a new look at the fruit of paradise. *HortScience*. Volume 42: pages 1088-1092
- Yonemori, K., Honsho, C., Kitajima, A., Aradhya, M.K., Giordani, E., Bellini, E., Parfitt, D. 2007. Relationship of European Persimmon (*Diospyros kaki* Thunb.) Cultivars to Asian Cultivars, Characterized Using AFLPs.. *Genetic Resources and Crop Evolution*. 55: 81-89
- Aradhya, M.K., Potter, D., Simon, C.J. 2007. Origin, evolution, and biogeography of *Juglans*: a phylogenetic perspective. *Tree Genetics and Genomes*. 3: 363-378.
- Aradhya, M.K., Simon, C.J., Potter, D. 2006. Origin, evolution, and biogeography of *Juglans*: a phylogenetic perspective. *Acta Horticulture*. 705: 85-94.
- Arroyo-Garcia, R., Ruiz-Garcia, L., Bolling, L., Octete, R., Lopez, M., Arnold, C., Ergul, A., Soylemezoglu, G., Uzun, H., Cabello, F., Ibanez, J., Aradhya, M.K., Antanassov, A., Balint, S., Cenis, J., Constantini, L., Gorislavets, S., Grando, M.S., Klein, B.Y., McGovern, P.E., Merdinoglu, D., Pejic, I., Pelsy, F., Primikrios, N., Risovannaya, V., Roubelakis-Angelakia, K.A., Snoussi, H., Sotiri, P., Tamhankar, S., This, P., Troshin, L., Malpica, J.M., Lefort, F., Martinez-Zapater, J.M. 2006. Multiple origins of cultivated grapevine (*Vitis vinifera* L. ssp. *sativa*) based on chloroplast DNA polymorphisms.. *Molecular Ecology* 15: 3707-3714
- Bhat, R., Colowit, P.M., Tai, T., Aradhya, M.K., Browne, G.T. 2006. Genetic and pathogenic variability in *Phytophthora cactorum* affecting fruit and nut crops in California. *Plant Disease*. 90: 161-169.
- Polito, V., Pinney, K., Weinbaum, S., Aradhya, M.K., Dangls, G., Vaknin, Y., Grant, J. 2006. Walnut pollination dynamics: Pollen flow in walnut orchards. *Acta Horticulture*. 705: 85-94.
- Stover, E.W., Ciliento, S., Myers, M. 2006. Trunk sprouting and growth of citrus as affected by naa, aluminum foil, and plastic trunk wraps. *HortScience*. Volume 41: pages 1612-1615
- Stover, E.W., Ciliento, S., Myers, M., Jackson, J., Still, M. 2006. Fruit size and yield of mandarins as influenced by spray volume and surfactant use in naa thinning. *HortScience*. Volume 41: pages 1435-1439

5310-21000-009-00D

MANAGEMENT OF CITRUS AND DATE GENETIC RESOURCES AND INFORMATION – Richard Lee (P) and Robert Krueger; Riverside, California

- Iracheta-Cardenas, M.M., Sandoval-Alejo, B.D., Roman-Calderon, M.E., Keremane, M.L., Lee, R.F., Rocha-Pena, M.A. 2008. Production of Polyclonal Antibodies to the Recombinant Coat Protein of Citrus tristeza virus and Their Effectiveness for Virus Detection. *Journal of Phytopathology*, 156: 243-250
- Keremane, M.L., Halbert, S.E., Ramadugu, C., Webb, S., Lee, R.F. 2008. Detection of *Candidatus Liberibacter asiaticus* in *Diaphorina citri* and its importance in the management of Citrus Huanglongbing in Florida. *Phytopathology*, 98: 387-396
- Lee, R.F., Febres, V.J., Moore, G.A. 2008. Transgenic Resistance to Citrus tristeza virus in Grapefruit. *Plant Cell Reports*, 27: 93-104
- Palmieri, M., Donis, I., Salazar, A.L., Cruz, N., Paniagua, A., Brlansky, R.H., Guerra-Moreno, A., Keremane, M.L., Ballance, P.M., Lee, R.F. 2007. Citrus Viruses in Guatemala: Application of Laboratory-Based Assays. *International Organization of Citrus Virologists Proceedings*, 16th Conference, Pgs. 386-391.
- Ramos, C., Castillo, J.C., Fernandez, O., Rangel, B., Keremane, M.L., Lee, R.F. 2007. Molecular Characterization of Citrus tristeza virus Isolates from Panama. *International Organization of Citrus Virologists Proceedings*, 16th Conference, Pgs. 159-164.
- Rangel, B., Krueger, R. 2007. Stubborn Disease of Citrus in California, *Subtropics newsletter*. 5(4):2-5, 2007, Trade Publication.p

- Guerra-Moreno, A.S., Keremane, M.L., Brilansky, R.H., Lee, R.F. 2007. Citrus Leprosis Symptoms Can be Associated with the Presence of Two Different Viruses: Cytoplasmic and Nuclear, the Former Having a Multipartite RNA Genome. *International Organization of Citrus Virologists Proceedings, 16th Conference*, Pgs. 230-239.
- Chao, C.T., Krueger, R. 2007. The date palm (*Phoenix dactylifera* L.); overview of biology, uses, varieties, and genetics. *HortScience* 42:1077-1082.
- De Azevedo, F.A., Mourao-Filho, De A., Schinor, E.H., De Paoli, L.G., Januzzi-Mendes, B.M., Harakava, R., Gabriel, D.W., Lee, R.F. 2006. GUS Gene Expression Driven by A Citrus Promoter in Transgenic Tobacco and 'Valencia' Sweet Orange. *Pesquisa Agropecuaria Brasileira*, 41:1623-1628.
- Cevik, B., Lee, R.F., Niblett, C.L. 2006. Genetic Transformation of Citrus Paradisi with Antisense and untranslatable RNA-dependent RNA Polymerase Genes of Citrus Tristeza Closterovirus. *Turkish Journal of Agriculture and Forestry*, 30 (3), Pgs. 173-182.
- Lee, R.F., Keremane, M.L., Williams, M., Vidalakis, G. 2006. Huanglongbing, citrus greening: preparing for the future. *Subtropics newsletter*.
- Sandoval-Alejo, B.D., Iracheta-Cardenas, M.M., Keremane, M.L., Lee, R.F., Rocha-Pena, M.A. 2006. Evaluación de Anticuerpos Desarrollados Contra la Proteína Recombinante de la Cápside del Virus Tristeza de los Cítricos. *Revista Mexicana de Fitopatología*, Vol. 24: 137-144.

5320-21000-012-00D

ASSURING LONG TERM PRESERVATION OF TROPICAL CROPS VIA GENETIC RESOURCE MANAGEMENT – Francis Zee (P), Lisa Keith, Tracie Matsumoto Brower, and Dennis Gonsalves; Hilo, Hawaii

- Matsumoto Brower, T.K., Zee, F.T., Suzuki, J.Y., Tripathi, S., Carr, J.B., Mackey, B.E. 2010. Determining sex and screening for the adventitious presence of transgenic material in *Carica papaya* L. seed germplasm. *HortScience*. 45:161-164.
- Keith, L.M. 2008. First report of *Pestalotiopsis virgata* on rambutan in Hawaii. *Plant Disease* 92(5):835.
- Cessna, S.G., Matsumoto Brower, T.K., Lamb, G.N., Rice, S.J., Hochstedler, W.W. 2007. The externally derived portion of the hyperosmotic shock-activated cytosolic calcium pulse mediates adaptation to ionic stress in suspension-cultured tobacco cells. *Journal of Plant Physiology*. 164:815 - 823.
- Huo, N., Lazo, G.R., Vogel, J.P., You, F., Ma, Y., Hayden, D.M., Coleman-Derr, D., Hill, T., Dvorak, J., Anderson, O.D., Luo, M., Gu, Y.Q. 2007. Characterizing the nuclear genome structure of *Brachypodium distachyon* using BAC end sequences. *Functional and Integrative Genomics*. 8:135-147.
- Agbayani, R., Nishijima, W.T., Moore, P.H., Zhu, Y.J. 2007. Elicitor-induced Plant Defense and Pathogenesis-Related Gene Expression in *Carica papaya* L. 19th Annual CTAHR Student Research Symposium. April 7-8, 2006. Honolulu, Hawaii. Abstract #84. pg 51.
- T.K. Matsumoto, M.A. Nagao and B. Mackey. 2007. Off-season flower induction of longan with potassium chlorate and sodium hypochlorite. *HortTechnology* 17(3): 296-300.
- Keith, L.M., Ko, W., Sato, D.M. 2006. Identification Guide for Diseases of Tea (*Camellia sinensis*). Extension Publications. PD-33. Available: <http://www.ctahr.hawaii.edu/oc/freepubs/pdf/SCM-14.pdf>
- Schenk, A., Berger, M., Keith, L.M., Bender, C.L., Muskhelishvili, G., Ullrich, M.S. 2006. The algT gene of *Pseudomonas syringae* pv. *glycinea* and new insights into the transcriptional organization of the algT-muc gene cluster. *Journal of Bacteriology*. 188(23):8013-8021
- Agbayani, R., Nishijima, W.T., Moore, P.H., Zhu, Y.J. 2006. Characterization and Improvement of Disease Response to *Phytophthora Palmivora* in Papaya. 18th Annual CTAHR Student Research Symposium [abstract]. Paper No. 104-63.
- Gu, Y., Lazo, G., Vogel, J., Coleman-Derr, D., Luo, M.C., Thilmony, R., Garvin, D., Anderson, O., Huo, N. 2006. Construction and characterization of two BAC libraries from *Brachypodium distachyon*, a new model for grass genomics. *Genome*. 49(9):1099-1108.
- Eustice, M., Lai, C.W., Yu, Q., Presting, G., Moore, P.H., Ming, R. 2006. Variation of microsatellite markers among selected papaya varieties. 18th Annual CTAHR Student Research Symposium. (Abstracts)#77, pg. 49.

5320-21000-013-00D

MOLECULAR RESOURCES FOR THE IMPROVEMENT OF TROPICAL ORNAMENTAL AND FRUIT CROPS – Dennis Gonsalves (P), Roxana Cabos, and Jon Suzuki; Hilo, Hawaii

- Tripathi, S., Suzuki, J.Y., Carr, J.B., McQuate, G.T., Ferreira, S.A., Manshardt, R.M., Pitz, K.Y., Wall, M.M., Gonsalves, D. 2011. Nutritional Composition of Rainbow Papaya, the First Commercialized Transgenic Fruit Crop. *Journal of Food Composition and Analysis*. 24(2):140-147.
- Cabos, R.Y., Sipes, B.S., Nagai, C., Serracin, M., Schmitt, D.P. 2010. Evaluation of Coffee Genotypes for Root-knot Nematode Resistance. *Nematopica* 40:191-202.

Paret, M. L., Cabos, R., Kratky, B. A., and Alvarez, A. M. 2010. Effect of plant essential oils on *Ralstonia solanacearum* race 4 and bacterial wilt of edible ginger. *Plant Dis.* 94:521-527.

5325-21000-014-00D

AN INTEGRATED DATABASE AND BIOINFORMATICS RESOURCE FOR SMALL GRAINS – Olin Anderson (P), Yong Gu, Gerard Lazo, and David Matthews; Albany, California

- You, F., Huo, N., Deal, K., Gu, Y.Q., Luo, M., McGuire, P., Dvorak, J., Anderson, O.D. 2011. Annotation-based genome-wide SNP discovery in the large and complex *Aegilops tauschii* genome using next-generation sequencing without a reference genome sequence. *Biomed Central (BMC) Genomics.* 12:59.
- You, F., Wanjugi, H., Huo, N., Lazo, G.R., Luo, M., Anderson, O.D., Dvorak, J., Gu, Y.Q. 2010. RJPrimers: unique transposable element insertion junction discovery and PCR primer design for marker development. *Nucleic Acids Research.* 38:Suppl:W310-320.
- Anderson, O.D., Coleman-Derr, D., Gu, Y.Q., Heath, S. 2010. Structural and transcriptional analysis of plant genes encoding the bifunctional lysine ketoglutarate reductase saccharopine dehydrogenase enzyme. *Biomed Central (BMC) Genomics.* 10:113.
- Anderson, O.D. 2009. EST mining for structure and expression of genes in the region of the wheat high-molecular-weight glutenin loci. *Genome.* 52:1-15.
- Gu, Y.Q., Ma, Y., Huo, N., Vogel, J.P., You, F., Lazo, G.R., Nelson, W., Soderlund, C., Dvorak, J. 2009. A BAC-based physical map of *Brachypodium distachyon* and its comparative analysis with rice and wheat. *Biomed Central (BMC) Genomics.* 10:496
- Luo, M.C., Deal, K.R., Akhunov, E.D., Akhunova, A.R., Anderson, O.D., Anderson, J.A., Blake, N., Clegg, M.T., Coleman-Derr, D., Conley, E.E., Crossman, C.C., Dubcovsky, J., Gill, B.S., Gu, Y.Q., Hadam, J., Heo, H., Huo, N., Lazo, G.R., Lundy, K.E., Ma, Y., Matthews, D.E., McGuire, P.E., Morrell, P.L., Nicolet, C.M., Qualset, C.O., Renfro, J., Tabano, D., Talbert, L.E., Tian, A., Toleno, D.M., Warburton, M.L., You, F.M., Zhang, W.J., Dvorak, J. 2009. Genome Comparisons Reveal a Dominant Mechanism of Chromosome Number Reduction in Grasses and Accelerated Genome Evolution in Triticeae. *Proceedings of the National Academy of Sciences.* 106:15780-15785.
- Michalak, M.K., Ghavami, F., Lazo, G.R., Gu, Y.Q., Kianian, S.F. 2009. Evolutionary relationship of nuclear genes encoding mitochondrial proteins across grasses. *Maydica.* 54: 471-483
- Wanjugi, H., Coleman-Derr, D., Huo, N., Kianian, S.F., Luo, M., Wu, J., Anderson, O.D., Gu, Y.Q. 2009. Rapid development of PCR-based genome-specific repetitive DNA junction markers in wheat. *Genome.* 52:576-587.
- Yu, G., Cai, X., Harris, M., Gu, Y.Q., Luo, M., Xu, S.S. 2009. Saturation and comparative mapping of genomic region harboring Hessian fly resistance gene H26 in wheat. *Theoretical and Applied Genetics.* 118(8):1589-1599.
- Wanjugi, H., Coleman-Derr, D., Huo, N., Kianian, S.F., Luo, M., Wu, J., Anderson, O.D., Gu, Y.Q. 2009. Rapid development of PCR-based genome-specific repetitive DNA junction markers in wheat. *Genome.* 52:576-587.
- Anderson, O.D., Gu, Y.Q., Kong, K., Lazo, G.R., Wu, J. 2009. Structure of the omega-gliadin gene family. *Functional and Integrative Genomics.* 9: 397-410.
- Bai, W., Cai, A., Rz, Z., Li, A., Huo, N., Li, S., Gu, Y.Q., Yabut, L., Jia, J., Qi, Y. 2009. Novel microRNAs uncovered by deep sequencing of small RNA transcriptomes in bread wheat (*Triticum aestivum* L.) and *Brachypodium distachyon* (L.) Beauv. *Functional and Integrative Genomics.* 9:532-541.
- Bortiri, E., Coleman-Derr, D., Lazo, G.R., Anderson, O.D., Gu, Y.Q. 2008. The complete chloroplast genome sequence of *Brachypodium distachyon*: sequence comparison and phylogenetic analysis of eight grass plastomes. *BMC Research Notes.* 1:61.
- You, F.M., Huo, N., Gu, Y.Q., Luo, M., Ma, Y., Hane, D., Lazo, G.R., Dvorak, J., Anderson, O.D. 2008. BatchPrimer3: A high throughput web application for PCR and sequencing primer design. *BMC Bioinformatics.* 9:253.
- Chingcuanco, D.L., Vensel, W.H. 2008. Globulins are the main seed storage proteins in *brachypodium distachyon*. *Theoretical and Applied Genetics.* 117:555-563.
- Huo, N., Lazo, G.R., Vogel, J.P., You, F., Ma, Y., Hayden, D.M., Coleman-Derr, D., Hill, T., Dvorak, J., Anderson, O.D., Luo, M., Gu, Y.Q. 2007. Characterizing the nuclear genome structure of *Brachypodium distachyon* using BAC end sequences. *Functional and Integrative Genomics.* 8:135-147.
- You, F.M., Luo, M.C., Gu, Y.Q., Lazo, G.R., Thomas, C., McGuire, P.E., Dvorak, J., and Anderson, O.D. 2007. Genoprofiler: Automated processing of high throughput capillary fingerprinting data. *Bioinformatics.* 23:240-242.
- Gao, S., Gu, Y.Q., Wu, J., Huo, N., Crossman, C., Jia, J., Ren, Z., Anderson, O.D., Xiuying, K. 2007. Rapid evolution and complex structural organization in genomic regions harboring multiple prolamin genes in the polyploid wheat genome. *Plant Molecular Biology.* 65:189-203.

- Kalavacharla, V., Hossain, K., Gu, Y.Q., Riera, L.O., Isabel, V.M., Bhamidimarri, S., Gonzalez-Hernandez, J., Maan, S., Kianian, S. 2006. High-resolution radiation hybrid map of wheat chromosome 1d. *Genetics*. 173:1-11.
- Szucs, P., Karsai, I., Von Zitzewitz, J., Meszaros, K., Copper, L., Gu, Y.Q., Chen, T., Hayes, P., Skinner, J. 2006. Positional relationship between photoperiod response qtl and photoreceptor and vernalization genes in barley. *Theoretical and Applied Genetics*. 112: 1,277-1,285.
- Gu, Y., Salse, J., Coleman-Derr, D., Dupin, A., Crossman, C., Lazo, G., Huo, N., Belcram, H., Ravel, C., Charmet, G., Charles, M., Anderson, O., Chalhou, B. 2006. Types and Rates of Sequence Evolution at the HMW-Glutelin Locus in Hexaploid Wheat and Its Ancestral Genomes. *Genetics*. 174:1493-1504.
- Gu, Y., Lazo, G., Vogel, J., Coleman-Derr, D., Luo, M.C., Thilmony, R., Garvin, D., Anderson, O., Huo, N. 2006. Construction and characterization of two BAC libraries from *Brachypodium distachyon*, a new model for grass genomics. *Genome*. 49(9):1099-1108.

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GENOMICS AND MOLECULAR APPROACHES TO WHEAT TRAITS – Olin Anderson (P), Yong Gu, and Debbie Chingcuanco; Albany, California

- Akhunov, E.D., Akhunova, A., Anderson, O.D., Anderson, J., Blake, N., Clegg, M., Coleman-Derr, D., Conley, E., Crossman, C., Deal, K., Dubcovsky, J., Gill, B., Gu, Y.Q., Hadam, J., Heo, H., Huo, N., Lazo, G.R., Luo, M., Ma, Y., Matthews, D.E., McGuire, P., Morrell, P., Qualset, C., Renfro, J., Tabanao, D., Talbert, L., Tian, C., Toleno, D., Warburton, M., You, F., Zhang, W., Dvorak, J. 2010. Nucleotide diversity maps reveal variation in diversity among wheat genomes and chromosomes. *Biomed Central (BMC) Genomics*. 11:702-710.
- Anderson, O.D., Coleman-Derr, D., Gu, Y.Q., Heath, S. 2010. Structural and transcriptional analysis of plant genes encoding the bifunctional lysine ketoglutarate reductase saccharopine dehydrogenase enzyme. *Biomed Central (BMC) Genomics*. 10:113.
- Saathoff, A.J., Chow, E.K., Dien, B.S., Tobias, C.M. 2011. Downregulation of Cinnamyl-Alcohol Dehydrogenase in Switchgrass by RNA Silencing Results in Enhanced Glucose Release after Cellulase Treatment. *PLoS One*. 6(1): e16416. doi:10.1371/journal.pone.0016416.
- Okada, M., Lanzatella-Craig, C., Saha, M., Bouton, J., Wu, R., Tobias, C.M. 2010. Complete Switchgrass Genetic Maps Reveal Subgenome Collinearity, Preferential Pairing and Multilocus Interactions. *Genetics*. 185:745-760.
- Fleury, D., Luo, M., Dvorak, J., Ramsay, L., Gill, B., Anderson, O.D., You, F., Schoaei, Z., Deal, K., Langridge, P. 2010. Physical mapping of a large plant genome using global high-information content fingerprinting: a distal region of wheat chromosome 3DS. *Genetics*. 11:382.
- Fleury, D., Luo, M., Dvorak, J., Ramsay, L., Gill, B., Anderson, O.D., You, F., Schoaei, Z., Deal, K., Langridge, P. 2010. Physical mapping of a large plant genome using global high-information-content-fingerprinting: the distal region of the wheat ancestor *Aegilops tauschii* chromosome 3DS. *Biomed Central (BMC) Plant Biology*. 10:113.
- Bevan, M.W., Garvin, D.F., Vogel, J.P. 2010. *Brachypodium distachyon* genomics for sustainable food and fuel production. *Current Opinion in Biotechnology*. 21:211-217.
- Anderson, O.D., Gu, Y.Q., Kong, K., Lazo, G.R., Wu, J. 2009. Structure of the omega-gliadin gene family. *Functional and Integrative Genomics*. 9: 397-410.
- Bai, W., Cai, A., Rz, Z., Li, A., Huo, N., Li, S., Gu, Y.Q., Yabut, L., Jia, J., Qi, Y. 2009. Novel microRNAs uncovered by deep sequencing of small RNA transcriptomes in bread wheat (*Triticum aestivum* L.) and *Brachypodium distachyon* (L.) Beauv. *Functional and Integrative Genomics*. 9:532-541.
- Gu, Y.Q., Wanjugi, H., Coleman-Derr, D., Kong, X., Anderson, O.D. 2009. Conserved globulin gene across eight grass genomes identify fundamental units of the loci encoding seed storage proteins. *Functional and Integrative Genomics*. 10:111-122.
- Huo, N., Vogel, J.P., Lazo, G.R., You, F., Ma, Y., McMahon, S.A., Dvorak, J., Anderson, O.D., Luo, M., Gu, Y.Q. 2009. Structural characterization of *Brachypodium* genome and its syntenic relationship with rice and wheat. *Plant Molecular Biology*. 70: 47-61.
- Kalavacharla, V., Hossain, K., Gu, Y.Q., Maan, S., Kianian, S. 2009. Radiation hybrid mapping in crop plants. *Advances in Agronomy*. 102:199-219.
- Gu, Y.Q., Ma, Y., Huo, N., Vogel, J.P., You, F., Lazo, G.R., Nelson, W., Soderlund, C., Dvorak, J. 2009. A BAC-based physical map of *Brachypodium distachyon* and its comparative analysis with rice and wheat. *Biomed Central (BMC) Genomics*. 10:496
- Michalak, M.K., Ghavami, F., Lazo, G.R., Gu, Y.Q., Kianian, S.F. 2009. Evolutionary relationship of nuclear genes encoding mitochondrial proteins across grasses. *Maydica*. 54: 471-483
- Gu, Y.Q., Wanjugi, H., Coleman-Derr, D., Kong, X., Anderson, O.D. 2009. Conserved globulin gene across eight grass genomes identify fundamental units of the loci encoding seed storage proteins. *Functional and Integrative Genomics*. 10:111-122.

- Stamova, B., Roessner, U., Suren, S., Chingcuanco, D.L., Basic, A., Beckles, D. 2008. Metabolic profiling of transgenic wheat over-expressing the high-molecular weight Dx5 glutenin subunit. *Metabolomics*. 5(2):239-252.
- Scossa, F., Chingcuanco, D.L., Anderson, O.D., Vensel, W.H., Lafiandra, D., D'Ovidio, R., Masci, S. 2008. Comparative transcriptional and proteomic profiling of bread wheat cultivar and its derived transgenic line over-expressing a low molecular weight glutenin subunit gene in the endosperm. *Molecular and Cellular Proteomics*. 8(14):2948-2966.
- Chingcuanco, D.L., Vensel, W.H. 2008. Globulins are the main seed storage proteins in brachypodium distachyon. *Theoretical and Applied Genetics*. 117:555-563.
- Zhang, S., Gu, Y., Singh, J., Coleman-Derr, D., Brar, D., Jiang, N., Lemaux, P. 2007. New Insights into *Oryza* genome evolution: high gene colinearity and differential retrotransposon amplification. *Plant Molecular Biology*. 64:589-600.
- Gao, S., Gu, Y.Q., Wu, J., Huo, N., Crossman, C., Jia, J., Ren, Z., Anderson, O.D., Xiuying, K. 2007. Rapid evolution and complex structural organization in genomic regions harboring multiple prolamin genes in the polyploid wheat genome. *Plant Molecular Biology*. 65:189-203.
- Szucs, P., Karsai, I., Von Zitzewitz, J., Meszaros, K., Cooper, L., Gu, Y.Q., Chen, T., Hayes, P., Skinner, J. 2006. Positional relationship between photoperiod response qtl and photoreceptor and vernalization genes in barley. *Theoretical and Applied Genetics*. 112: 1,277-1,285.
- Gu, Y., Salse, J., Coleman-Derr, D., Dupin, A., Crossman, C., Lazo, G., Huo, N., Belcram, H., Ravel, C., Charmet, G., Charles, M., Anderson, O., Chalhou, B. 2006. Types and Rates of Sequence Evolution at the HMW-Glutenin Locus in Hexaploid Wheat and Its Ancestral Genomes. *Genetics*. 174:1493-1504.

5325-21000-018-00D

MOLECULAR TOOLS TO MINIMIZE RISK IN GENETICALLY ENGINEERED CROPS – Ann Blechl (P), Roger Thilmony, and James Thomson; Albany, California

- Thomson, J.G., Cook, M.A., Guttman, M.E., Smith, J.D., Thilmony, R.L. 2011. Novel null binary vectors enable an inexpensive foliar selection method in *Arabidopsis*. *BMC Research Notes*. 4(44) Available: <http://www.biomedcentral.com/1756-0500/4/44>.

5325-21420-004-00D

MOLECULAR GENETIC TOOLS FOR POTATO AND FRUIT TREE IMPROVEMENT – William Belknap (P) and Kent McCue; Albany, California

- Dupont, F.M., Vensel, W.H., Tanaka, C.K., Hurkman II, W.J., Altenbach, S.B. 2011. Deciphering the complexities of the wheat flour proteome using quantitative two-dimensional electrophoresis, three proteases and tandem mass spectrometry. *Proteome Science*. 9(10) available:<http://www.proteomesci.com/content/9/1/10>.
- Oosumi, T., Rockhold, D.R., Maccree, M.M., Deahl, K.L., Mc Cue, K.F., Belknap, W.R. 2009. Gene Rpi-bt1 from *Solanum bulbocastanum* confers resistance to late blight in transgenic potatoes. *American Journal of Potato Research*. 86(6):456-465.
- Mc Cue, K.F. 2009. Potato Glycoalkaloids, Past Present and Future. *Fruit, Vegetable, and Cereal Science and Biotechnology*. 3:65-71.
- Harmon, F.G., Imaizumi, T., Gray, W.M. 2008. CUL1 Regulates TOC1 Protein Stability in the *Arabidopsis* Circadian Clock. *Plant Journal*. 55(4):568-579.
- Rockhold, D.R., Chang, S., Taylor, N., Allen, P.V., Mc Cue, K.F., Belknap, W.R. 2008. Structure of two *Solanum bulbocastanum* polyubiquitin genes and expression of their promoters in transgenic potatoes. *American Journal of Potato Research*. 85:219-226.
- Rommens, C.M., Haring, M.A., Swords, K., Davies, H.V., Belknap, W.R. 2007. The Intragenic Approach as a New Extension to Traditional Plant Breeding. *Trends in Plant Science*. 12:397-403.
- Mc Cue, K.F., Allen, P.V., Shepherd, L.V., Blake, A., Maccree, M.M., Rockhold, D.R., Novy, R.G., Stewart, D., Davies, H.V., Belknap, W.R. 2007. Potato glycosyltransferase, the terminal step in triose side-chain biosynthesis. *Phytochemistry*. 68 (3): 327-334.
- Mc Cue, K.F., Allen, P.V., Shepherd, L.V., Blake, A., Rockhold, D.R., Novy, R.G., Stewart, D., Davies, H.V., Belknap, W.R. 2007. Manipulation and compensation of steroidal glycoalkaloid biosynthesis in potatoes. *Acta Horticulturae*. 745: 343-349.
- Faigon-Soverna, A., Harmon, F.G., Storani, L., Karayekov, E., Staneloni, R.J., Gassmann, W., Mias, P., Casal, J.J., Kay, S.A., Yanovsky, M.J. 2006. A Constitutive Shade-Avoidance Mutant Implicates TIR-NBS-LRR Proteins in *Arabidopsis* Photomorphogenic Development. *The Plant Cell* 18(11):2919-2928.

5325-21430-012-00D

PRODUCTION OF WHEAT GERMPLASM WITH ENHANCED BAKING QUALITY – Ann Blechl (P); Albany, California

Di, R., Blechl, A.E., Dill-Macky, R., Tortora, A., Tumer, N.E. 2010. Expression of a truncated form of yeast ribosomal protein L3 in transgenic wheat improves resistance to Fusarium head blight. *Plant Science*. 178: 374-380.

5325-43000-027-00D

MOLECULAR ANALYSIS OF EFFECTS OF ENVIRONMENT ON WHEAT FLOUR QUALITY AND ALLERGENIC POTENTIAL – William Hurkman (P), William Vensel, and Susan Altenbach; Albany, California

Dupont, F.M., Vensel, W.H., Tanaka, C.K., Hurkman II, W.J., Altenbach, S.B. 2011. Deciphering the complexities of the wheat flour proteome using quantitative two-dimensional electrophoresis, three proteases and tandem mass spectrometry. *Proteome Science*. 9(10) available:<http://www.proteomesci.com/content/9/1/10>.

Altenbach, S.B., Vensel, W.H., Dupont, F.M. 2010. Integration of transcriptomic and proteomic data from a single wheat cultivar provides new tools for understanding the roles of individual alpha gliadin proteins in flour quality and celiac disease. *Journal of Cereal Science*. 52:143-151.

Altenbach, S.B., Vensel, W.H., Dupont, F.M. 2010. Analysis of expressed sequence tags from a single wheat cultivar facilitates interpretation of tandem mass spectrometry data and discrimination of gamma gliadin proteins that may play different functional roles in flour. *Biomed Central (BMC) Plant Biology*. 10(7):1471-2229. Available: <http://www.biomedcentral.com/1471-2229/10/7>.

5335-21000-026-00D

GENOMIC AND GENETIC ANALYSIS OF THE CEREAL CIRCADIAN SYSTEM AND CROP PRODUCTION – Frank Harmon (P); Albany, California

Harmon, F.G., Rowe, S.C., Schnable, J. 2010. Genome Sequencing and Analysis of the Model grass *Brachypodium distachyon*. *Nature*. 463(7282):763-8.

Khan, S., Rowe, S.C., Harmon, F.G. 2010. Coordination of the maize transcriptome by a conserved circadian clock. *Biomed Central (BMC) Plant Biology*. 10:126.

Thines, B., Harmon, F.G. 2010. Ambient temperature response establishes ELF3 as a required component of the *Arabidopsis* core circadian clock. *Proceedings of the National Academy of Sciences*. 107(7):3257-3262.

Thines, B., Harmon, F.G. 2010. Four easy pieces: mechanisms underlying circadian regulation of growth and development. *Current Opinion in Plant Biology*. 14:1–7.

Harmon, F.G., Imaizumi, T., Gray, W.M. 2008. CUL1 Regulates TOC1 Protein Stability in the *Arabidopsis* Circadian Clock. *Plant Journal*. 55(4):568-579.

Para, A., Farre, E.M., Imaizumi, T., Pruneda-Paz, J., Harmon, F.G., Kay, S.A. 2007. PRR3 Is a Vascular Regulator of TOC1 Stability in the *Arabidopsis* Circadian Clock. *The Plant Cell*. Published on November 30, 2007; 10.1105/tpc.107.054775.

Faigon-Soverna, A., Harmon, F.G., Storani, L., Karayekov, E., Staneloni, R.J., Gassmann, W., Mias, P., Casal, J.J., Kay, S.A., Yanovsky, M.J. 2006. A Constitutive Shade-Avoidance Mutant Implicates TIR-NBS-LRR Proteins in *Arabidopsis* Photomorphogenic Development. *The Plant Cell* 18(11):2919-2928.

5335-21000-032-00D

MOLECULAR MECHANISMS OF PHYTOCHROME SIGNALING AND GENE REGULATION – Sarah Hake (P); Albany, California

Quail, P.H. 2010. Phytochromes. *Current Biology*. 20(12):R504-R507.

Leivar, P., Teppermann, J.M., Monte, E., Calderon, R.H., Liu, T.L., Quail, P.H. 2009. Definition of Early Transcriptional Circuitry Involved in Light-Induced Reversal of PIF-Imposed Repression of Photomorphogenesis in Young *Arabidopsis* Seedlings. *The Plant Cell*. 21:3535-3553.

5335-21000-034-00D

POSITIONAL CLONING IN MAIZE OF GENES THAT REGULATE PLANT ARCHITECTURE– Sarah Hake (P); Albany, California

- Johnston, R., Candela, H., Hake, S.C., Foster, T. 2010. The maize milkweed pod1 mutant reveals a mechanism to modify organ morphology. *Genesis*. Published online 10.1002/dvg.20622.
- Chuck, G., Whipple, C., Jackson, D., Hake, S.C. 2010. The maize SBP-box transcription factor encoded by tasselsheath4 regulates bract development and the establishment of meristem boundaries. *Development*. 137(8):1243-1250.

5335-21000-036-00D

MOLECULAR DEVELOPMENTAL GENETICS OF POLLEN AND POLLEN-PISTIL INTERACTIONS IN CROP PLANTS – Shelia McCormick (P); Albany, California

- Zheng, B., Chen, X. and McCormick, S. 2011. The anaphase-promoting complex/cyclosome is a dual integrator that regulates both miRNA-mediated transcriptional regulation of Cyclin B1 and degradation of Cyclin B1 during Arabidopsis male gametophyte development. *Plant Cell* 23: 1033-1046, doi: 10.1105/tpc.111.083980.
- Muschietti, J., McCormick, S.M. 2010. Abscissic acid (ABA) receptors: light at the end of the tunnel. *Faculty of 1000 Biology*. 2:15 (doi: 10.3410/B2-15).
- Wengier, D., Mazzeckla, M., Salem, T., McCormick, S.M., Muschietti, J. 2010. STIL, a peculiar molecule from styles, specifically dephosphorylates the pollen receptor kinase LePRK2 and stimulates pollen tube growth in vitro. *Biomed Central (BMC) Plant Biology*. Online 1471-2229/10/33.
- Zhang, Y., He, J., Lee, D., McCormick, S.M. 2010. Interdependence of endomembrane trafficking and actin dynamics during polarized growth of Arabidopsis pollen tubes. *Plant Physiology*. 152:2200-2210.
- Ron, M., Alandete-Saez, M., Williams, L.C., Fletcher, J.C., McCormick, E.J. 2010. Proper regulation of a sperm-specific cis-nat-siRNA is essential for double fertilization in Arabidopsis. *Genes and Development*. 24(10):24: 1010-1021.
- Guan, Y., Boavida, L.C., McCormick, S.M. 2010. Gametogenesis. *Encyclopedia for Life Sciences*. Online 1/15/2010. DOI: 10.1002/9780470015902.a0002037.pub2.

5335-21000-038-00D

FUNCTIONAL GENOMICS OF PLANT ARCHITECTURE – Jennifer Fletcher (P); Albany, California

- Ha, C.M., Jun, J.H., Fletcher, J.C. 2010. Control of Arabidopsis leaf morphogenesis through regulation of the YABBY and KNOX families of transcription factors. *Genetics*. 186(1):197-206. Epub 2010 Jul 6.
- Jun, J.H., Fiume, E., Roeder, A.H., Meng, L., Sharma, V.K., Osmont, K.S., Baker, C., Ha, C.M., Meyerowitz, E.M., Feldman, L.J., Fletcher, J.C. 2010. Comprehensive analysis of CLE polypeptide signaling gene expression and over-expression activity in Arabidopsis. *Plant Physiology*. 154:1721-1736.
- Jun, J., Ha, C., Fletcher, J.C. 2010. BLADE-ON-PETIOLE1 coordinates organ determinacy and axial polarity in Arabidopsis by directly activating ASYMMETRIC LEAVES2. *The Plant Cell*. 22:62-76.
- Meng, L., Ruth, K.C., Fletcher, J.C., Feldman, L. 2010. The Roles of Different CLE Domains in Arabidopsis CLE Polypeptide Activity and Functional Specificity. *Molecular Plant*. Published online doi:10.1093/mp/ssq021.

5341-21000-004-00D

ARCTIC AND SUBARCTIC PLANT GENETIC RESOURCES CONSERVATION, RESEARCH, AND INFORMATION MANAGEMENT – Nancy Robertson (P), Danny Barney, and Kim Hummer; Palmer, Alaska

- Robertson, N.L., Smeenk, J., Anderson, J. 2011. Molecular Characterization of Potato Leafroll virus, Potato virus A, and Potato virus X Isolates from Potatoes in Alaskan Cities and Villages. *Plant Health Progress*. Available: <http://www.plantmanagementnetwork.org/php/>.
- Barney, D.L., Hummer, K.E. 2011. Blueberry Trials on Alaska's Kenai Peninsula: First Year Report. Extension Reports. <http://www.ars.usda.gov/Main/docs.htm?docid=19218>.
- Pantoja, A., Hagerty, A.M., Emmert, S.Y., Kuhl, J.C., Pike, K., Alvarez, J.M., Jensen, A. 2010. Aphids (Homoptera: Aphididae) associated with rhubarb (*Rheum* spp.) in the Matanuska Valley, Alaska: species composition, seasonal abundance, and potential virus vectors. *Journal of British Columbia Entomological Society*. 107:75-81.

- Robertson, N.L., Brown, K.L. 2010. First Report of Bean Yellow Mosaic Virus in Alaska from Clover (*Trifolium* spp.). *Plant Disease*. 94(3):372.
- Robertson, N.L., Furman, B.J. 2010. Serological Detection and Molecular Analysis of Tobacco Ringspot Virus and Strawberry Latent Ringspot Virus in Mint (*Mentha* sp.). *Phytopathology*. 100(6):S109.
- Robertson, N.L., Brown, K.L., Winton, L.M., Holloway, P.S. 2009. First Report of Tobacco Rattle Virus in Peony in Alaska. *Plant Disease*. 93(6):675.
- Pantoja, A., Kuhl, J.C. 2009. Morphologic Variation in the USDA/ARS Rhubarb Germplasm Collection. *Plant Genetic Resources*. Available: <http://journals.cambridge.org/action/displayAbstract?aid=5946564>.
- Robertson, N.L., Coyne, C.J. 2009. First Report of Bean Yellow Mosaic Virus from Diseased *Lupinus luteus* L. in Eastern Washington. *Plant Disease*. 93(3):319.
- Robertson, N.L., Brown, K.L. 2009. Identification and Molecular Characterization of a Potyvirus Isolated from Native Larkspur (*Delphinium glaucum*) in Alaska. *Plant Disease*. 93(4):428.
- Deridder, B.P., Crafts-Brandner, S.J. 2008. Chilling stress response of post-emergent cotton seedlings. *Physiologia Plantarum* 134: 430-439
- Dyer, J.M., Mullen, R.T. (2008) Engineering plant oils as high-value industrial feedstocks for biorefining: the need for underpinning cell biology research. *Physiologia Plantarum*, (132):11-22.
- Kang, H., Kuhl, J.C., Kachroo, P., Klessig, D.F. 2008. CRT1, an Arabidopsis ATPase that Interacts with Diverse Resistance Proteins and Modulates Disease Resistance to Turnip Crinkle Virus. *Cell Host and Microbe*. 3:48-57.
- Kuhl, J.C., Deboer, V.L. 2008. Genetic Diversity of Rhubarb Cultivars. *Journal of the American Society for Horticultural Science*. 133(4):1-6.
- Deridder, B.P., Salvucci, M.E. 2007. Recovery of rubisco activase gene expression during heat stress involves post-transcriptional control. *Plant Science*, 172: 246-254.
- Kuhl, J.C., Zarka, K., Coombs, J., Kirk, W.W., Douches, D.S. 2007. Late Blight Resistance of RB Transgenic Potato Lines. *Journal of the American Society for Horticultural Science*. 132(6):783-789.
- Robertson, N.L. 2007. Identification and characterization of a new virus in the genus Potyvirus from wild populations of *Angelica lucida* L. and *A. genuflexa* Nutt., family Apiacea. *Archives of Virology*. 152(9):1603-1611. DOI 10.1007/s00705-007-1003-4.
- Robertson, N.L., Cote, F., Pare, C., Leblanc, E., Bergeron, M., Leclerc, D. 2007. Complete nucleotide sequence of Nootka lupine vein-clearing virus. *Virus Genes*. 35(3):807-814.

5347-21000-011-00D

PHYSIOLOGICAL AND GENETIC BASIS OF COTTON ACCLIMATION TO ABIOTIC STRESS – Michael Gore (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

- Barta, C., Chausse, A., Wachter, R.M., Salvucci, M.E. Structural Changes Associated with the Acute Thermal Instability of Rubisco Activase. *Archives of Biochemistry and Biophysics* (2010) 499:17-25.
- Gore, M.A., Coyle, G.G., Friebe, B., Coffelt, T.A., Salvucci, M.E., 2010. Complex Ploidy Level Variation in Guayule Breeding Programs. *Crop Science*, 51:210-216.

5347-21410-005-00D

COMMERCIALIZATION OF NEW INDUSTRIAL CROP GERmplasm AND CROPPING SYSTEMS – Terry Coffelt (P), Michael Salvucci, and John Dyer; Maricopa, Arizona

- Gidda, S.K., Shockey, J.M., Rothstein, S.J., Dyer, J.M., Mullen, R.T., 2011. Hydrophobic-domain-dependent protein-protein interactions mediate the localization of GPAT enzymes to ER subdomains. *Traffic*. 12:452-472.
- Coffelt, T.A. and Johnson, L. (2010). A Set of Descriptions for Evaluating Guayule Germplasm. *Industrial Crops and Products*, doi.org/10.1016/j.indcrop.2010.04.2004.
- Coffelt, T.A. and Ray, D.T. (2010). Cutting height effects on guayule latex, rubber and resin yields. *Industrial Crops and Products*, DOI:10.1016/j.indcrop.2010.04.018.
- Ray, D.T., Foster, M.A., Coffelt, T.A., McMahan, C.M. *Guayule: Culture, Breeding and Rubber Production*, *Industrial Crops and Uses* (2010), pp. 384-410.
- Bekaardt, C.R., Coffelt, T.A., Fenwick, J.R., Wiesner, L.E. Environmental, irrigation and fertilization impacts on the seed quality of guayule (*Parthenium argentatum* Gray). *Industrial Crops and Products* (2010), Vol. 31, Issue 3, pp. 427-436.
- James, C.N., Horn, P.J., Case, C.R., Gidda, S.K., Zhang, D., Mullen, R.T., Dyer, J.M., Anderson, R.G., and Chapman, K.D. (2010). Disruption of the Arabidopsis CGI-58 homologue produces Chanarin-Dorfman-like lipodystrophy in plants. *Proceedings of the National Academy of Sciences of the United States of America*, 107:17833-17838.

- Coffelt, T.A., Williams, C.F. 2009. Characterization and recycling of waste water from guayule latex extraction. *Industrial Crops and Products* (29), pp. 648-653.
- Gidda, S.K., Shockey, J.M., Rothstein, S.J., Dyer, J.M., Mullen, R.T. (2009). Arabidopsis thaliana GPAT8 and GPAT9 are localized to the ER and possess distinct ER retrieval signals: functional divergence of the dilysine ER retrieval motif in plant cells. *Plant Physiology and Biochemistry*, 47:867-879.
- Jenderek, M.M., Dierig, D.A., Isbell, T. 2009. Fatty acid profile of Lesquerella germplasm in the National Plant Germplasm System collection. *Industrial Crops and Products*, Vol. 29, Issue 1, pp. 154-164.
- Coffelt, T.A. and Johnson, L. (2009). A set of descriptors for evaluating guayule germplasm. In: Cermack, S.C. and M. Berti, Eds. 21st Annual Meeting of the Association for Advancement of Industrial Crops (AAIC), Chillan, Chile, p. 18.
- Lessire, R., Cahoon, E., Chapman, K., Dyer, J.M., Eastmond, P., Heinz, E. 2009. Highlights of Recent Progress in Plant Lipid Research. *Plant Physiology and Biochemistry*. 47:443-447.
- O'Quin, J.B., Mullen, R.T., Dyer, J.M. 2009. Addition of an N-terminal epitope tag significantly increases the activity of plant fatty acid desaturases expressed in yeast cells. *Applied Microbiology and Biotechnology*. 83:117-125.
- Slocombe, S.P., Cornah, J., Pinfield-Wells, H., Soady, K., Zhang, Q.Y., Gilday, A., Dyer, J.M., Graham, I.A. (2009). Oil accumulation in leaves directed by modification of fatty acid breakdown and lipid synthesis pathways. *Plant Biotechnology Journal*, 7:694-703.
- Coffelt, T.A., Nakayama, F.S., Ray, D.T., Cornish, K., McMahan, C.M. 2009. Post-harvest storage effects on guayule latex and rubber yields. *Industrial Crops and Products*. 29(2-3):326-335.
- Chow, P., Nakayama, F.S., Blahnik, B., Youngquist, J., Coffelt, T.A. 2008. Chemical constituents and physical properties of guayule wood and bark. *Industrial Crops and Products*, 28:303-308.
- Attanayake, K., Glawe, D., McPhee, K.E., Dugan, F.M., Chen, W. 2008. First report of powdery mildew of chickpea (*Cicer arietinum*) caused by *Leveillula taurica* in Washington State. *Plant Health Progress*. doi:10.1094/PHP-2008-0702-01-BR.
- Lessire, R., Beisson, F., Chapman, K., Dyer, J.M., Eastmond, P. (2008). Highlights of the International Symposium on Plant Lipids, 18th ISPL 2008, Bordeaux July 20–25th, 2008. *European Journal of Lipid Science and Technology* 110:1190-1194.
- Bhardwaj, H.L., Hamama, A.A., Dierig, D.A. 2007. Fatty acids in vernonia produced in the mid-atlantic region of usa. *J. Amer. Oil Chem. Soc.*, 84:393-397.
- Foster, M.A., Dierig, D.A., Wintermeyer, M. 2007. A weed control program for establishing lesquerella. *Issues in New Crops and New Uses*. In: Janick, J. and Whipkey, A. (eds.). ASHS Press, Alexandria, VA, pp. 95-96. 2007.
- Dierig, D.A., Dahlquist, G.H., Tomasi, P. 2006. Registration of wcl-l03 high oil lesquerella fendleri germplasm. *Crop Science*. 46(4):1832-1833.
- Dierig, D.A., Salywon, A.M., Rodriguez, D. 2006. Registration of a mutant lesquerella genetic stock for cream flower color. *Crop Science*. 46(4):1836-1837.
- Dierig, D.A., Thompson, A.E., Ray, D.T., Coffelt, T.A. 2006. Registration of three day-neutral flowering vernonia germplasms. *Crop Science*. 46(5):2339-2340
- Mozingo, R.W., Coffelt, T.A., Phipps, P.M., Coker, D.L. 2006. Registration of 'champs' peanut. *Crop Science* 46:2711-2712.

5348-21000-022-00D

ACQUISITION, EVALUATION AND CONSERVATION OF TEMPERATE FORAGE LEGUME GENETIC RESOURCES – Stephanie Greene (P) and Jinguo Hu; Prosser, Washington

- Boller, B., S.L. Greene. 2010. Genetic Resources. p. 13-38. /In/ B. Boller, U.K. Posselt, F. Veronesi (ed.) *Handbook of Plant Breeding- Vol. 5 Fodder and Amenity Grasses*, Springer, NY.
- Greene, S.L., Kisha, T.J., Dzyubenko, N.L. 2008. Conserving alfalfa wild relatives: is past introgression with Russian varieties evident today?. *Crop Science* 46:1853-1864.
- Nair, R., Hughes, S.J., Ellwood, S., Oliver, R., Greene, S.L., Delalande, M., Wen, J., Oldroyd, G. 2006. *Medicago truncatula* Stock Center. In: *Medicago Truncatula Handbook*. Available: <http://www.noble.org/MedicagoHandbook>
- Greene, S.L., Hughes, S.J., Nair, R., Hugué, T., Aouani, M.E., Prosperi, J.M., Delalande, M. 2006. Wild Accessions and Populations. In: *Medicago Truncatula Handbook*. Available: <http://www.noble.org/MedicagoHandbook>

5348-21000-023-00D

CHARACTERIZATION OF STRESS RESISTANCE GENES AND MECHANISMS, AND IMPROVEMENT AND GENOTYPING OF WHEAT AND BARLEY GERMPLASM FOR THE WESTERN U.S. – Camille Steber (P), Deven See, Daniel Skinner, and Kimberly Garland Campbell; Pullman, Washington

- Ariizumi, T., Lawrence, P.K., Steber, C.M. 2011. The Role of Two F-box Proteins, SLEEPY1 and SNEEZY, in Arabidopsis GA Signaling. *Plant Physiology*. Vol. 155, pp. 765-775.
- Baek, K., Skinner, D.Z. 2010. Molecular Cloning and Expression of Sequence Variants of Manganese Superoxide Dismutase Genes from Wheat. *Korean Journal of Environmental Agriculture*. 29(1):77-85.
- Schramm, E.C., Abellera, J.C., Strader, L., Garland Campbell, K.A., Steber, C.M. 2010. Isolation of ABA-responsive mutants in allohexaploid bread wheat (*Triticum aestivum* L.): Drawing connections to grain dormancy, preharvest sprouting, and drought tolerance. *Plant Science*. 2010.06.004
- Murphy, L.R., Santra, D., Kidwell, K.K., Yan, G.P., Chen, X., Garland Campbell, K.A. 2009. Linkage maps of wheat stripe rust resistance genes Yr5 and Yr15 for use in marker assisted selection. *Crop Sci*. 49:1-5.
- Kuraparthi, V., Sood, S., See, D.R., Gill, B.S. 2009. Development of a PCR Assay and Marker-Assisted Transfer of Leaf Rust and Stripe Rust Resistance Genes Lr57 and Yr40 into Hard Red Winter Wheats. *Crop Sci*. 49:120-126.
- Skinner, D.Z. 2009. Phospholipase A2 activity during cold acclimation of wheat. *Crop Sci*. 49:323-328. DOI:10.2135/cropsci2008.06.0359
- Zale, J., Agarwal, S., Loar, S., Steber, C.M. 2009. Evidence for stable transformation of wheat by floraldip in *Agrobacterium tumefaciens*. *Plant Cell Reports* 28(6):903-913. DOI 10.1007/s00299-009-0696-0.
- Ariizumi, T., Murase, K., Sun, T.P., Steber, C.M. 2008. Proteolysis-independent down-regulation of DELLA repression by the gibberellin receptor GID1. *The Plant Cell*, 20(9):2447-59.
- Leonard, J.M., Watson, C.J., Carter, A.H., Hansen, J.L., Zemetra, R.S., Santra, D.K., Garland Campbell, K.A., Riera-Lizarazu, O. 2008. Identification of a candidate gene for the wheat endopeptidase Ep-D1 locus and two other STS markers linked to the eyespot resistance gene Pch1. *Journal of Theoretical and Applied Genetics*. 116: 261-270.
- Zale, J., Agarwal, S., Loar, S., Steber, C.M. 2008. Floral Transformation of Wheat, Chapter 6, pp. 105-113. IN: Huw D. Jones and Peter R. Shewry (eds.) *Methods in Molecular Biology, Transgenic Wheat, Barley and Oats*. Humana Press, New York, NY.
- Schramm, E., Abellera, J., Strader, L., Garland Campbell, K.A., Steber, C.M. Can ABA Signaling be Used to Develop Drought Tolerant Wheat?. *Wheat Genetics International Symposium Proceedings*. Sydney Univ Press, O38, pg 1-3.
- Sherman, J.D., Souza, E.J., See, D.R., Talbert, L.E. 2008. Microsatellite markers for kernel color genes in wheat. *Crop Science*. 48:1419-1424.
- Walker, C., Garland Campbell, K.A., Carter, B., Kidwell, K. 2008. Using the solvent retention capacity test when breeding wheat for diverse production environments. *Crop Sci*. 48: 495-506.
- Finkelstein, R., Reeves, W., Ariizumi, T., Steber, C.M. 2008. Molecular aspects of seed dormancy. *Annu. Rev. Plant Biol*. 59:387-415.
- Skinner, D.Z., Garland Campbell, K.A. 2008. Evidence of a major genetic factor conditioning freezing sensitivity in winter wheat. *Plant Breeding* 127:228-234.
- Skinner, D.Z., Garland Campbell, K.A. 2008. The relationship of LT50 to prolonged freezing survival in winter wheat. *Can. J. Plant Sci*. 88: 885-889.
- Ariizumi, T., Steber, C.M. 2007. Seed germination of GA-insensitive sleepy1 mutants does not require RGL2 protein disappearance in Arabidopsis. *The Plant Cell* 19, 791-804.
- Jones, S.S., Lyon, S.R., Balow, K.A., Gollnick, M.A., Burns, J.W., Schillinger, W.F., Reisenauer, P.E., Murray, T.D., Chen, X., Garland Campbell, K.A., Morris, C.F., Goates, B. 2007. Registration of 'bauermeister' wheat. *Crop Sci*. 47:430-431.
- Murphy, K.M., Garland Campbell, K.A., Lyon, S.R., Jones, S.S. 2007. Evidence of varietal adaptation to organic farming systems. *Field Crops Research* 102:172-177.
- Murphy, L.R., Barroca, J., Franceschi, V.R., Lee, R., Roalson, E.H., Edwards, G.E., Ku, M.S. 2007. Diversity and plasticity of C4 photosynthesis in Eleocharis (Cyperaceae). *Functional Plant Biology* 34: 571-580.
- Jones, S.S., Lyon, S.R., Balow, K.A., Gollnick, M.A., Burns, J.W., Schillinger, W.F., Reisenauer, P.E., Murray, T.D., Chen, X., Garland Campbell, K.A., Morris, C.F., Goates, B. 2007. Registration of 'bauermeister' wheat. *Crop Sci*. 47:430-431.
- Jones, S.S., Lyon, S.R., Balow, K.A., Murray, T.D., Chen, X., Carter, B.P., Morris, C.F., Garland Campbell, K.A., Burns, J.W., Schillinger, W.F., Reisenauer, P.E., Goates, B. 2006. Registration of 'Masami' wheat. *Crop Sci*. 46:476-477.
- Skinner, D.Z. 2006. Plant abiotic stress. *Environmental Quality* 35:955A.

5348-21000-024-00D

COOL SEASON GRAIN LEGUME GENETIC ENHANCEMENT AND PATHOLOGY – George Vandemark (P), Kevin Mcphee, and Weidong Chen; Pullman, Washington

- Njambere, E., Vandemark, G.J., Chen, W. 2010. Development and characterization of microsatellite markers of the fungal plant pathogen *Sclerotinia trifoliorum*. *Genome*. 53:494-200.
- Vandemark, G.J., Porter, L. 2010. First report of root rot of lentil caused by *Aphanomyces euteiches* in Idaho. *Plant Disease*. 94:480.
- Vandemark, G.J., Ariss, J., Hughes, T. 2010. Real-time PCR suggests that *Aphanomyces euteiches* is associated with reduced amounts of *Phytophthora medicaginis* in alfalfa that is co-inoculated with both pathogens. *Journal of Phytopathology*. 158:117-124.
- Attanayake, R., Glawe, D., Dugan, F.M., Chen, W. 2009. *Erysiphe trifolii* causing powdery mildew of lentil (*Lens culinaris*). *Plant Disease*. 93:797-803.
- Attanayake, K., Glawe, D., Mcphee, K.E., Dugan, F.M., Chen, W. 2008. First report of powdery mildew of chickpea (*Cicer arietinum*) caused by *Leveillula taurica* in Washington State. *Plant Health Progress*. doi:10.1094/PHP-2008-0702-01-BR.
- Chilvers, M., Peever, T., Akamatsu, H., Chen, W., Muehlbauer, F.J. 2008. *Didymella rabiei* primary inoculum release from chickpea debris in relation to weather variables in the Pacific Northwest of the United States. *Canadian Journal of Plant Pathology*. 2007. 29: 4, 365-371. 33 ref.
- Njambere, E.N., Chen, W., Frate, C., Wu, B., Temple, S.R., Muehlbauer, F.J. 2008. Stem and Crown Rot of Chickpea in California Caused by *Sclerotinia trifoliorum*. *Plant Disease*. 92:917-922.
- Vandemark, G.J., Fourie, D., Miklas, P.N. 2008. Genotyping with Real-time PCR Reveals Recessive Epistasis Between Independent QTL Conferring Resistance to Common Bacterial Blight in Dry Bean. *Journal of Theoretical and Applied Genetics*. 117:513-522.
- Chen, W. 2008. Chickpea Diseases: Ecology and Control. *Encyclopedia of Pest Management*, 1:1,1-5.
- Chen, W., Mcphee, K.E., Muehlbauer, F.J. 2007. Field evaluation of fungicides for control of *Ascochyta* blight of chickpea, 2006.. *Plant Disease Management Reports*.
- Cutforth, H., McGinn, S., Mcphee, K.E., Miller, P. 2007. Adaptation of Pulse Crops to the Changing Climate of the Northern Great Plains. *American Society of Agronomy Meetings*. vol99 pg 1684-1699
- Mcphee, K.E., Chen, C., Wichman, D., Muehlbauer, F.J. 2007. Registration of 'Windham' Winter Feed Pea.. *Crop Science*..1(2):117-118.
- Mcphee, K.E., Croser, J., Sarmah, B., Ali, S.S., P.N., R., Zhang, H.B., Higgins, T.J. 2007. Development of Transgenics in Chickpea. In: S.S. Yadav, B. Redden, W. Chen and B. Sharma (Eds.) *Chickpea Breeding and Management*. CABI International, UK. pp. 458-473.
- P.N., R., Mcphee, K.E., Ford, R., Pittock, C., Kumar, J., Muehlbauer, F.J. 2007. Ciceromics: Advancement in Genomics and Recent Molecular Techniques. In: S.S. Yadav, B. Redden, W. Chen and B. Sharma (Eds.) *Chickpea Breeding and Management*. CABI International, UK. pp. 445-457.
- P.N., R., Muehlbauer, F.J., Mcphee, K.E. 2007. Stability analysis of chickpea large genomic DNA inserts in *Agrobacterium*.. *International Chickpea and Pigeonpea Newsletter*.online
- White, D., Chen, W. 2007. Towards identifying pathogenic determinants of the chickpea pathogen *Ascochyta rabiei*.. *European Journal of Plant Pathology*.
- Yadav, S.S., Longnecker, N., Dusunceli, F., Bejiga, M., Yadav, M., Rizi, A.H., Manahor, M., Reddy, A.A., Xaxiao, Z., Chen, W. 2007. Uses, consumption, and utilization. Book Chapter. p. 72-100.
- Chen, C., Miller, P., Muehlbauer, F.J., Neill, K., Wichman, D., Mcphee, K.E. 2006. Winter pea and lentil response to seeding date and micro- and macro-environments. *Agronomy Journal*. 98:1655-1663.
- Muehlbauer, F.J., Chen, W. 2007. Resistance to *ascochyta* blights of cool season food legumes.. *European Journal of Plant Pathology*.

5348-21000-026-00D

RESEARCH, ACQUISITION, MANAGEMENT, AND DOCUMENTATION OF PLANT GENETIC RESOURCES – Jinguo Hu (P), Frank Dugan, Stephen Clement, Richard Johnson, Clarice Coyne, Barbara Hellier, Molly Welsh, Theodore Kisha, and Vicki Bradley; Pullman, Washington

- Mallikarjuna, N., Coyne, C.J., Cho, S., Rynearson, S., Rajesh, P., Jadhav, D.R., Muehlbauer, F. 2011. *Cicer*. Wild Crop Relatives: Genomic and Breeding Resources. *Legume Crops and Forages*, pp. 63-82. Springer, New York, N.Y.
- Dugan, F.M. 2011. Three new host-fungus records for *Golovinomyces* species in Montana and Washington. *North American Fungi*. 6(3): 1-7.
- Dugan, F.M., Hellier, B.C., Lupien, S.L. 2011. Resistance to *Penicillium allii* in accessions from a National Plant Germplasm System *Allium* collection. *Crop Protection Journal*. 30: 483-488.

- Skoglund, L.G., Harveson, R.M., Chen, W., Dugan, F.M., Schwartz, H.F., Markell, S.G., Porter, L., Burrows, M.L., Goswami, R. 2011. Ascochyta Blight of Peas. Plant Health Progress.
<http://www.plantmanagementnetwork.org/sub/php/diagnosticguide/2011/pea/>.
- Johnson, R.C., Johnston, W.J., Bertoli, F.L., Golob, C. 2011. Seed yield, Development, and Variation in Diverse *Poa Pratensis* Accessions. Crop Science. 50:337-344.
- Kisha, T.J., Cramer, C.S. 2011. Determining Redundancy of Short-day, Onion Accessions in a Germplasm Collection Using Microsatellite and Targeted Region Amplified Polymorphic Markers. Journal of the American Society for Horticultural Science. vol. 136(2):1-6.2011.
- Bensch, K., Groenewald, J.Z., Dijksterhuis, J., Starink-Willemse, M., Anderson, B., Summerell, B.A., Shin, H.D., Dugan, F.M., Schroers, H.J., Braun, U., Crous, P.W. 2010. Species and ecological diversity within the *Cladosporium cladosporioides* complex (Davidiellaceae, Capnoidiales). Studies in Mycology 67: 1-94.
- Bradley, V.L. 2010. Isolation Distance, Inflorescence Sampling, and Population Size: Maintaining Genetic Diversity in the U.S. Temperate Grass Germplasm Collection. International Symposium on Forage, Turfgrass and Biofuel Germplasm Research p.100-103.
- Clement, S.L., Elbertson, L.R. 2010. Variable Effects of Grass-Neotyphodium Associations on Cereal Leaf Beetle (Coleoptera:Chrysomelidae) Feeding, Development and Survival. Journal of Entomological Science 45:1-7.
- Hellens, R.P., Moreau, C., Wang, K., Schwinn, K., Thomson, S., Fiers, M., Frew, T.J., Murray, S.R., Hofer, J., Jacobs, J., Davies, K.M., Allan, A.C., Bendahmane, A., Coyne, C.J., Timmerman, G.M., Ellis, N.T. 2010. Identification of Mendel's white flower character. PLoS One. 5 (10).
- Hu, J. 2010. Genetic Linkage Maps: Strategies, Resources and Achievements. Genetics, Genomics and Breeding Series: Sunflower. In: Hu, J., Seiler, G., Kole, C., Editors. Genetics, Genomics and Breeding of Sunflower. Enfield, NH: Science Publishers p 79-109.
- Hu, J., Seiler, G.J., Kole, C. 2010. Genetics, Genomics and Breeding in Sunflower. Genetics, Genomics and Breeding Series: Enfield, NH: Science Publishers pp 335.
- Kwon, S.J., Hu, J., Coyne, C.J. 2010. Genetic diversity and relationship among faba bean (*Vicia faba* L.) germplasm entries as revealed by TRAP markers. Plant Genetic Resources: Characterization and Utilization. 8(3); 204-213.
- Newcombe, G., Dugan, F.M. 2010. Fungal pathogens of plants in the Homogocene. Current Advances in Molecular Mycology. book chapter.
- Johnson, R.C., Erickson, V.J., Mandel, N.L., St Clair, J.B., Vance-Borland, K.W. 2010. Mapping genetic variation and seed zones for *Bromus carinatus* in the Blue Mountains of eastern Oregon, USA.. Botany 88: 725-736
- Smykal, P., Kenicer, G., Flavell, A.J., Kosterin, O., Redden, R.J., Ford, R., Coyne, C.J., Maxted, N., Ambrose, M.J., Ellis, T.N. 2010. Phylogeny, phylogeography and genetic diversity of *Pisum* genus. Plant Genetic Resources. 1-15 doi:10.1017/S147926211000033X.
- Clement, S.L., Huseby, D.S., Eigenbrode, S.D. 2010. Ecological Factors Influencing Pea Aphid Outbreaks in the U.S. Pacific Northwest. Book Chapter.
- Pecetti, L., Johnson, R.C., Romani, M., Bassignana, M., Marianna, D.G. 2010. Ecological characterisation of *supina* bluegrass (*Poa supina* Schrad.) germplasm from the Italian Alps. Grass and Forage Science. doi: 10.1111/j.1365-2494.2010.00766.x.
- Clement, S.L. 2009. Variable effects of fungal endophyte-infected grasses on the performance of pestiferous aphids. Redia Journal of Zoology. 92:205-209.
- Clement, S.L., Bradley, V.L., Elbertson, L.R., Bragg, D.E., Phillips, T.D. 2009. Cereal Leaf Beetle Colonizes Grass Germplasm Nurseries and Impacts Seed Production Activities. Online. Forage and Grazinglands. doi:10.1094/FG-2009-1214-01-RS.
- Clement, S.L., McPhee, K.E., Elbertson, L.R., Evans, M.A. 2009. Pea weevil, *Bruchus pisorum* (L.) (Coleoptera: Bruchidae), resistance in *Pisum sativum* x *P. fulvum* interspecific crosses. Plant Breeding. Online, doi:10.1111/j.1439-0523.2008.01603.x.
- Clement, S.L., Sharma, H.C., Muehlbauer, F.J., Elbertson, L.R., Mattinson, D.S., Fellman, J.K. 2009. Resistance to Beet Armyworm in a Chickpea Recombinant Inbred Line Population. Journal of Applied Entomology. Online, doi:10.1111/j.1439-0418.2009.01411.x.
- Clement, S.L., Smith, L.J., Prena, J., Kleene, M.D., Johnson, R.C. 2009. Non-Target Plant Use by a Weed Biocontrol Agent in Idaho: Host Expansion or Opportunistic Behavior? Biocontrol Science and Technology 19:455-461.
- Coyne, C.J., Brown, A.F., Timmerman-Vaughn, G. 2009. TaqMan assays exploring allelic diversity in USDA pea core collection. Plant and Animal Genome XVII Conference Abstract p. 150.
- Coyne, C.J., Diesso, D., Vandemark, G.J., Timmerman-Vaughan, G., Brown, A.F. 2009. Development of genetic resources for association mapping in pea. Meeting Proceedings. Model Legume Congress, Asilomar Conference Grounds, July 12-16, 2009.
- Coyne, C.J., Taylor, L.D., Varshney, R.K. 2009. Allelic diversity of USDA chickpea core collection of quantitative trait loci for resistance to ascochyta blight. Second International Ascochyta Workshop, June 29-July 2, 2009.

- Dugan, F.M. 2009. Dregs of our forgotten ancestors: fermentative microorganisms in the prehistory of Europe, the steppes and Indo-Iranian Asia, and their contemporary use in traditional and probiotic beverages. *Fungi* 2(4):16-39.
- Dugan, F.M., Akamatsu, H., Lupien, S.L., Chen, W., Chilvers, M., Peever, T.L. 2009. Ascochyta blight of chickpea reduced 38% by application of *Aureobasidium pullulans* (anamorphic Dothioraceae, Dothideales) to post harvest debris. *Biocontrol Science and Technology* 19:537-545.
- Dugan, F.M., Glawe, D.A., Attanayake, R.N., Chen, W. 2009. The Importance of Reporting New Host-Fungus Records for Ornamental and Regional Crops. APS Net Plant Pathology Online. Plant Health Progress doi:10.1094/PHP-2009-0512-01-RV.
- Dugan, F.M., Hellier, B.C. 2009. Managing diseases in seed garlic: What are the options?. Garlic Press. newsletter note.
- Hu, J., Mwengi, J.E., Coyne, C.J., Pan, W.L. 2009. First-year results of evaluating winter-hardiness of 55 faba bean (*Vicia faba* L.) accessions from the NPGS collection. *Pisum Genetics* 41:57-58.
- Furman, B.J., Coyne, C.J., Redden, B., Sharma, S.K., Vishnyakova, M. 2009. Chapter 6. Genetic Resources: Collection, Characterization, Conservation and Documentation. pages 64-75. In: *The Lentil: Botany, Production and Uses*, Erskine, W., Muehlbauer, F.J., Sarker, A. and Sharma, B. (eds.). CAB International, Oxfordshire, United Kingdom. Book Chapter.
- Murray, S., Falloon, L., Frew, T., Coyne, C.J., Butler, R., Timmerma-Vaughn, G. 2009. Association mapping of yield candidate gene homologs in a diverse collection of pea (*Pisum sativum* L.) lines. Meeting Proceedings. 19th Annual queenstown Molecular Biology Meetings 30 Aug-1st Sept., 2009.
- Newcombe, G., Gaylord, R., Yenish, J.P., Mastrogioseppe, J., Dugan, F.M. 2009. New Records for Pathogenic Fungi on Weedy or Non-Indigenous Plants. *North American Fungi* 4(8):1-12 doi: 10.2509/naf2009.004.008
- Nieves-Rivera, A.M., Santos-Flores, C.J., Dugan, F.M., Miller, T.E. 2009. Guanophilic fungi in three caves of southwestern Puerto Rico. *International Journal of Speleology* 38: 61-71.
- Robertson, N.L., Coyne, C.J. 2009. Evaluation of USDA *Lupinus* sp. collection for seed-borne potyviruses. *Plant Genetic Resources*. Published online 07 May 2009. doi:10.1017/S1479262109257923
- Smykal, P., Coyne, C.J., Ford, R., Redden, R., Hybl, M., Flavell, A.J., Warkentin, T., Burstin, J., Duc, G., Ambrose, M., Ellis, T., 2009. EFFORT TOWARDS A WORLD PEA (*Pisum sativum* L.) GERMPLASM CORE COLLECTION: The case for common markers and data compatibility. *Pisum Genetics* 40:11-14.
- Wang, F., Yue, B., Hu, J., Stewart, J.M., Zhang, J. 2009. A Target Region Amplified Polymorphism (TRAP) Marker for Fertility Restorer Gene Rf1 and Chromosomal Localization of Rf1 and Rf2 in Cotton 2009. *Crop Science* 49:1602-1608.
- Yue, B., Cai, X., Vick, B.A., Hu, J. 2009. Genetic diversity and relationships among 177 public sunflower inbred lines assessed by TRAP markers. *Crop Science*, 49:1242-1249 (2009).
- Dugan, F.M., Groenewald, J.Z., Crous, P.W., Braun, U. 2008. Morphological plasticity in *Cladosporium sphaerospermum*. *Persoonia: Molecular Phylogeny and Evolution of Fungi* 21:9-16.
- Johnson, D.A., Pimentel, G., Dugan, F.M. 2008. *Cladosporium herbarum* causes a leaf spot on *Caltha leptosepala* (marshmarigold) in western North America. Online Plant Health Progress doi:10.1094/PHP-2008-1121-01-RS.
- Johnson, R.C. 2008. Genebanks Pay Big Dividends to Agriculture, the Environment, and Human Welfare. Public Library of Science Biology. <http://biology.plosjournals.org/perlserv/?request=get-document&doi=10.1371/journal.pbio.0060148>(See URL).
- Johnson, R.C., Bradley, V.L., Kisha, T.J. 2008. Safflower Germplasm: Past, Present, and Future.. International Safflower Conference Proceedings.
- Clement, S.L., Martin, R.C., Dombrowski, J.E., Elbertson, L.R., Kynaston, M., Azevedo, M.D. 2008. Neotyphodium endophytes in tall fescue seed: infection frequencies after seed production and prolonged cold storage. *Seed Science and Technology* 36:710-720.
- Coyne, C.J., Pilet-Nayel, M.-L. 2008. Candidate genes associated with QTL controlling resistance to fusarium root rot in pea. Plant and Animal Genome VX Conference Abstracts. January 12-16, San Diego, CA.
- Coyne, C.J., Porter, L., Inglis, D.A., Grunwald, N.J., McPhee, K.E., Muehlbauer, F.J. 2008. Registration of W6 26740, W6 26743 and W6 26745 Green Pea Germplasm Resistant to Fusarium Root Rot. *Journal of Plant Registrations* (2008) Vol 2, No. 2, 137-139.
- Dugan, F.M. 2008. Fungi, Folkways and Fairy Tales: Mushrooms & Mildews in Stories, Remedies & Rituals, from Oberon to the Internet. *North American Fungi* 3 (7): 23-72. http://www.pnwfungi.org/pdf_files/manuscripts_volume_3/naf20087/naf200874.pdf
- Dugan, F.M. *Fungi in the Ancient World: How Mushrooms, Mildews, Molds and Yeast Shaped the Early Civilizations of Europe, the Mediterranean, and the Near East*. APS Press, St. Paul, MN.. 2008. Complete Book.
- Johnson, R.C., Dajue, L. 2008. Safflower winter survival and selection response relates to Fall Growth morphology and acclimation capacity. *Crop Science* 48:1872-1880.
- Johnson, R.C., Hopkins, A.A., Evans, M.A. 2008. Carbon Isotope Discrimination, Selection Response, and Forage Production of Tall Fescue in Contrasting Environments. *Crop Science* 48:10481054.

- Pappu, H., Hellier, B.C., Dugan, F.M. 2008. Evaluation of the National Plant Germplasm System's Garlic Collection for Seven Viruses. *Plant Health Progress* doi:10.1094/PHP-2008-0919-01-RS.
- Wise, K.A., Bradley, C.A., Pasche, J.S., Gudmestad, N.C., Dugan, F.M., Chen, W. 2008. Baseline Sensitivity of *Ascochyta rabiei* to Azoxystrobin, Pyraclostrobin, and Boscalid. *Plant Disease* 92:295-300.
- Sharma, H.C., Gowda, C.L., Stevenson, P.C., Ridsdill-Smith, T.J., Clement, S.L., Ranga Rao, G.V., Romies, J., Miles, M., Bouhssini, M. 2007. Host Plant Resistance and Insect Pest Management in Chickpea. *Chickpea Breeding and Management*. In: Yadav S.S., R.J. Redden, W. Chen, B. Sharma, editors. *Chickpea Breeding and Management*. CAB International, Wallingford, U.K., pp. 520-537.
- Varshney, R.K., Coyne, C.J., Swamy, P., Hoisington, D. 2007. Molecular identification of genetically distinct accessions in the USDA chickpea core collection. *Pisum Genetics* 39: 32-33.
- Clement, S.L., Elbersen, L.R., Kynaston, M. 2007. High *Neotyphodium* infection frequencies in tillers and seed of infected tall fescue plants, pp. 49-52. *Proceedings of the 6th International Symposium on Fungal Endophytes of Grasses* (A.J. Popay and E.R. Thom, eds.). New Zealand Grassland Association, Dunedin, New Zealand.
- Clement, S.L., Elbersen, L.R., Waldron, B.L., Quisenberry, S.S. 2007. Variable Performance of Bird Cherry-Oat Aphid on *Neotyphodium*-infected Wild Tall Fescue from Tunisia pgs. 337-340. *6th International Symposium on Fungal Endophytes of Grasses*.
- Clement, S.L., Hellier, B.C., Elbersen, L.R., Bradley, V.L., Johnson, R.C. 2007. Insect Pollination in the Regeneration of Germplasm at the Western Regional Plant Introduction Station. *Proceedings 9th International Pollination Symposium on Plant-Pollinator Relationships, Diversity in Action*. Iowa State University, Ames, Iowa. p. 62-63.
- Clement, S.L., Hellier, B.C., Elbersen, L.R., Staska, R.T., Evans, M.A. 2007. Flies (Diptera: Muscidae, Calliphoridae) are efficient pollinators of *Allium ampeloprasum* L. (Alliaceae) in field cages. *Journal of Economic Entomology*. 100(1):131-135.
- Coyne, C.J., McClendon, M.T., Walling, J.G., Timmerman-Vaughn, G.M., Murray, S., Meksem, K., Lightfoot, D.A., Shultz, J.L., Keller, K.E., Martin, R.R., Inglis, D.A., Rajesh, P.N., McPhee, K.E., Weeden, N.F., Grusak, M.A., Storlie, E.W. 2007. Construction and characterization of two bacterial artificial chromosome libraries of pea (*Pisum sativum* L.) for the isolation of economically important genes. *Genome* 50:871-875.
- Dugan, F.M. 2007. Blastomycetes. In: In: McGraw-Hill Book Company editor. *McGraw-Hill Encyclopedia of Science & Technology*, 10th edition. New York, NY: McGraw-Hill. p. 154-155.
- Dugan, F.M. 2007. Diseases and Disease Management in Seed Garlic: Problems and Prospects. *The Americas Journal of Plant Science and Biotechnology*. 1: 47-51.
- Dugan, F.M. 2007. Hyphomycetes. In: McGraw-Hill Book Company editor. *McGraw-Hill Encyclopedia of Science & Technology*, 10th edition. New York, NY: McGraw-Hill. p. 780-782.
- Dugan, F.M., Glawe, D.A. 2007. Powdery mildews on weeds in the Pacific Northwest: a miscellany of new records. *Pacific Northwest Fungi*. 2(1):1-7. DOI: 10.2509/pnwf.2007.002.001
- Dugan, F.M., Hellier, B.C., Lupien, S.L. 2007. Pathogenic fungi in garlic seed cloves from the United States and China, and efficacy of fungicides against pathogens in garlic germplasm in Washington State.. *Journal of Phytopathology*. 155: 437-45.
- Dugan, F.M., Newcombe, G. 2007. New records for powdery mildews and *Taphrina* spp. in Idaho and Washington. *Pacific Northwest Fungi* 2(8): 1-5, doi: 10.2509/pnwf.2007.002.008..
- Dugan, F.M., Rector, B.G. 2007. Mycoflora of seed of common teasel (*Dipsacus fullonum*) in Washington State. *Pacific Northwest Fungi* 2 (6):1-10.
- Johnson, R.C., Kisha, T.J., Evans, M.A. 2007. Characterizing Safflower Germplasm with AFLP Molecular Markers. *Crop Science*. 47:1728-1736.
- Mallikarjuna, N., McGrew, S.L., Ryneerson, S., Rajest, P.N., Coyne, C.J., Muehlbauer, F.J. 2007. Pollen as a means of international transfer of germplasm. *Journal of Semi-Arid Tropical Agricultural Research (eJournal of J.SAT Agric. Res.* 3(1):www.icrisat.org/Journal/chickpea_pigeonpea3.htm.
- Redden, B., Furman, B., Upadhyaya, H.D., Pundir, R.S., Gowda, C.L., Coyne, C.J., Enneking, D. 2007. Biodiversity Management. In: Yadav S.S., R.J. Redden, W. Chen, B. Sharma, editors. *Chickpea Breeding and Management*. p. 355-368.
- Redden, B., Maxted, N., Furman, B., Coyne, C.J. 2007. Lens Biodiversity, p. 11-22.in: SS Yadav, D. McNeil, P.C. Stevenson (eds). *Lentil: An Ancient Crop for Modern Times*.
- Sharma, H.C., Clement, S.L., Ridsdill-Smith, T.J., Ranga Rao, G.V., El Bouhssini, M., Ujagir, R., Srivastava, C.P., Miles, M. 2007. *Proceedings Fourth International Food Legume Research Conference*, Indian Society of Genetics and Plant Breeding, New Delhi, India, pp. 1-23.
- Tadych, M., Bergen, M., Dugan, F.M., White, J.F. 2007. The potential role of water in spread of conidia of the *Neotyphodium* endophyte of *Poa annua*. *Mycological Research*. 111: 466-472.
- Babcock, C., Chen, X., Crous, P.W., Dugan, F.M., Goates, B., Green, P.N. 2007. *Plant Germplasm Centers and Microbial Culture Collections: A User's Guide to Key Genetic Resources for Plant Pathology*. *Plant Disease*. 91: 476-484.
- Dugan, F.M. 2007. Agonomycetes. In: McGraw-Hill Encyclopedia of Science & Technology, 10th edition. p.214-215

- Upadhyaya, H.D., Coyne, C.J., Singh, S., Gowda, C., Lalitha, N., Muehlbauer, F.J. 2006. Identification of Large-Seeded High Yielding Diverse Kabuli Accessions in Newly Assembled Chickpea Germplasm. *International Chickpea and Pigeonpea Newsletter*.13:2-5
- Johnson, R.C., Li, D. Registration of WSRC01, WSRC02, and WSRC03 Winter Hardy Safflower Germplasm. *Journal of Plant Registrations* 2:140-142.
- Mimura, M., Coyne, C.J., Bambuck, M.W., Lumpkin, T.A. 2006. Ssr diversity of vegetable soybean [glycine max (l.) merr.]. *Genetic Resources and Crop Evolution*. DOI: 10.1007/s10722-006-0006-4.
- Clement, S.L. 2006. Pea Aphid Outbreaks and Virus Epidemics on Peas in the U.S. Pacific Northwest: Histories, Mysteries, and Challenges. *Plant Health Progress*. doi:10.1094/PHP-2006-1018-01-RV.
- Dugan, F.M., Glawe, D.A. 2006. First report of powdery mildew on *dipsacus sylvestris* caused by *sphaerotheca dipsacearum* in north america. *Plant Health Progress*. doi:10.1094/PHP-2006-0607-02-BR.
- Glawe, D.A., Dugan, F.M. 2006. First report of Erysiphe (Uncinuliella) flexuosa in western North America. *Pacific Northwest Fungi*. 1(11):1-11. DOI 10.2509/pnwf.2006.001.011

5348-21430-004-00D

GENETIC IMPROVEMENT OF FREEZING TOLERANCE IN WINTER WHEAT – Daniel Skinner (P); Pullman, Washington

- Skinner, D.Z., Bellinger, B.S. 2010. Differential response of wheat cultivars to components of the freezing process in saturated soil. *Crop Science*. Vol. 51 No. 1, page 69-74 (2011).
- Skinner, D.Z., Bellinger, B.S. 2010. Exposure to subfreezing temperature and a freeze-thaw cycle affect freezing tolerance of winter wheat in saturated soil. *Plant and Soil*. 332(1):289.
- Baek, K., Skinner, D.Z. 2010. Molecular Cloning and Expression of Sequence Variants of Manganese Superoxide Dismutase Genes from Wheat. *Korean Journal of Environmental Agriculture*. 29(1):77-85.
- Skinner, D.Z. 2009. Post-acclimation transcriptome adjustment is a major factor in freezing tolerance of winter wheat. *Funct Integr Genomics*. 9:513–523. DOI 10.1007/s10142-009-0126-y.
- Skinner, D.Z., Mackey, B.E. 2009. Freezing tolerance of winter wheat plants frozen in saturated soil. *Field Crops Research* 113:335-341.
- Skinner, D.Z. 2009. Phospholipase A2 activity during cold acclimation of wheat. *Crop Sci*. 49:323-328. DOI:10.2135/cropsci2008.06.0359.
- Skinner, D.Z., Garland Campbell, K.A. 2008. Evidence of a major genetic factor conditioning freezing sensitivity in winter wheat. *Plant Breeding* 127:228-234.
- Skinner, D.Z., Garland Campbell, K.A. 2008. The relationship of LT50 to prolonged freezing survival in winter wheat. *Can. J. Plant Sci*. 88: 885-889.

5354-21220-010-00D

POTATO GERmplasm ENHANCEMENT THROUGH TRAIT DISCOVERY, GENETIC EVALUATION AND INCORPORATION – Duroy Navarre (P), James Crosslin, and Charles Brown; Prosser, Washington

- Brown, C. R., K. G. Haynes, M. Moore, M. J. Pavék, D. C. Hane, S. L. Love, R. G. Novy, and J. C. Miller, Jr. 2011. Stability and broad-sense heritability of mineral content in potato: Zinc. *American Journal of Potato Research*. 88:238-244
- Brown, C.R., Haynes, K.G., Moore, M., Pavék, M.J., Hane, D.C., Love, S.L., Novy, R.G., Miller, Jr, J.C. 2010. Stability and Broad-sense Heritability of Mineral Content in Potato: Iron. *American Journal of Potato Research*. 87: 390–396.
- Crosslin, J., Eastwell, K.C., Davitt, C.M., Abad, J.A. 2010. First report of seed-borne cherry leaf roll virus in wild potato, *Solanum acaule*, from South America. *Plant Disease*. 94:782.
- Crosslin, J., Hamlin, L. 2010. First report of Impatiens necrotic spot virus infecting greenhouse-grown potatoes in Washington State. *Plant Disease*. 94:1507.
- Crosslin, J., Munyaneza, J.E., Brown, J.K., Liefting, L. 2010. Potato zebra chip disease: a phytopathological tale. *Plant Health Progress*. http://www.apsnet.org/online/feature/zebra/PlantHealthProgress_Zebra.pdf.
- Navarre, D.A., Shakyia, R., Holden, M., Kumar, S. 2010. The Effect of Different Cooking Methods on Phenolics and Vitamin C in Developmentally Young Potato Tubers. *American Journal of Potato Research*. 87:350-359.
- Zhang, L., Brown, C.R., Culley, D., Baker, B.J., Kunibe, E., Denny, H., Smith, C., Ward, N., Beavert, T., Coburn, J., Pavék, J.J., Dauenhauer, N., Dauenhauer, R. 2010. Inferred origin of several Native American potatoes from the Pacific Northwest using SSR markers. *Euphytica*. 174:15-29.
- Brown, C.R., Mojtahedi, H., Crosslin, J., James, S., Charlton, B., Novy, R., Love, S., Vales, M.I., Hamm, P. 2009. Characterization of Resistance to Corky Ringspot Disease in Potato: A Case for Resistance to Infection by Tobacco Rattle Virus. *American Journal of Potato Research*. 86:49–55

- Brown, C.R., Mojtahedi, H., Zhang, L.H., Riga, E. 2009. Independent Resistant Reactions Expressed in Root and Tuber of Potato Breeding Lines with Introgressed Resistance to *Meloidogyne chitwoodi*. *Phytopathology*. 99:1085-1089.
- Nitzan, N., Boydston, R.A., Batchelor, D., Crosslin, J., Hamlin, L., Brown, C.R. 2009. Hairy Nightshade is an Alternative Host of *Spongospora subterranea*, the Potato Powdery Scab Pathogen. *American Journal of Potato Research*. 86:297-303.
- Goyer, A., Navarre, D.A. 2009. Vitamin B9 is Higher in Developmentally Younger Potato Tubers. *Journal of the Science of Food and Agriculture*. 89:579-583.
- Crosslin, J., Mallik, I., Gudmestad, N.C. 2009. First report of tomato spotted wilt virus causing potato tuber necrosis in Texas. *Plant Disease*. 93(8):845.
- Crosslin, J., Munyaneza, J.E. 2009. Evidence that the Zebra Chip Disease and the Putative Causal Agent Can be Maintained in Potatoes by Grafting and In Vitro. *American Journal of Potato Research*. 86:183-187.
- Arif, M., Thomas, P.E., Crosslin, J., Brown, C.R. 2009. Development of molecular resistance in potato against potato leaf roll virus and potato virus Y through *Agrobacterium*-mediated double transgenesis. *Pakistan Journal of Botany*. 41:945-954.
- Crosslin, J. 2009. Detection of tobacco rattle virus RNA in processed potato chips displaying symptoms of corky ringspot disease. *HortScience*. 44:1790-1791.
- Crosslin, J., Bester, G. 2009. First report of *Candidatus Liberibacter psyllaureus* in zebra chip symptomatic potatoes from California. *Plant Disease*. 93:551.
- Navarre, D.A., Goyer, A., Shakya, R. 2009. Developing the Nutritional Potential of Potato. In : Nigel Yee and William Bussell (Eds) *Potato IIII. Global Science Books. Food 3 (Special Issue 1)*, 118-124.
- Navarre, D.A., Shakya, R., Holden, M., Crosslin, J. 2009. LC-MS Analysis of Phenolic Compounds in Tubers Showing Zebra Chip Symptoms. *American Journal of Potato Research*. 86:88-95.
- Ottoman, R.J., Hane, D.C., Brown, C.R., Yilma, S., James, S.R., Mosley, A.R., Crosslin, J., Vales, M.I. 2009. Validation and Implementation of Marker-Assisted Selection (MAS) for PVY Resistance (Ryadg gene) in a Tetraploid Potato Breeding Program. *American Journal of Potato Research*. 86:304-314.
- Rondon, S.I., Hane, D., Brown, C.R., Vales, M.I., Dogramaci, M. 2009. Resistance of potato germplasm to the potato tuberworm (Lepidoptera: Gelechiidae). *Journal of Economic Entomology*. 102:1649-1653.
- Venugopal, S.C., Jeong, R., Chandra-Shekara, A.C., Zhu, S., Mandal, M., Hersh, M., Stromberg, A.J., Navarre, D.A., Kachroo, A., Kachroo, P. 2009. ENHANCED DISEASE SUSCEPTIBILITY 1 and SALICYLIC ACID act redundantly to regulate resistance gene-mediated signaling. *PLoS Genetics*. 5(7): e1000545
DOI:10.1371/journal.pgen.1000545.
- Xia, Y., Gao, Q., Yu, K., Lapchuk, L., Navarre, D.A., Hildebrand, D., Kachroo, A., Kachroo, P. 2009. An intact cuticle in distal tissues is essential for the induction of systemic acquired resistance in plants. *Cell Host and Microbe*. 5:151-65.
- Arif, M., Thomas, P.E., Crosslin, J., Brown, C.R. 2009. *Agrobacterium*-mediated transformation of potato using PLRV-rep and PVY CP genes and assessment of replicase mediated resistance against natural infection of PLRV. *Pakistan Journal of Botany*. 41:1477-1488.
- Nitzan, N., Evans, M.A., Cummings, T.F., Johnson, D.A., Batchelor, D.L., Olsen, C., Brown, C.R. 2009. Field Resistance to Potato Stem Colonization by the Black Dot Pathogen *Colletotrichum coccodes*. *Plant Disease*. 93(11):1116-1122.
- Nitzan, N., Cummings, T., Johnson, D., Miller, J., Batchelor, D., Olsen, C., Brown, C.R. 2008. Resistance to Root Gallings Caused by the Powdery Scab Pathogen *Spongospora subterranea* in Potato. *Plant Disease*. 92:1643-1649.
- Gudmestad, N.C., I. Mallik, J.S. Pasche, and J.M. Crosslin. 2008. First report of tobacco rattle virus causing corky ringspot in potatoes grown in Minnesota and Wisconsin. *Plant Disease* 92:1254.
- Brown, C.R. 2008. Variability of Phytonutrient Content of Potato In Relation to Growing Location and Cooking Method. *Potato Research*. 51:259-270.
- Chanda, B., Venugopal, S.C., Kulshrestha, S., Navarre, D.A., Downie, B., Vaillancourt, L., Kachroo, A., Kachroo, P. 2008. Glycerol-3-Phosphate Levels Are Associated with Basal Resistance to the Hemibiotrophic Fungus *Colletotrichum higginsianum* in *Arabidopsis*. *Plant Physiology*. 147(4): 2017-2029.
- Kachroo, A., Da-Qi, F., Havens, W., Navarre, D.A., Kachroo, P., Ghabrial, S.A. 2008. An oleic acid-mediated pathway induces constitutive defense signaling and enhanced resistance to multiple pathogens in soybean. *Molecular Plant-Microbe Interactions*. 21:564-575.
- Kirk, W.W., S.L. Gieck, J.M. Crosslin, and P.B. Hamm. 2008. First report of corky ringspot caused by Tobacco rattle virus on potatoes (*Solanum tuberosum*) in Michigan. *Plant Disease* 92:485.
- Nchimbi-Msolla, S., Misangu, R., Mabagala, R., Magayane, F., Kweka, S., Butler, L.M., Miklas, P.N., Myers, J.R. 2008. Mshindi Kablanketi Dry Bean for East Africa. *Bean Improvement Cooperative Annual Report*. 51:278-279.

- Rommens, C.M., Richael, C.M., Yan, H., Navarre, D.A., Ye, J., Krucker, M., Swords, K. 2008. Engineered Native Pathways for High Kaempferol and Caffeoylquinic Production in Potato. *Plant Biotechnology Journal*. DOI: 10.1111/j.1467-7652.2008.00362. 6:870-886.
- Shakya, R., Navarre, D.A. 2008. Lc-ms analysis of solanidane glycoalkaloid diversity among four wild potato species and three cultivars. *Journal of Agricultural and Food Chemistry*. DOI 10.1021/jf8006618. 56:6949-58.
- Bhattarai, K.K., Xie, Q., Mantelin, S., Bishnoi, U., Girke, T., Navarre, D.A., Kaloshian, I. 2008. Tomato susceptibility to root-knot nematodes requires an intact jasmonic Acid signaling pathway. *Molecular Plant-Microbe Interactions*. 21(9):1205-14.
- Brown, C.R. 2007. Breeding for Phytonutrient Enhancement of Potato. *American Journal of Potato Research*. 85:298-307. DOI 10.1007/s12230-008-9028-0.
- Zhang, L., Mojtahedi, H., Kuang, H., Baker, B.J., Brown, C.R. 2007. The Use of STS Markers in the Marker-assisted Selection of Columbia Root-knot Nematode Resistance Introgressed from *Solanum bulbocastanum*. *Crop Science*. 47:2021-2026.
- Nitzan, N., Johnson, D., Batchelor, D., Brown, C.R. 2007. Powdery Scab. *Potato Progress*. July/August 2007, pp 6-7.
- Akamatsu, H. O., Grünwald, N. J., Chilvers, M. I., Porter, L. D., and Peever, T. L. 2007. Development of codominant simple sequence repeat, single nucleotide polymorphism and sequence characterized amplified region markers for the pea root rot pathogen, *Aphanomyces euteiches*. *Journal of Microbiological Methods* 71: 82-86.
- Mojtahedi, H., Boydston, R.A., Crosslin, J., Brown, C.R., Riga, E., Anderson, T.L., Spellman, D.E., Quick, R.A. 2007. Establishing a Corky Ringspot Disease Plot for Research Purposes. *Journal of Nematology*. 39(4):313-316.
- Gieck, S.L., David, N.L., Hamm, P.B., Crosslin, J., Ingham, R.E. 2007. Delayed emergence, stem distortion, stunting, and foliar symptoms associated with Tobacco Rattle Virus and *Paratrichodorus allius* in potatoes grown in the Pacific Northwest. Online. *Plant Health Progress* doi: 10.1094/PHP-2007-0917-01-BR.
- Goyer, A., Navarre, D.A. 2007. Determination of folate concentrations in diverse potato germplasm using a trienzyme extraction and microbiological assay. *Journal of Agricultural and Food Chemistry*. 55(9):3523-3528.
- "Inglis, D.A., Brown, C.R., Gundersen, B.G., Porter, L., Miller, J.S., Johnson, D.A., Lozoya-Saldana, H., Haynes, K.G. 2007. Assessment of *Solanum hougasii* in Washington and Mexico as a Source of Resistance to Late Blight. *American Journal of Potato Research*. Vol 54: 217-228."
- Moeder, W., Del Pozo, O., Navarre, D.A., Martin, G.B., Klessig, D.F. 2007. Plant aconitase functions as an rna-binding protein and plays a role in regulating resistance to oxidative stress and hypersensitive cell death. *Plant Molecular Biology*. 63(2): 273-287.
- Joyce, P.A., Constable, F.E., Crosslin, J., Eastwell, K., Howell, W.E., Rodoni, B.C. 2006. Characterization of Pear Blister Canker Viroid Isolates from Australian Pome Fruit Orchards. *Australasian Plant Pathology Society*. 35:465-471.
- Lorenzen, J.H., Meacham, T., Berger, P.H., Shiel, P.J., Crosslin, J., Hamm, P.B., Kopp, H. 2006. Whole genome characterization of Potato virus Y isolates collected in the western USA and their comparison to isolates from Europe and Canada. *Archives of Virology*. 151:1055-1074.

5354-21220-016-00D

DEVELOP STRESS-RESISTANT DRY BEAN GERMLASM AND SUSTAINABLE PEST MANAGEMENT STRATEGIES FOR EDIBLE LEGUMES – Phillip Miklas (P), Lydon Porter, and Richard Porter; Prosser, Washington

- Bassett, M., Miklas, P.N., Caldas, G.V., Blair, M.W. 2010. A dominant gene for garnet brown seed coats at the Rk locus in Dorado common bean and mapping Rk to linkage group 1. *Euphytica*. 176:281–290.
- Talukder, Z.I., Anderson, E., Miklas, P.N., Blair, M.W., Osorno, J., Dilawari, M., Hossain, K.G. 2010. Genetic diversity and selection of genotypes to enhance Zn and Fe content in common bean. *Canadian Journal of Plant Science*. 90: 49-60.
- Pascual, A., Campa, A., Pérez-Vega, E., Giraldez, R., Miklas, P.N., Ferreira, J. 2010. Screening common bean for resistance to four *Sclerotinia sclerotiorum* isolates collected in northern Spain. *Plant Disease*. 94: 885-890.
- Miklas, P.N., Porch Clay, T.G. 2010. Guidelines for Common Bean QTL Nomenclature. *Annual Report of the Bean Improvement Cooperative* 53:202-204.
- Bassett, M.J., Miklas, P.N. 2009. Blue Pattern Flower in Common Bean Expressed by Interaction of Prpi-2 with a New Gene tbp. *Journal of the American Society for Horticultural Science*. 134: 423-427.
- Miklas, P.N., Seo, Y.S., Gilbertson, R.L. 2009. Quantitative Resistance to Bean dwarf mosaic virus in Common Bean is Associated with the Bct gene for Resistance to Beet curly top virus. *Plant Disease*. 93:645-648.
- Porter, L., Hamm, P., David, N., Gieck, S., Miller, J. 2009. Metalaxyl-M-Resistant *Pythium* Species in Potato Cropping Systems in the Pacific Northwest, USA. *Plant Disease*. Published online DOI 10. A 1007/s 12230-009-0985-z.

- Miklas, P.N., Fourie, D., Wagner, J., Larsen, R.C., Mienie, C. 2009. Tagging and mapping Pse-1 gene for resistance to halo blight in common bean differential cultivar UI-3. *Crop Science*. 49:41-48.
- Mutlu, N., Urrea, C.A., Miklas, P.N., Steadman, J.R., Pastor Corrales, M.A., Lindgren, D.T., Reiser, J., Vidaver, A.K., Coyne, D.P. 2008. Registration of Common Bacterial Blight, Rust and Bean Common Mosaic Resistant Great Northern Bean Germplasm Line ABC - Weihing. *Journal of Plant Registrations*. 2:120-124.
- Nchimbi-Msolla, S., Misangu, R., Mabagala, R., Magayane, F., Kweka, S., Butler, L.M., Miklas, P.N., Myers, J.R. 2008. Pesa Large Red Dry Bean. *Bean Improvement Cooperative Annual Report*. 51:280-281.
- Porter, L., Coffman, V.A. 2008. Assessment of organic seed treatments to manage seed and root rot on peas, 2007. *Plant Disease Management Reports (online)*. Report 2:V153. DOI:10.1094/PDMR02. The American Phytopathological Society, St. Paul, MN.
- Porter, L., Coffman, V.A. 2008. Efficacy of seed treatments in reducing seed and root rot of peas in the presence of metalaxyl-resistant *Pythium*, 2007. *Plant Disease Management Reports (online)*. Report 2:V154. DOI:10.1094/PDMR02. The American Phytopathological Society, St. Paul, MN.
- Porter, L., Hoheisel, G., Coffman, V.A. 2008. Resistance of Peas to *Sclerotinia sclerotiorum* in the *Pisum* Core Collection. *Plant Pathology*. 58:52-60.
- Riga, E., Porter, L.D., Mojtahedi, H., and Brocke, G.F. 2008. First report of *Pratylenchus neglectus*, *Pratylenchus thornei* and *Paratylenchus hametis* nematodes causing yield reduction to dry land peas and lentils in Idaho. *Plant Disease* 92: 979.
- Weeden, N.F., Porter, L. 2008. The Genetic Basis of *Fusarium* Root Rot Tolerance in the 'Afghanistan' Pea. *Pisum Genetics*. 39: 35-36.
- Singh, S. P., H. Teran, M. Lema, H.F. Schwartz, and P. N. Miklas. 2007. Registration of white mold resistant dry bean germplasm line A 195. *Journal of Plant Registrations* 1:62-63.
- Porter, L.D., Johnson, D.A. 2007. Survival of Sporangia of New Clonal Lineages of *Phytophthora infestans* in Soil under Semiarid Conditions. *Plant Disease*. 91: 835-841.
- Osorno, J. M., C. G. Muñoz, J. S. Beaver, F. H. Ferwerda, M. J. Bassett, P. N. Miklas, T. Olczyk, and B. Bussey. 2007. Two genes from *Phaseolus coccineus* L. confer resistance to Bean golden yellow mosaic virus in common bean. *J. Amer. Soc. Hort. Sci.* 132:530-533.
- Miklas, P.N. 2007. Marker-assisted backcrossing QTL for partial resistance to *Sclerotinia* white mold in dry bean. *Crop Science*. 47:935-942.
- Miklas, P.N., Larsen, K.M., Terpstra, K.A., Hauf, D.C., Grafton, K.F., Kelly, J.D. 2007. Qtl analysis of ica bunsii-derived resistance to white mold in a pinto x navy bean cross. *Crop Science*. 47: 174-179.
- Bassil, N.V., Gilmore, B.S., Oliphant, J.M., Hummer, K.E., Henning, J.A. 2007. Genic SSRs for European and North American Hop (*Humulus lupulus* L.). *Genetic Resources and Crop Evolution*. Available: www.springerlink.com/content/d000k04454n55v2p/fulltext.pdf
- Bassett, M.J., Miklas, P.N. 2007. A new gene, bic, with pleiotropic effects (with T P V) for bicolor flowers and dark olive brown seed coat in common bean. *Journal of the American Society for Horticultural Science*. 132: 352-356.
- Akamatsu, H. O., Grünwald, N. J., Chilvers, M. I., Porter, L. D., and Peever, T. L. 2007. Development of codominant simple sequence repeat, single nucleotide polymorphism and sequence characterized amplified region markers for the pea root rot pathogen, *Aphanomyces euteiches*. *Journal of Microbiological Methods* 71: 82-86.
- Miklas, P.N., Grafton, K.F., Hauf, D., Kelly, J.D. 2006. Registration of Partial White Mold Resistant Pinto Bean Germplasm Line USPT-WM-1. *Crop Science*.46:2339
- Miklas, P.N., Smith, J.R., Singh, S.P. 2006. Registration of Common Bacterial Blight Resistant White Kidney Bean Germplasm Line SWK-CBB-17. *Crop Science*. 46:2338-2339
- Hang, A.N., Miklas, P.N., Silbernagel, M.J., Hosfield, G.L. 2006. Registration of 'silver cloud' white kidney dry bean. *Crop Science*. 46:991-992.
- Hang, A.N., Miklas, P.N., Silbernagel, M.J., Hosfield, G.L. 2006. Registration of 'quincy' pinto bean. *Crop Science*.46:991.

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PHYSIOLOGY AND GENETIC IMPROVEMENT OF SMALL FRUIT CROPS – David Bryla (P) and Chad Finn;
Corvallis, Oregon

- Finn, C.E., Strik, B.C., Yorgey, B.M., Martin, R.R. 2011. 'Onyx' trailing blackberry. *HortScience*. 46:657-659.
- Bryla, D.R., Gartung, J.L., Strik, B. 2011. Evaluation of irrigation methods for highbush blueberry. I. Growth and water requirements of young plants. *HortScience*. 46(1):95-101.
- Bryla, D.R., Machado, R., Shireman, A.D. 2010. Effects of method and level of nitrogen fertilizer application on soil pH, electrical conductivity, and availability of ammonium and nitrate in blueberry. *Acta Horticulturae*. 868:95-101.

- Bryla, D.R., Trout, T.J., Ayars, J.E. 2010. Weighing lysimeters for developing crop coefficients and efficient irrigation practices for vegetable crops. *HortScience*. 45(11):1597-1604.
- Charlebois, D., Byers, P., Finn, C.E., Thomas, A.L. 2010. Elderberry: botany, horticulture, potential. *Horticultural Reviews*. 37:213-280.
- Clark, J.R., Finn, C.E., editors. 2010. Register of New Fruit and Nut Cultivars List 45. *HortScience*. 45:716-756.
- Dossett, M., Finn, C.E. 2010. Identification of resistance to the large raspberry aphid in black raspberry. *Journal of the American Society for Horticultural Science*. 135(5):438-444.
- Du, X., Finn, C.E., Qian, M. 2010. Bound volatile precursors in genotypes in the pedigree of 'Marion' Blackberry (*Rubus* Sp.). *Journal of Agricultural and Food Chemistry*. 58:3694-3699.
- Du, X., Finn, C.E., Qian, M. 2010. Distribution of volatile composition in 'Marion' (*Rubus* Species Hyb) blackberry pedigree. *Journal of Agricultural and Food Chemistry*. 1860-1869.
- Du, X., Finn, C.E., Qian, M. 2010. Volatile composition and odour-activity value of thornless 'Black Diamond' and 'Marion' blackberries. *Food Chemistry*. 119:1127-1134.
- Du, X., Kurnianta, A., Mcdaniel, M., Finn, C.E., Qian, M.C. 2010. Flavour profiling of 'Marion' and thornless blackberries by instrumental and sensory analysis. *Food Chemistry*. 121:1080-1088.
- Finn, C.E., Clark, J.R., Sleezer, S. 2010. Blackberry. *HortScience*. 45:720-721.
- Finn, C.E., Strik, B.C., Yorgey, B., Martin, R.R., Stahler, M. 2010. 'Newberry' trailing blackberry. *HortScience*. 45(3):437-440.
- Finn, C.E., Strik, B.C., Yorgey, B., Qian, M., Martin, R.R., Peterson, M.E. 2010. 'Wild Treasure' thornless trailing blackberry. *HortScience*. 45(3):434-436.
- Hancock, J.F., Finn, C.E., Dale, A., Luby, J.J., Callow, P.W., Serce, S. 2010. Reconstruction of the strawberry, *Fragaria* × *ananassa*, using native genotypes of *F. virginiana* and *F. chiloensis*. *HortScience*. 45:1006-1013.
- Bassil, N.V., Bunch Jr, T.R., Nyberg, A.M., Zee, F.T., Hummer, K.E. 2010. Microsatellite Markers Distinguish Hawaiian Ohelo from Other *Vaccinium* L. Section *Myrtillus* Species. *Acta Horticulturae*. 859:81-88.
- Hancock, J.F., Finn, C.E., Dale, A., Luby, J.J., Callow, P.W., Serce, S. 2010. Reconstruction of the strawberry, *Fragaria* × *ananassa*, using native genotypes of *F. virginiana* and *F. chiloensis*. *HortScience*. 45:1006-1013.
- Dossett, M., Finn, C.E. 2010. Identification of resistance to the large raspberry aphid in black raspberry. *Journal of the American Society for Horticultural Science*. 135(5):438-444.
- Charlebois, D., Byers, P., Finn, C.E., Thomas, A.L. 2010. Elderberry: botany, horticulture, potential. *Horticultural Reviews*. 37:213-280.
- Finn, C.E., Moore, P.P., Kempler, C., Yorgey, B., Strik, B., Martin, R.R. 2009. 'Valley Red' Strawberry. *HortScience*. 44(5):1468-1471.
- Bryla, D.R., Yorgey, B., Shireman, A.D. 2009. Irrigation management effects on yield and fruit quality of highbush blueberry. *Acta Horticulturae*. 810:649-656.
- "Thompson, E., Strik, B., Finn, C.E., Zhao, Y., Clark, J.R. 2009. High tunnel versus open field: management of primocane-fruiting blackberry using pruning and tipping to increase yield and extend the fruiting season. *HortScience*. 44(6):1581-1587."
- Clark, J.R., McCall, C., Finn, C.E. 2008. Blackberry. In: Finn, C., Clark, J., editors. Register of newfruit and nut cultivars- List 44. *HortScience*. 43:1323-1324.
- Finn, C.E., Clark, J.R. 2008. Register of new fruit and nut cultivars- List 44. *HortScience*. 43:1321-1343.
- Bryla, D.R., Strik, B.C. 2008. Water requirements of young blueberry plants irrigated by sprinklers, microsprays, and drip. *Acta Horticulturae*. 792:135-139.
- Bryla, D.R., Yang, W.Q., Linderman, R.G. 2008. Incidence of *Phytophthora* and *Pythium* infection and the relation to cultural conditions in commercial blueberry fields. *HortScience*. 43(1):260-263.
- Dossett, M., Lee, J., Finn, C.E. 2008. Inheritance of phenological, vegetative, and fruit chemistry traits in black raspberry. *Journal of the American Society for Horticultural Science*. 133:408-417.
- Finn, C.E., Thomas, A.L., Byers, P.L., Serce, S. 2008. Evaluation of American (*Sambucus canadensis*) and European (*S. nigra*) elderberry genotypes grown in Missouri and Oregon and impact on cultivar development. *HortScience*. 43(5):1385-1391.
- Machado, R., Bryla, D.R., Verissimo, M.L., Sena, A.M., Oliverira, M. 2008. Nitrogen requirements for growth and early fruit development of drip-irrigated processing tomato (*Lycopersicon esculentum* Mill.) in Portugal. *Journal of Food Agriculture and Environment*. 6(3&4):215-218.
- Resendes, M.L., Bryla, D.R., Eissenstat, D.M. 2008. Early events in the life of apple roots: variation in root growth rate is linked to mycorrhizal and nonmycorrhizal fungal colonization. *Plant and Soil Journal*. Available: <http://www.springerlink.com/content/275k6l4m56234623/>
- Strik, B.C., Clark, J.R., Finn, C.E., Banados, M.P. 2008. Worldwide production of blackberries. *Acta Horticulturae*. 777:209-218.
- Strik, B.C., Finn, C.E., Clark, J.R., Buller, G. 2008. Management of Primocane-Fruiting Blackberry to Maximize Yield and Extend the Fruiting Season. *Acta Horticulturae*. 777:423-428.
- Thomas, A.L., Byers, P.L., Finn, C.E., Chen, Y.C., Rottinghaus, G.E., Malone, A.M., Applequist, W.L. 2008. Occurrence of Rutin and Chlorogenic Acid in Elderberry Leaf, Flower, and Stem in Response to Genotype, Environment, and Season. *Acta Horticulturae*. 765:197-206.

- Weebadde, C.K., Wang, D., Finn, C.E., Lewers, K.S., Luby, J.J., Bushakra, J., Sjulín, T.M., Hancock, J.F. 2008. Using a linkage mapping approach to identify QTL for day-neutrality in the octoploid strawberry. *Plant Breeding*. 127:94-101.
- Thompson, J.R., Strik, B.C., Clark, J.R., Finn, C.E. 2007. Flowering and fruiting morphology of primocane-fruiting blackberries. *Acta Horticulturae*. 777:281-288.
- Jetti, R.R., Yang, E., Kurnianta, A., Finn, C.E., Qian, M.C. 2007. Quantification of selected aroma-active compounds in strawberries by headspace solid-phase microextraction gas chromatography and correlation with sensory descriptive analysis. *Journal of Food Science*. S487-S496.
- Thompson, E., Strik, B.C., Clark, J.R., Finn, C.E. 2007. Flowering and fruiting patterns of primocane-fruiting blackberries. *HortScience*. 42:1174-1176.
- Strik, B.C., Clark, J.R., Finn, C.E., Banados, M.P. 2007. Worldwide blackberry production. *HortTechnology*. 17:205-213.
- Moore, P.P., Finn, C.E. 2007. 'Cascade Bounty' red raspberry. *HortScience*. 42:393-396.
- Kempler, C., Daubeney, H.A., Harding, B., Baumann, T., Finn, C.E., Moore, P.P., Sweeney, M., Walters, T. 2007. 'Saanich' red raspberry. *HortScience*. 42:176-178.
- Bryla, D.R. and Linderman, R.G. 2007. Implications of irrigation method and amount of water application on Phytophthora and Pythium infection and severity of root rot in highbush blueberry. *HortScience*. 42:1463-1467.
- Benedict, C., Skinner, J.S., Meng, R., Chang, Y., Bhalerao, R., Huner, N.A., Finn, C.E., Chen, T.H., Hurry, V. 2006. The cbf1-dependent low temperature signalling pathway, regulon and increase in freeze tolerance are conserved in populus spp. *Plant Cell and Environment*. 29(7):1259-1272.
- Clark, J.R., Finn, C.E. 2006. Blackberry And Hybrid Berry. *HortScience*. 41(5):1104-1106.
- Clark, J.R., Finn, C.E. editors. 2006. Register Of New Fruit And Nut Cultivars, List 43. *HortScience*. 41(5): 1101-1133.

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- Hummer, K.E., Bassil, N.V., Njuguna, W. 2011. *Fragaria*. Book Chapter. 6(WCR):17-44.
- Bassil, N.V., Martin, R.C. 2010. Proceedings of the Second International Symposium on Molecular Markers in Horticulture *Acta Horticulturae*. 49(4):53-54.
- Bassil, N.V., Muminova, M., Njuguna, W. 2010. Microsatellite-Based Fingerprinting of Western Blackberries from Plants, IQF Berries and Puree. *Acta Horticulturae*. 859:73-80.
- Bassil, N.V., Volk, G.M. 2010. Standardized Phenotyping: Advantages to Horticulture, Introduction to the Workshop. *HortScience*. 45(9):1306.
- Bassil, N.V., Bunch Jr, T.R., Nyberg, A.M., Zee, F.T., Hummer, K.E. 2010. Microsatellite Markers Distinguish Hawaiian Ohelo from Other *Vaccinium* L. Section *Myrtillus* Species. *Acta Horticulturae*. 859:81-88.
- Dalton, D., Postman, J.D., Hummer, K.E. 2010. Comparative Infectivity of *Cronartium ribicola* Aeciospores and Urediniospores in Genotypes of *Ribes nigrum*. *Plant Disease*. 94(4):461-464.
- Dalton, D.T., Hummer, K.E. 2010. *Ribes* Bloom Phenology: Section *Botrycarpum* and *Ribes*. *Journal of American Pomological Society*. 64(3):140-150.
- Dossett, M., Bassil, N.V., Finn, C.E. 2010. Transferability of *Rubus* Microsatellite Markers to Black Raspberry. *Acta Horticulturae*. 859:103-109.
- Flores, N., Reed, B.M., Graham, J., Bassil, N.V. Microsatellite Markers for Raspberries, Blackberries, and their Hybrids. 2010. *Journal of the American Society for Horticultural Science*. 135(3):271-278.
- Flores, N.R., Reed, B.M., Graham, J., Fernandez-Fernandez, F., Bassil, N.V. 2010. Microsatellite Markers for Raspberries and Blackberries. *Journal of the American Society for Horticultural Science*. 135(3):271-278.
- Hoefer, M., Reed, B.M. 2010. Conservation Strategy of Strawberry Genetic Resources in Germany. Abstract of International Horticultural Congress. 2:524.
- Hummer, K.E. 2010. Maxine M. Thompson - Dedication. *Horticultural Reviews*. 38:xvii-xix.
- Hummer, K.E. 2010. *Rubus* Pharmacology: Antiquity to the Present. *HortScience*. 45:1587-1591.
- Hummer, K.E., Dale, A. 2010. Horticulture of *Ribes*. *Forest Pathology*. 40(2010):251-263.
- Hummer, K.E., Postman, J.D., Iketani, H., Imanishi, H. 2010. Japan - United States Expedition to Hokkaido and Northern Honshu to collect Fruit and Nut Genetic Resources. International Horticultural Congress. 2:524.
- Iezzoni, A., Weebadde, C., Luby, J., Yue, C., Weg, E., Fazio, G., Bassil, N.V., Main, D., Peace, C., Mcferson, J. 2010. RosBREED: Enabling Marker-Assisted Breeding in Rosaceae. *Acta Horticulturae*. 859:389-394.
- Iketani, H., Hummer, K.E., Postman, J.D., Imanishi, H., Mase, N. 2010. Collaborative Exploration between NIAS Genebank and USDA ARS for the Collection of Genetic Resources of Fruit and Nut Species in Hokkaido and the Northern Tohoku Region. *Journal of Japanese Botany*. 26:13-26.
- Janick, J., Hummer, K.E. 2010. Healing, Health, and Horticulture: Introduction to the Workshop. *HortScience*. 45(11):1584-1586.

- Kushnarenko, S., Kovalchuk, I., Mukhitdinov, Z., Rakhimov, E., Reed, B.M. 2010. Cold Acclimation and Cryopreservation Treatments Increase the Cell Size of Apple Meristem Cells During Cryopreservation. *Asian and Australasian Journal of Plant Science and Biotechnology*. p. 10-20.
- Mota, J.A., Hummer, K.E., Williams, R. 2010. Berry Trials in the Azores. *Acta Horticulturae*. 2:579.
- Nathewet, P., Yanagi, T., Hummer, K.E., Iwatsubo, Y., Sone, K. 2010. Karyotype Analysis in Octoploid and Decaploid Wild Strawberries *Fragaria* (Rosaceae). *Cytologia*. 74(3):355-364.
- Nathewet, P., Hummer, K.E., Iwatsubo, Y., Sone, K., Yanagi, T. 2010. Karyotype Analysis in Octoploid and Decaploid Wild Strawberries, *Fragaria* (Rosaceae). *Cytologia*. 75(3):277-288.
- Njuguna, W., Liston, A., Cronn, R., Bassil, N.V. 2010. Multiplexed *Fragaria* Chloroplast Genome Sequencing. *Acta Horticulturae*. 859:315-320.
- Postman, J.D., Bretting, P.K., Kinard, G.R., Cyr, P.D., Weaver, B., Millard, M.J., Gardner, C.A., Bohning, M.A., Emberland, G.P., Sinnott, Q.P., Ayala Silva, T., Hummer, K.E., Franco, T., Mackay, M., Guarino, L. 2010. GRIN-Global: An International Project to Develop a Global Plant Genebank Information Management System. *Acta Horticulturae*. 859:49-55.
- Postman, J.D., Volk, G.M., Aldwinckle, H. 2010. Standardized Plant Disease Evaluations will Enhance Resistance Gene Discovery. *HortScience*. 45:1317-1320.
- Reed, B.M. 2010. Techniques for Short and Long-Term Preservation of Plant Tissues and Organs. *In Vitro*. 46:S284.
- Reed, B.M., Castillo, N., Wada, S., Bassil, N.V. 2010. Genetic Stability of Cryopreserved Shoot Tips of *Rubus* Germplasm. *In Vitro Cellular and Developmental Biology - Plants*. 46:246-256.
- Reed, B.M., Uchendu, E., Muminova, M., Traber, M. 2010. Improving Shoot Tip Cryopreservation with Antioxidant and Antistress Compounds. *In Vitro Cellular and Developmental Biology - Plants*. 46:S280.
- Tzanetakis, I.E., Postman, J.D., Samad, A., Martin, R.R. 2010. Mint Viruses: Beauty, Stealth and Disease. *Plant Disease*. 94(1):4-12.
- Uchendu, E., Leonard, S.W., Traber, M.G., Reed, B.M. 2010. Vitamins C and E Improve Regrowth and Reduce Lipid Peroxidation of Blackberry Shoot Tips Following Cryopreservation. *Plant Cell Reports*. 29:25-35.
- Uchendu, E.E., Muminova, M., Gupta, S., Reed, B.M. 2010. Antioxidant and Anti-stress Compounds Improve Regrowth of Cryopreserved *Rubus* Shoot Tips. *In Vitro Cellular and Developmental Biology - Plants*. 46:246-256.
- Wada, S., De Noma, J.S., Niedz, R.P., Evens, T.J., Reed, B.M. 2010. Improved Mineral Nutrition for Pear Shoot Cultures. *In Vitro Cellular and Developmental Biology - Plants*. 46:s282.
- Wada, S., Nonogaki, H., Reed, B.M. Identification of Blackberry Cultivars by Seed Structure. Oregon State University Extension Publications Online Catalog #EM9002. pp. 25.
- Wada, S., Reed, B.M. 2010. Seed Coat Morphology Differentiates Blackberry Cultivars. *Journal of American Pomological Society*. 64(3):151-160.
- Dalton, D.T., Hummer, K.E. 2009. Inheritance of the Cr Gene in *Ribes nigrum*. *Journal of American Pomological Society*. 63(4):142-144.
- Bretting, P.K., Kinard, C.E., Cyr, P.D., Weaver, B., Millard, M.J., Gardner, C.A., Bohning, M.A., Emberland, G.P., Sinnott, Q.P., Postman, J.D., Hummer, K.E., Ayala Silva, T., Franco, T., Mackay, M., Gaurino, L. 2009. GRIN-Global: An International Project to Develop a Global Plant Genebank and Information Management System. Meeting Abstract. 44(4):1179.
- Bassil, N.V., Postman, J.D. 2009. Identification of European and Asian Pears Using EST-SSRs From *Pyrus*. *Genetic Resources and Crop Evolution*. 57:357-370.
- Bassil, N.V., Lewers, K.S. 2009. Genomics Opportunities, New Crops and New Products. In: Kevin M. Folta, Susan E. Gardiner. *Genetics and Genomics of Rosaceae*. New York, NY: Springer-Verlag, LLC. Chapter 3 p.55-70.
- Bassil, N.V., Postman, J.D., Hummer, K.E., Sezer, A., Botu, M. 2009. SSR Fingerprinting Panel Verifies Identities of Clones in Backup Hazelnut Collection of USDA Genebank. *Acta Horticulturae*. 845:95-102.
- Hall, H., Hummer, K.E., Jaimieson, A., Jennings, S., Weber, C. 2009. Plant Breeding Reviews: Raspberry Breeding and Genetics. New Jersey: Wiley Blackwell. 32:39-382
- Hinrichsen, P., M. Herminia, C., Ravest, G., Rojas, G., Mendez, M., Bassil, N.V., Munoz, C. 2009. Minimal Microsatellite Marker Panel for Fingerprinting Blueberry Cultivars. *Acta Horticulturae*. 810:173-180.
- Hummer, K.E. 2009. *Rubus* Pharmacology: Antiquity to the Present. *HortScience*. 44(4):985.
- Hummer, K.E., Strik, B.C., Finn, C.E. 2009. Proceedings of the Ninth International Vaccinium Symposium Vol. 1. International Society of Horticultural Science Proceedings. 810(1):1-438.
- Hummer, K.E., Strik, B.C., Finn, C.E. 2009. Proceedings of the Ninth International Vaccinium Symposium Vol.2. International Society of Horticultural Science Proceedings. 810(2):439-909.
- Njuguna, W., Hummer, K.E., Bassil, N.V. 2009. DNA Barcoding in *Fragaria* L. (Strawberry) Species [Abstract]. *HortScience*. 44(4):1090.
- Postman, J.D. *Cydonia oblonga*: The Unappreciated Quince. *Arnoldia*. 67(1):2-9.

- Postman, J.D., Aldwinckle, H., Volk, G.M. 2009. Standardized Plant Disease Evaluations will Enhance Resistance Gene Discovery. Meeting Proceedings. HortScience 44:975.
- Reed, B.M. 2009. Applying Cryopreservation Techniques to Diverse Biological Materials. University of Kebangsaan Malaysia Monograph Series. p. 18.
- Bassil, N.V., Hummer, K.E. 2009. Blueberry Microsatellite Markers Identify Cranberries. [Abstract] International Society for Horticultural Science Meeting. Acta Horticulturae. 810:181-187.
- Bassil, N.V., Oda, A.K., Hummer, K.E. 2009. Blueberry Microsatellite Markers Identify Cranberry Cultivars. Acta Horticulturae. 810:181-186.
- Hummer, K.E. 2009. Global Conservation of Strawberries: A Strategy is Formed. Acta Horticulturae. 842:577-580
- Hummer, K.E., Hall, H. 2009. Dedication: Hugh Daubeny, Berry Breeder and Enthusiast Extraordinaire. In: Janick, J. Plant Breeding Reviews Volume 32. Raspberry Breeding and Genetics. Hoboken, New Jersey: John Wiley & Sons, Inc. p.1-37.
- Hummer, K.E., Hancock, J. 2009. Strawberry Genomics: Botanical History, Cultivation, Traditional Breeding, and New Technologies. In: K. Folta and S. Gardiner (eds.) Genetics and Genomics of Rosaceae. New York, NY: Springer. Chapter 11, p.413-435.
- Hummer, K.E., Janick, J. 2009. Rosaceae: Taxonomy, Economic Importance, Genomics. In: K. Folta and S. Gardiner (eds.) Genetics and Genomics of Rosaceae. New York, NY: Springer. Chapter 1, p. 1-17.
- Hummer, K.E., Williams, R., Mota, J. 2009. Pests of Blueberries on Sao Miguel, Azores, Portugal. Acta Horticulturae. 810:287-292.
- Kovalchuk, I., Lyudvikoval, Y., Volgenial, M., Reed, B.M. 2009. Medium Container and Genotype All Influence In Vitro Cold Storage of Apple Germplasm. Plant Cell Tissue And Organ Culture. 96:127-136.
- Njuguna, W., Bassil, N.V., Hummer, K.E., Richards, C.M., Davis, T. 2009. Genetic Diversity of Japanese Strawberry Species Based on Microsatellite Markers. Acta Horticulturae. 842:581-584.
- Peredo, E.L., Arroyo-Garcia, R., Reed, B.M., Revilla, M. 2009. Genetic Stability of In Vitro Conserved Germplasm of Humulus Lupulus L.. Experiment Station Bulletins. 18(2):144-151(8).
- Postman, J.D., Oliphant, J.M., Hummer, K.E. 2009. Plant Diseases Impact USDA Clonal Vaccinium Genebank. Acta Horticulturae. 810:319-324.
- Uchendu, E., Reed, B.M. 2009. Desiccation Tolerance and Cryopreservation of In-Vitro Grown Blueberry and Cranberry Shoot Tips. Acta Horticulturae. 810:567-574.
- Bassil, N.V., Hummer, K.E., Postman, J.D., Fazio, G., Baldo, A.M., Armas, I., Williams, R. 2008. Nomenclature and Genetic Relationships of Apples and Pears from Terceira Island. Genetic Resources and Crop Evolution (2009). 56:339-352.
- Hummer, K.E., Nathewet, P., Yanagi, T. 2008. Decaploidy in Fragaria Iturupensis Staudt (Rosaceae). American Journal of Botany. 96(3)713-716.
- Hummer, K.E., Postman, J.D. 2008. Pyrus L. Pear. In: Bonner, F.T. and Karrfalt, R.P., editors. The Woody Plant Seed Manual. Washington, D.C.: United States Government Printing Office. p. 922-927.
- Bassil, N.V., Postman, J.D., Hummer, K.E., Dolan, S., Lawliss, L., Oda, A. 2008. Molecular Fingerprints Identify Historic Pear Trees in Two U.S. National Parks. Acta Horticulturae. 800:417-422.
- Brevis, P.A., Bassil, N.V., Ballington, J., Hancock, J. 2008. Impact of Wide Hybridization in Southern Highbush Blueberry Breeding. Journal of the American Society for Horticultural Science. 133(3):427-437.
- Dalton, D., Hummer, K.E. 2008. Ribes Bloom Phenology in a Diverse Field Genebank [abstract]. HortScience. 43(4):1135.
- Hummer, K.E. 2008. Currants. HortScience. 43(5):1325.
- Hummer, K.E. 2008. Pawpaws. HortScience. 43(5):1329.
- Hummer, K.E., Bassil, N.V. 2008. Unexpected Polyploidy in Wild Asian Strawberries [abstract]. HortScience. 43(4):1187.
- Hummer, K.E., Gent, D.H., Henning, J.A., Mahaffee, W.F., Ocamb, C. 2008. Powdery Mildew Resistant Hop Germplasm Release: 'Kazak 2000' [abstract]. Acta Horticulturae. p.59
- Hummer, K.E., Sabitov, A. 2008. Strawberry Species of Iturup and Sakhalin Islands. HortScience. 43(5)2008:1623-1625.
- Njuguna, W., Bassil, N.V., Hummer, K.E., Slovin, J.P. 2008. Cross Species Amplification of Microsatellite Markers in Fragaria. North American Strawberry Conference Proceedings 2007. p. 122-123.
- Peredo, E.L., Arroyo-Garcia, R., Reed, B.M., Revilla, M. 2008. Epigenetic Stability of Cryopreserved and Cold-Stored Hops. Cryobiology. 57:234-241.
- Postman, J.D. 2008. Pears (In: Brooks & Olmo Register of New Fruit and Nut Varieties List 44). HortScience. 43(5):1334.
- Postman, J.D. 2008. The USDA Quince and Pear Genebank in Oregon, A World Source of Fire Blight Resistance. Acta Horticulturae. 793:357-362.
- Postman, J.D. 2008. World Pyrus Collection at USDA Germplasm in Corvallis, Oregon. Acta Horticulturae. 800:527-533.
- Reed, B.M., Hummer, K.E., Chang, Y., Gupta, S. 2008. MEDIUM AND LONG-TERM STORAGE OF RUBUS GERMPLASM. Acta Horticulturae. 777:91-98.

- Uchendu, E., Reed, B.M. 2008. A comparative study of three cryopreservation protocols for effective storage of in vitro-grown mint (*mentha* spp.). *CryoLetters*. 29(3):181-188.
- Wada, S., Reed, B.M. 2008. Morphological Analysis of *Rubus* Seed. *Acta Horticulturae*. (782)p. 67-74.
- Bassil, N.V., Gilmore, B.S., Oliphant, J.M., Hummer, K.E., Henning, J.A. 2007. Genic SSRs for European and North American Hop (*Humulus lupulus* L.). *Genetic Resources and Crop Evolution*. Available: www.springerlink.com/content/d000k04454n55v2p/fulltext.pdf
- Castillo, N., Bassil, N.V., Reed, B.M. 2007. Microsatellite Markers Identify Raspberries and Blackberries. *Plant and Animal Genome*. p. 134.
- Castillo, N.F., Reed, B.M., Bassil, N.V. 2007. Fingerprinting and Genetic Stability of *Rubus* Using Molecular Markers. *HortScience*. 42(4):914.
- Hummer, K.E. 2007. Development of a Global Conservation Strategy for Strawberries. *Acta Horticulturae*. 760:49-54.
- Hummer, K.E. 2007. Global Conservation of Strawberries: A Strategy is Formed. *International Strawberry Symposium*.
- Hummer, K.E. 2007. Introduction to the workshop. *HortScience*. 42:190.
- Hummer, K.E. 2007. Second International Symposium on Plant Genetic Resources of Horticultural Crops - Volume 2. *Acta Horticulturae*. Leuven, Belgium. ISHS. 308 p.
- Hummer, K.E. 2007. Second International Symposium on Plant Genetic Resources of Horticultural Crops - Volume I. *Acta Horticulturae*. Leuven, Belgium. ISHS. 368 p.
- Hummer, K.E., Durst, R. 2007. Fruit Anthocyanins In *Vaccinium praestans* Lamb.. *HortScience*. 42(4):1008.
- Hummer, K.E., Janick, J. 2007. *Rubus* Iconography: Antiquity to the Renaissance. *Acta Horticulturae*. 759:89-106.
- Hummer, K.E., Layne, D.R. 2007. Introduction to the Workshop Blueberries - Now and the Future". *Journal of American Pomological Society*. 61(3):138-139."
- Hummer, K.E., Zee, F.T., Strauss, A.J., Keith, L.M., Nishijima, W. 2007. Evergreen Production of Southern Highbush Blueberries in Hawaii. *Journal of American Pomological Society*. 61(4):188-195.
- Njuguna, W., Bassil, N.V., Davis, T. 2007. Genetic Diversity of *Fragaria inumae* and *F. nipponica* Based on Microsatellite Markers. *HortScience*. 42(4):915.
- Postman, J.D., Hummer, K.E., Stover, E.W., Krueger, R., Forsline, P.L., Grauke, L.J., Zee, F.T., Ayala Silva, T., Irish, B.M. 2007. Fruit and Nut Genebanks in the US National Plant Germplasm System. *Pomona*.
- Reed, B.M. 2007. Cryopreservation of Temperate Berry Crops. In: Reed, B.M. *Plant Cryopreservation, A Practical Guide*. New York, NY:Springer. p. 333-364.
- Reed, B.M. 2007. Cryopreservation-Practical Considerations. In: Reed, B.M. *Plant Cryopreservation, A Practical Guide*. New York, NY: Springer. p. 3-14.
- Reed, B.M. 2007. Culture and Storage Medium Iron Concentration Influence In-Vitro Cold Storage Duration of Hops (*Humulus*). *In Vitro Cellular and Developmental Biology - Plants*. 43:529-530.
- Reed, B.M. 2007. *Plant Cryopreservation, A Practical Guide*. New York, NY: Springer. 515 p.
- Reed, B.M., Uchendu, E. 2007. Controlled Rate Cooling. In: Reed, B.M. *Plant Cryopreservation, A Practical Guide*. New York, NY. p. 77-92.
- Sabitov, A., Cherbukin, P., Hummer, K.E. 2007. Plant Exploration for Fruit Genetic Resources in Sakhalin Territory. *Acta Horticulturae*. 760:381-388.
- Uchendu, E.E., Reed, B.M. 2007. A comparative study of three cryopreservation protocols for effective storage of mint (*mentha* spp.). *In Vitro Cellular and Developmental Biology*. 43:545.
- Zhao, Y., Wu, Y., Chang, Y., Reed, B.M. 2007. Cryopreservation of Fruit and Ornamental Trees. In: Reed, B.M., editor. *Plant Cryopreservation A Practical Guide*. New York, NY: Springer. p. 387-420.
- Blom, P.E., Tarara, J.M. 2007. Rapid, non-destructive estimation of leaf area on field-grown *Vitis labruscana* grapevines. *American Journal of Enology and Viticulture*. 58:393-397.
- Sardani, M., Spotts, R.A., Calabro, J.M., Postman, J.D., Qu, A.P. 2006. Evaluation of the USDA National Clonal *Pyrus* Germplasm Collection for Resistance to *Podosphaera leucotricha*. *HortScience*. 41(3):717-720.
- Sargent, D.J., Clarke, J., Simpson, D.W., Arus, P., Monfort, A., Vilanova, S., Denoyes-Rothan, B., Folta, K.M., Bassil, N.V., Battey, N.H. 2006. An enhanced microsatellite map of diploid *fragaria*. *Theoretical and Applied Genetics*. 112:1349-1359.

5358-21000-040-00D

GENOMICS, GERMLASM DEVELOPMENT AND IPM OF HOP – John Henning (P), David Gent, and Gary Banowetz; Corvallis, Oregon

- Gent, D.H., Ocamb, C.M. 2009. Prediction of infection risk of hop by *Pseudoperonospora humuli*. *Phytopathology*. 99:1190-1198.

- Schwartz, H.F., Gent, D.H., Fichtner, S.M., Hammon, R., Cranshaw, W.S., Mahaffey, L., Camper, M., Otto, K., Mcmillan, M. 2009. Straw Mulch and Reduced-Risk Pesticide Impacts upon Thrips and Iris yellow spot virus on Western-Grown Onions. *Southwestern Entomologist*. 34(1):13-29.
- Gent, D.H., Bull, C.T. 2009. Bacterial Diseases of Minor Importance. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium of Hop Diseases and Pests*. St. Paul, MN: APS Press. p. 56.
- Gent, D.H., Bull, C.T. 2009. Crinkle Disease. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium of Hop Diseases and Pests*. St. Paul, MN: APS Press. p. 54.
- Gent, D.H., Bull, C.T. 2009. Crown Gall. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium of Hop Diseases and Pests*. St. Paul, MN: APS Press. P. 55.
- Gent, D.H., James, D.G., Wright, L.C., Brooks, D.J., Barbour, J.D., Dreves, A.J., Fisher, G.C., Walton, V.M. 2009. Effects of Powdery Mildew Fungicide Programs on Twospotted Spider Mite (Acari: Tetranychidae), Hop Aphid (Homoptera: Aphididae), and Their Natural Enemies in Hop Yards. *Journal of Economic Entomology*. 102(1):274-286.
- Gent, D.H., Nelson, M.E., Farnsworth, J.L., Grove, G.G. 2009. PCR detection of *Pseudoperonospora humuli* in air samples from hop yards. *Plant Pathology*. p. 1365-1369.
- Henning, J.A., Pethybridge, S.J., Gent, D.H. 2009. Hop Cultivars and Breeding. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium on Hop Diseases and Pests*. St. Paul, MN: APS Press. p. 11-13.
- Johnson, D.A., Engelhard, B., Gent, D.H. 2009. Downy Mildew. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium of Hop Diseases and Pests*. St. Paul, MN: APS Press. p. 18-22.
- Solarska, E., Grudzinska, M., Kaminska, M., Sliwa, H., Gent, D.H. 2009. Hop Shoot Proliferation. In: Mahaffee, W.F., Pethybridge, S.J., Gent, D.H., editors. *Compendium on Hop Diseases and Pests*. St. Paul, MN: APS Press. p. 57.
- Townsend, M.S., Henning, J.A. 2009. AFLP Discrimination of Wild American and Cultivated Hop. *Crop Science*. 49:600-607.
- Gent, D.H., Nelson, M.E., Grove, G.G. 2008. Persistence of Phenylamide Insensitivity in *Pseudoperonospora humuli*. *Plant Disease*. 92(3):463-468.
- Gent, D.H., Turechek, W., Mahaffee, W.F. 2008. Spatial and Temporal Stability of the Estimated Parameters of the Binary Power Law. *Phytopathology*. 98:1107-1117.
- Henning, J.A., Haunold, A., Townsend, M.S., Gent, D.H., Parker, T. 2008. Registration of 'Teamaker' Hop. *Journal of Plant Registrations*. 2(1):13-14.
- Gent, D. H., Nelson, M. E., George, A. E., Grove, G. G., Mahaffee, W. F., Ocamb, C. M., Barbour, J. D., Peetz, A., and Turechek, W. W. 2008. A decade of hop powdery mildew in the pacific northwest. Online. *Plant Health Progress* doi:10.1094/PHP-2008-0314-01-RV.
- Gent, D.H., Martin, R.R., Ocamb, C.M. 2007. First Report of Iris yellow spot virus on Onion and Leek in Western Oregon. *Plant Disease*. V91: P 468.
- Gent, D.H., Turechek, W., Mahaffee, W.F. 2007. Sequential sampling for estimation and classification of the incidence of hop powdery mildew I: Leaf sampling. *Plant Disease*. 91:1002-1012.
- Gent, D.H., Turechek, W., Mahaffee, W.F. 2007. Sequential sampling for estimation and classification of the incidence of hop powdery mildew II: Cone sampling. *Plant Disease*. 91:1013-1020.

5366-21000-025-00D

PLANT AND SEED CHEMISTRY GENETICS – Victor Raboy (P) and John Bonman; Aberdeen, Idaho

- Raboy, V. 2009. Approaches and Challenges to Engineering Seed Phytate and Total Phosphorus. *Plant Science* 177:281-296.
- Veum, T.L., Ledoux, D.R., Shannon, M.C., Raboy, V. 2009. Effect of graded levels of iron, zinc, and copper supplementation in diets with low-phytate or normal barley on growth performance, bone characteristics, hematocrit volume, and zinc and copper balance of young swine. *Journal of Animal Science*. 87:2625-2634
- Raboy, V. 2008. Forward genetics studies of seed phytic acid. *Israel Journal of Plant Sciences* 55:171-181.
- Erickson, C.A., Obert, D.E., Burrup, D.E., Blake, K., Dillon, M.A., Wesenberg, D.M., Whitmore, J.C. 2008. Registration of 'maverick' oat. *Journal of Plant Registrations*. 2(3):194-196.
- Fu, J., Peterson, K., Guttieri, M., Souza, E.J., Raboy, V. 2008. Barley (*Hordeum vulgare* L.) inositol monophosphatase: gene structure and enzyme characteristics. *Plant Molecular Biology*. 67:629-642.
- Salarmoini, M., Campbell, G.L., Rosnagel, B.G., Raboy, V. 2008. Retention and growth performance of chicks given low-phytate conventional or hull-less barleys. *British Poultry Science*. 49:321-328
- Linares, L.B., Broomhead, J.N., Guaiume, E.A., Ledoux, D.R., Veum, T.L., Raboy, V. 2007. Effects of Low Phytic Acid Barley on Zinc Utilization in Young Broiler Chicks. *Poultry Science*. v 86:299-308
- Edney, M.J., Rosnagel, B.G., Raboy, V. 2007. Effect of low-phytate barley on malt quality including mineral loss during fermentation. *Journal of American Society of Brewing Chemists*. 65: 81-85
- Raboy, V. 2007. The ABCs of low-phytate crops. *Nature Biotechnology* 25: 874-875

- Veum, T.L., Ledoux, D.R., Raboy, V. 2007. Low-phytate barley cultivars improve nutrient utilization in diets fed to young swine. *Journal of Animal Science*. 85: 961-971
- Hambidge, K.M., Westcott, J.E., Lei, S., Grunwald, G., Krebs, N.F., Mazariegos, M., Solomons, N.W., Raboy, V. 2007. Intestinal excretion of endogenous zinc in guatemalan school children. *Journal of Nutrition*. 137:1747-1749
- Mazariegos, M., Solomons, N.W., Hambidge, M.K., Krebs, N.F., Westcott, J.E., Sian, L., Campos, R., Barahona, B., Raboy, V. 2006. Zinc absorption in guatemalan school children fed normal or low-phytate maize. *American Journal of Clinical Nutrition*. v 83 p 59-64
- Raboy, V. 2006. Seed Phosphorus and the development of low-phytate crops. *Inositol Phosphates in the Plant and Soil System* 111-132
- Thacker, P.A., Rossnagel, B.G., Raboy, V. 2006. The effects of phytase supplementation on nutrient digestibility, plasma parameters, performance and carcass traits of pigs fed diets based on low-phytate barley fed without inorganic phosphorus. *Canadian Journal of Animal Science* 86: 245-254
- Bowen, D.E., Gutierrez, M.J., Peterson, K.L., Raboy, V., Souza, E.J. 2006. Phosphorus fractions in developing seeds of four low phytate barley (*hordeum vulgare* L.) genotypes. *Crop Science* 46: 2468-2473

5366-21000-026-00D

DEVELOPMENT OF POTATO VARIETIES AND GERMLASM WITH IMPROVED RESISTANCES, PRODUCTION EFFICIENCIES, AND TUBER QUALITIES FOR THE WESTERN U.S. – Richard Novy (P), Jonathan Whitworth, and John Bonman; Aberdeen, Idaho

- Novy, R.G., Whitworth, J.L., Stark, J.C., Love, S.L., Corsini, D.L., Pavék, J.J., Vales, M.I., James, S.R., Hane, D.C., Shock, C.C., Charlton, B.A., Brown, C.R., Knowles, N.R., Pavék, M.J., Brandt, T.L., Gupta, S., Olsen, N. 2010. Clearwater Russet: A Dual-Purpose Potato Cultivar with Cold Sweetening Resistance, High Protein Content, and Low Incidence of External Defects and Sugar Ends. *American Journal of Potato Research*. 87:458-471.
- Stark, J.C., R.G. Novy, J. L. Whitworth, N.R. Knowles, M.J. Pavék, S.L. Love, M.I. Vales, S.R. James, D.C. Hane, C.R. Brown, B.A. Charlton, D.L. Corsini, J.J. Pavék, N. Olsen and T. Brandt. 2010. Classic Russet: A Potato Cultivar with Excellent Fresh Market Characteristics and High Yields of U.S. No. 1 Tubers Suitable for Early Harvest or Full-Season Production. *Amer. J. Potato Res.* 87:360-373.
- Whitworth J.L., Hamm, P.B., McIntosh, C.S. 2010. Effect of potato virus Y on yield of a clonal selection of Russet Norkotah potato. *Amer J Potato Res.* 87:310-314.
- Whitworth, J.L., Novy, R.G., Stark, J., Pavék, J.J., Corsini, D.L., Love, S.L., Miller, J.S., Vales, M.I., Mosley, A.R., Yilma, S., James, S.R., Hane, D.C., Charlton, B.A., Brown, C.R., Knowles, N.R., Pavék, M.J. 2010. Yukon Gem: A yellow-fleshed potato cultivar suitable for fresh-pack and processing with resistances to PVYO and late blight. *Amer J Potato Res.* 87:327-336.
- Stark, J.C., R.G. Novy, J.L. Whitworth, S.L. Love, D.L. Corsini, J.J. Pavék, M.I. Vales, S.R. James, D.C. Hane, B.A. Charlton, C.R. Brown, N.R. Knowles, M.J. Pavék, T.L. Brandt, and N. Olsen. 2009. Highland Russet: A Full Season, Processing Variety with High Yields of Uniform U.S. No. 1 Tubers. *American Journal of Potato Research* 86:171-182
- Whitworth, J.L., Novy, R.G., Hall, D.G., Crosslin, J., Brown, C.R. 2009. Characterization of broad spectrum Potato virus Y resistance in a *Solanum tuberosum* ssp. andigena-derived population and select breeding clones using molecular markers, grafting, and field inoculations. *American Journal of Potato Research*. 86:286-296.
- Y-H., Druffel K.L., Whitworth J., Pavék M.J., Pappu H.R. 2009. Molecular characterization of two Potato Virus S isolates from late blight resistant genotypes of potato (*Solanum tuberosum*). *Archives of Virology*. 154:1861-1863
- Kelley, K., Whitworth, J.L., Novy, R.G. 2009. Mapping of the potato leafroll virus resistance gene, Rlretb, from *Solanum tuberosum* identifies interchromosomal translocations among its E-genome chromosomes 4 and 9 relative to the A-genome of *Solanum L. sect. Petota*. 2009 *Molecular Breeding* 23:489-500.
- Novy, R.G., Whitworth, J.L., Stark, J.C., Love, S.L., Corsini, D.L., Pavék, J.J., Vales, M.I., James, S.R., Hane, D.C., Shock, C.C., Charlton, B.A., Brown, C.R., Knowles, N.R., Pavék, M.J., Brandt, T.L., Olsen, N. 2008. Premier russet: a dual-purpose, potato cultivar with significant resistance to low temperature sweetening during long-term storage. *American Journal of Potato Research*. 85:198-209
- Srinivasan, R., Alvarez, J., Bosque-Perez, N.A., Eigenbrode, S.D., Novy, R.G. 2008. Effect of an alternate weed host, hairy nightshade, *solanum sarrachoides* (sendtner), on the biology of the two most important potato leafroll virus (luteoviridae: polerovirus) vectors. *myzus persicae* (sulzer) and *macrosiphum*. *Environmental Entomology*. 37 (2):592-600
- Mosley, A.R., Yilma, S., Hane, D.C., James, S.R., Rykbost, K.A., Shock, C.C., Love, S.L., Corsini, D.L., Pavék, J.J., Thornton, R.E., Charlton, B.A., Eldredge, E.P., Novy, R.G., Pavék, M.J., Knowles, N.R., Whitworth, J.L., Brown, C.R., Stark, J.C., Vales, M.I. 2008. Willamette: A Chipping Cultivar with High Yield and Specific

- Gravity, Low Incidence of Hollow Heart and Brown Center, and Suitability for Fresh-Market Usage. *American Journal of Potato Research* 85(1): 85-92
- Novy, R.G., Gillen, A.M., Whitworth, J.L. 2007. Characterization of the expression and inheritance of potato leafroll virus (PLRV) and potato virus Y (PVY) resistance in three generations of germplasm derived from *Solanum tuberosum*. *Theoretical and Applied Genetics*. 114:1161-1172
- Novy, R.G., Miller, J.S., Porter, L., Whitworth, J.L., Lozoya-Saldana, H. 2007. Segregation of progeny of *Solanum tuberosum* subsp. *andigena* for foliar and tuber resistance to late blight. *American Journal of Potato Research*. 84(1): 106-107
- Stark, J.C., Novy, R.G., Love, S.L., Whitworth, J.L., Corsini, D.L., Pavek, J.J., Mosley, A.R., Pavek, M.J., Knowles, N.R., Thornton, R.E., James, S.R., Hane, D.C., Olsen, N., Vales, M.I., Brown, C.R. 2007. Blazer Russet: An early to mid-season potato cultivar with high U.S. No. 1 yields and good processing and culinary qualities. *American Journal of Potato Research*. 84 (6):467-477
- Thompson, A.L., Taylor, R.J., Pasche, J.S., Novy, R.G., Gudmestad, N.C. 2007 Resistance to *Phytophthora erythroseptica* and *Pythium ultimum* in a potato clone derived from *S. berthaultii* and *S. tuberosum*. *American Journal of Potato Research* 84:149-160
- Whitworth, J.L., Novy, R.G., Pavek, J.J., Corsini, D.L., Lozoya-Saldana, H., Yilma, S., Mosley, A.R. 2007. Multiple-Site Identification of Potato Parent Clones Conferring High Levels of Late Blight Resistance with a Corresponding Genetic Model for Resistance. *American Journal of Potato Research*. 84: 313-321
- Bonman, J.M., Bockelman, H.E., Jin, Y., Hijmans, R.J., Gironella, A. 2007. Geographic Distribution of Stem Rust Resistance in Wheat Landraces. *Crop Science*. 47:1955-1963.
- Kelley, K., Gillen, A.M., Whitworth, J.L., Novy, R.G. Development of molecular markers for introgression of potato leafroll virus resistance from *S. tuberosum* into cultivated potato. *American Journal of Potato Research*. 84(1):97
- Novy, R.G., Alvarez, J.M., Sterret, S.B., Kuhar, T.P., Horton, D.R. 2006. Progeny of a tri-species potato somatic hybrid express resistance to wireworm in eastern and western potato production regions of the U.S.. *American Journal of Potato Research*. 83(1):126
- Novy, R.G., Love, S.L., Corsini, D.L., Pavek, J.J., Whitworth, J.L., Mosley, A.R., James, S.R., Hane, D.C., Shock, C.C., Rykboost, R.Y., Brown, C.R., Thornton, R.E., Knowles, N.R., Pavek, M.J., Olsen, N., Inglis, D.A. 2006. Defender: a high-yielding, processing potato cultivar with foliar and tuber resistance to late blight. *American Journal of Potato Research*. 83(1):126
- Novy, R.G., Miller, J.S., Porter, L., Yilma, S. 2006. Identification and use of *Solanum tuberosum* subsp. *andigena* clones having both foliar and tuber resistance to late blight. *American Journal of Potato Research*. 83(1):126-127

5366-21000-027-00D

MANAGEMENT OF NATIONAL SMALL GRAINS COLLECTION RESOURCES – Harold Bockelman (P), John Bonman, and Blair Goates; Aberdeen, Idaho

- Adhikari, T.B., Hansen, J.M., Gurung, S., Bonman, J.M. 2011. Identification of New Sources of Resistance to *Xanthomonas translucens* pv. *undulosa* in Winter Wheat. *Plant Disease*. PDIS-10-10-0760-RE.R2.
- Goates, B., Peterson, G.L., Bowden, R.L., Maddux, L.D. 2011. Analysis of induction and establishment of dwarf bunt of wheat under marginal climatic conditions. *Plant Disease*. 95:478-484.
- Bockelman, H.E., Valkoun, J. 2010. Barley germplasm conservation and resources" Chapter 7 in "Barley: Improvement production and uses" Blackwell Publishing ed.s.e. Ullrich. *Barley: Improvement Production and Uses*. 144-159."
- Goates, B. J. 2010. Survival of secondary sporidia of floret-infecting *Tilletia* species: Implications for epidemiology. *Phytopathology* 100:655- 662.
- Goates, B.J., Peterson, G.L., Bowden R.L., and Maddux, L.D. 2010. Analysis of induction and establishment of dwarf bunt of wheat under marginal climatic conditions. *Phytopathology* 100 No. 6 (Supplement): S41.
- Gurung, S., Bonman, J.M., Ali, S., Patel, J., Myrfield, M., Mergoum, M., Singh, P.K., Adhikari, T.B. 2009. New and Diverse Sources of Multiple Disease Resistance in Wheat. *Crop Sci*. 49:1655-1666.
- Goates, B., Mercier, J. 2009. Effect of biofumigation with volatiles from *Muscodora albus* on the viability of *Tilletia* spp. teliospores. *Canadian Journal of Microbiology* 2009. Can. J. Microbiol. 55:203-206.
- Karasev, A.V., Meacham, T., Hu, X., Whitworth, J.L., Gray, S.M., Olsen, N., Nolte, P. 2008. Identification of Potato virus Y Strains Associated with Tuber Damage During a Recent Virus Outbreak in Potato in Idaho. *Plant Disease*. 92:1371.
- Erickson, C.A., Obert, D.E., Burrup, D.E., Blake, K., Dillon, M.A., Wesenberg, D.M., Whitmore, J.C. 2008. Registration of 'monico' oat. *Journal of Plant Registrations*. 2(3):197-199.
- Erickson, C.A., Obert, D.E., Burrup, D.E., Blake, K., Dillon, M.A., Wesenberg, D.M., Whitmore, J.C. 2008. Registration of 'maverick' oat. *Journal of Plant Registrations*. 2(3):194-196.

- Goates, B. 2008. Quantification of *Tilletia indica* teliospores in sori of commercially harvested wheat grains. *Phytopathology* 98 (6):S60.
- Mosley, A.R., Yilma, S., Hane, D.C., James, S.R., Rykbost, K.A., Shock, C.C., Love, S.L., Corsini, D.L., Pavék, J.J., Thornton, R.E., Charlton, B.A., Eldredge, E.P., Novy, R.G., Pavék, M.J., Knowles, N.R., Whitworth, J.L., Brown, C.R., Stark, J.C., Vales, M.I. 2008. Willamette: A Chipping Cultivar with High Yield and Specific Gravity, Low Incidence of Hollow Heart and Brown Center, and Suitability for Fresh-Market Usage. *American Journal of Potato Research* 85(1): 85-92.
- Goates, B., Mercier, J. 2007. Mortality of *Tilletia* spp. teliospores caused by volatiles from the biofumigant fungus *Muscodor albus*. *Phytopathology*.
- Goates, B. 2006. Anomalous and new epidemiological relationships of seed diseases caused by *tilletia* species. *Phytopathology* 96:S132.

5366-21000-028-00D

SMALL GRAINS GENETICS AND GERMPLASM ENHANCEMENT – John Bonman (P), Donald Obert, Paul Bregitzer, and Eric Jackson; Aberdeen, Idaho

- Verhoeven, E.C., Bonman, J.M., Bregitzer, P.P., Brunick, B., Cooper, B., Corey, A.E., Cuesta-Marcos, A., Filichkina, T., Mundt, C.C., Obert, D.E., Rossmagel, B., Richardson, K.L., Hayes, P.M. 2011. Registration of the BISON genetic stocks in *Hordeum vulgare* L.. *Journal of Plant Registrations*. 5:135-140.
- Gaylord, G.T., Barrows, F.T., Overturf, K.E., Liu, K., Hu, G. (2010) An Overview of Progress toward developing and all Plant-based Diet for Rainbow Trout. *Bull. Fish. Res. Agen.* 31:9-14.
- Acevedo, M., Jackson, E.W., Chong, J., Rines, H.W., Harrison, S., Bonman, J.M. 2010. Identification and Validation of Quantitative Trait Loci for Partial Resistance to Crown Rust in Oat. *Phytopathology*. 100(5):511-521.
- Acevedo, M., Jackson, E. W., Sturbaum, A., Ohm, H. W. and Bonman, J. M.(2010) 'An improved method to quantify *Puccinia coronata* f. sp. *avenae* DNA in the host *Avena sativa*', *Canadian Journal of Plant Pathology*, 32(2):215-224.
- Bonman, J.M., Gu, Y.Q., Coleman Derr, D., Jackson, E.W., Bockelman, H.E. 2010. Inferring geographic origin of barley accessions using molecular markers. *Genetic Resources and Crop Evolution*. DOI: 10.1007/s10722-010-9574-4.
- Bregitzer, P.P., Raboy, V., Obert, D.E. 2010. Registration of LP1-2581, LP1-2163H, LP3-1159, and LP640-1304 low phytate spring barley germplasm lines. *Journal of Plant Registrations*. 4:228-231.
- Oliver, R.E., Obert, D.E., Hu, G., Bonman, J.M., O'Leary-Jepsen, E., Jackson, E.W. 2010. Development of oat-based markers from barley and wheat microsatellites. *Genome*. 53(6): 458–471.
- Gurung, S., Bonman, J.M., Ali, S., Patel, J., Myrfield, M., Mergoum, M., Singh, P.K., Adhikari, T.B. 2009. New and Diverse Sources of Multiple Disease Resistance in Wheat. *Crop Sci.* 49:1655-1666.
- Jackson, C.A., Windes, J.M., Bregitzer, P.P., Obert, D.E., Price, W., Bradford, B. 2009. Low Phytic Acid Barley Responses to Phosphorus Rates. *Crop Science*. 49:1800-1806.
- Obert, D.E., Evans, C.P., Windes, J.M., Wesenberg, D.M., Ullrich, S., Budde, A.D., Chen, X., Jackson, E.W. 2009. Registration of Endeavor barley. *Journal of Plant Registrations*. 3:124-126.
- Tinker, N.A., Kilian, A., Wright, C.P., Heller-Uszynska, K., Wenzl, P., Rines, H.W., Bjornstad, A., Howarth, C., Jannink, J., Anderson, J.M., Rossmagel, B.G., Stuthman, D.D., Sorrells, M.E., Jackson, E.W., Tuveeson, S., Kolb, F.L., Olsson, O., Federizzi, L.C., Carson, M.L., Ohm, H.W., Molnar, S.J., Scoles, G.J., Eckstein, P.E., Bonman, J.M., Ceplitis, A., Langdon, T. 2009. New DArT markers for oat provide enhanced map coverage and global germplasm characterization. *BMC Medical Genetics*. 10:39.
- Raboy, V. 2009. Approaches and Challenges to Engineering Seed Phytate and Total Phosphorus. *Plant Science* 177:281-296.
- Veum, T.L., Ledoux, D.R., Shannon, M.C., Raboy, V. 2009. Effect of graded levels of iron, zinc, and copper supplementation in diets with low-phytate or normal barley on growth performance, bone characteristics, hematocrit volume, and zinc and copper balance of young swine. *Journal of Animal Science*. 87:2625-2634.
- Kelley, K., Whitworth, J.L., Novy, R.G. 2009. Mapping of the potato leafroll virus resistance gene, *Rlretb*, from *Solanum tuberosum* identifies interchromosomal translocations among its E-genome chromosomes 4 and 9 relative to the A-genome of *Solanum* L. sect. *Petota*. 2009 *Molecular Breeding* 23:489-500.
- Smith, S., Fenn, P., Chen, P., Jackson, E.W. 2008. Inheritance of resistance to *Phomopsis* seed decay in PI 360841 soybean. *Journal of Heredity*.
- Bonman, J.M., Gu, Y., Coleman Derr, D., Jackson, E.W., Chao, S., Bockelman, H.E. 2008. Origin of barley accessions with multiple disease resistance determined by SSR analysis.. *International Congress of Plant Pathology Abstracts and Proceedings*.

- Bregitzer, P.P., Dahleen, L.S., Neate, S., Schwarz, P., Manoharan, M. A. 2008. single backcross is an effective tool for the elimination of agronomic and quality alterations caused by somaclonal variation in barley . Crop Science. 48: 471-479.
- Bregitzer, P.P., Mornhinweg, D.W., Obert, D.E., Windes, J.M. 2008. Registration of 'RWA 1758' Russian wheat aphid-resistant spring barley. Journal of Plant Registrations. 2: 5-9.
- Bregitzer, P.P., Raboy, V., Obert, D.E., Windes, J., Whitmore, J. 2008. Registration of 'Clearwater' Low-Phytate Hulless Spring Barley. Journal of Plant Registrations. 2: 1-4.
- Jackson, E.W., Obert, D.E., Hu, G., Menz, M., Bonman, J.M. 2008. Qualitative and Quantitative trait loci conditioning resistance to Puccinia coronata races NQMG and LGCG in the oat (*Avena sativa* L.) cultivars Ogle and TAM O-301. Theoretical and Applied Genetics.v116:517-527.
- Jackson, E.W., Wise, M., Bonman, J.M., Obert, D.E., Hu, G. and Peterson, D.M.. 2008. QTLs Affecting -tocotrienol, -tocopherol, and total tocopherol concentrations detected in the Ogle/TAM O-301 oat mapping population. Crop Science 48:2141-2152.
- Obert, D.E., Evans, C.P., Wesenberg, D.M., Windes, J.M., Erickson, C.A., Jackson, E.W., Chen, X., Whitmore, J.C. 2008. Registration of 'Lenetah' barley. Journal of Plant Registrations. 2:85-87.
- Obert, D.E., Wesenberg, D.M., Burrup, D.E., Windes, J.M., Erickson, C.A., Whitmore, J.C. 2008. Registration of 'Tetonia' barley. Journal of Plant Registrations. 2:10-11.
- Raboy, V. 2008. Forward genetics studies of seed phytic acid. Israel Journal of Plant Sciences 55:171-181.
- Bonman, J.M., Bockelman, H.E., Jin, Y., Hijmans, R.J., Gironella, A. 2007. Geographic Distribution of Stem Rust Resistance in Wheat Landraces. Crop Science. 47:1955-1963.
- Bregitzer, P.P., Cooper, L.D., Hayes, P.M., Lemaux, P.G., Singh, J., Sturbaum, A.K. 2007. Viability and bar expression are negatively correlated in oregon wolfe barley dominant hybrids. Transgenic Research. Plant Biotechnology Journal v 5 p 381-388.
- Bregitzer, P.P., Obert, D.E., Wesenberg, D. 2007. Registration of 95SR316A Stripe rust-resistant Two-rowed Barley Germplasm. Journal of Plant Registrations 1:141-142.
- Bregitzer, P.P., Raboy, V. 2007. Registration of Four Low-Phytate/Wild Type Pairs of Barley Germplasm.. Journal of Plant Registrations 1:139-140.
- Bregitzer, P.P., Raboy, V., Obert, D.E., Windes, J.M., Whitmore, J.C. 2007. Registration of 'herald' barley. Crop Science 47:441-442, 2007.
- Jackson, E.W., Obert, D.E., Menz, M., Hu, G., Avant, J.B., Chong, J., Bonman, J.M. 2007. Characterization and mapping oat crown rust resistance using three assessment methods. Phytopathology.p.1063-1070.
- Jackson, E.W., Obert, D.E., Chong, J., Avant, J., Bonman, J.M. A detached leaf method for propagating Puccinia coronata races and assessing crown rust resistance in oat (*Avena sativa* L.). Plant Pathology 92:1400-1406.

5368-21220-003-00D

SUGARBEET GERMLASM DEVELOPMENT AND INNOVATIVE GENETIC AND MANAGEMENT APPROACHES TO REDUCING LOSSES CAUSED BY PATHOGENS – Carl Strausbaugh (P) and Imad Eujayl; Kimberly, Idaho

- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2011. Seed Treatments for the Control of Insects and Diseases in Sugarbeet. Journal of Sugarbeet Research. 47(3&4):105-125 (DOI: 10.5274/jsbr.47.3.105).
- Strausbaugh, C.A., Eujayl, I.A., Panella, L.W., Hanson, L.E. 2011. Virulence, Distribution and Diversity of Rhizoctonia Solani from Sugar Beet in Idaho and Oregon. Canadian Journal of Plant Pathology. 33(2): 210-226.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., and Foote, P. 2010. Commercial Sugar Beet Cultivars Evaluated for Rhizomania Resistance and Storability in Idaho, 2009. Plant Disease Management Reports online). Report 4:FC070 DOI:10.1094/PDMR04. The American Phytopathological Society, St. Paul, MN.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., and Foote, P. 2010. Experimental Sugar Beet Cultivars Evaluated for Rhizomania Resistance and Storability in Idaho, 2009. Plant Disease Management Reports online). Report 4:FC069 DOI:10.1094/PDMR04. The American Phytopathological Society, St. Paul, MN.
- Durrin, J.S., Nikolaeva, O.V., Strausbaugh, C.A., Karasev, A.V. 2010. Immunodetection of Two Curtoviruses Infecting Sugar Beet. Plant Disease. 94(8):972-976.
- Neher, O.T., and Strausbaugh, C.A. 2010. Evaluation of Fungicides as Root Dips for the Control of Root Rot in Storage, 2009. Plant Disease Management Reports (online). Report 4:FC075 DOI:10.1094/PDMR04. The American Phytopathological Society, St. Paul, MN. Available: <http://www.plantmanagementnetwork.org/pub/trial/pdmr/reports/2010/FC075.pdf>
- Strausbaugh, C.A. 2010. Improving Sugarbeet Storability. The Newsbeet. Spring Issue 23:22-23.
- Strausbaugh, C.A., Eujayl, I.A. 2010. Rhizoctonia and Bacterial Root Rot in Sugarbeet. The Sugarbeet. Spring Issue:11-12.
- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2010. Commercial Sugar Beet Cultivars Evaluated for Resistance to Bacterial Root Rot in Idaho, 2008. Plant Disease Management Reports. 4:FC010.

- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2010. Cultivar Selection for Sugar Beet Root Rot Resistance. *Phytopathology*. 100:S123.
- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2010. Experimental Sugar Beet Cultivars Evaluated for Resistance Bacterial Root Rot in Idaho, 2008. *Plant Disease Management Reports*. 4:FC011.
- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2010. Sugarbeet Cultivar Evaluation for Bacterial Root Rot. *Journal of Sugarbeet Research*. 47:51-64.
- Strausbaugh, C.A., Rearick, E., Eujayl, I.A., Foote, P. 2010. Effect of Insecticide Seed Treatments Improve Sugarbeet Storability. *Journal of Sugar Beet Research*. 47(3&4):65-88 (DOI: 10.5274/jsbr47.3.65).
- Eujayl, I.A., Simko, I., Strausbaugh, C.A. 2009. Association analysis of beet curly top disease resistant genes in sugarbeet [Abstract]. *Proceedings of 35th Biennial Meeting of American Society of Sugar Beet Technologists: Section B&E. Paper No. 1.*
- Strausbaugh, C.A. 2009. Reducing sucrose loss in sugarbeet storage. *Sugar Producer Magazine*. June/July:18-19.
- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2009. Insecticide seed treatments for sugarbeet. *The Sugarbeet*. Spring Issue:27.
- Eujayl, I.A., Strausbaugh, C.A. 2009. Evaluation of sugar beet germplasm and plant introductions response to rhizomania and storability in Idaho, 2008. *Plant Disease Management Reports*. 3:FC105.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., Foote, P. 2009. Commercial sugar beet cultivars evaluated for rhizomania resistance and storability in Idaho, 2008. *Plant Disease Management Reports*. 3:FC108.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., Foote, P. 2009. Experimental sugar beet cultivars evaluated for rhizomania resistance and storability in Idaho, 2008. *Plant Disease Management Reports*. 3:FC107.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., Foote, P., Ellison, D. 2009. Sugar beet cultivar evaluation for storability and rhizomania resistance. *Plant Disease*. 93:632-638.
- Strausbaugh, C.A., Gillen, A.M. 2009. Sugar Beet Root Rot at Harvest in the U.S. Intermountain West. *Canadian Journal of Plant Pathology*. 31:232-240.
- Eujayl, I.A., Morris, C.F. 2009. Identification of differentially expressed UniGenes in developing wheat seed using digital differential display. *Journal of Cereal Science*. 49:316-318.
- Strausbaugh, C.A., Eujayl, I.A. 2008. Influence of rhizomania on sugarbeet storability. *The Sugarbeet*. Spring Issue. p. 8.
- Strausbaugh, C.A., Eujayl, I.A., Foote, P. 2008. Transgenic sugar beet cultivars evaluated for resistance to bacterial root rot in Idaho, 2007. *Plant Disease Management Reports*. 2:FC108.
- Strausbaugh, C.A., Eujayl, I.A., Lewellen, R.T. 2008. Sugar beet germplasm evaluated for resistance to rhizomania and storability in Idaho, 2007. *Plant Disease Management Reports*. 2:FC105.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., Foote, P. 2008. Sugar beet cultivars evaluated for storability in Idaho during the 2006/2007 storage season. *Plant Disease Management Reports*. 2:FC022.
- Strausbaugh, C.A., Eujayl, I.A., Rearick, E., Foote, P. 2008. Transgenic sugar beet cultivars evaluated for resistance to rhizomania and storability in Idaho, 2007. *Plant Disease Management Reports*. 2:FC106.
- Strausbaugh, C.A., Gillen, A.M. 2008. Bacteria and yeast associated with sugar beet root rot at harvest in the Intermountain West. *Plant Disease*. 92:357-363.
- Strausbaugh, C.A., Rearick, E., Camp, S. 2008. Influence of curly top and Poncho Beta on sugar beet storability. *Journal of Sugarbeet Research*. 45(1-2):31-47.
- Strausbaugh, C.A., Rearick, E., Camp, S., Gallian, J.J., Dyer, A.T. 2008. Influence of beet necrotic yellow vein virus on sugar beet storability. *Plant Disease*. 92:581-587.
- Strausbaugh, C.A., Rearick, E., Eujayl, I.A., Foot, P. 2008. Sugar beet cultivar selection for storability and rhizomania resistance. *Phytopathology*. 98:S152.
- Strausbaugh, C.A., Wintermantel, W.M., Gillen, A.M., Eujayl, I.A. 2008. Curly top survey in the Western United States. *Phytopathology*. 98(11):1212-1217.
- Gillen, A.M., Novy, R.G. 2007. Molecular characterization of the progeny of *Solanum tuberosum* identifies a genomic region associated with resistance to potato leafroll virus. *Euphytica*. 155(3):403-415.
- Gillen, A.M., Strausbaugh, C.A., Gallian, J. 2007. USDA-ARS sugar beet germplasm developed in Salinas, CA, evaluated for rhizomania resistance in Idaho, 2005. *Plant Disease Management Reports*. 1:FC053.
- Singh, S.P., Teran, H., Lema, M., Webster, D.M., Strausbaugh, C.A., Miklas, P.N., Schwartz, H.F., Brick, M.A. 2007. Seventy-five years of breeding dry bean of the Western USA. *Crop Science*. 47:981-989.
- Strausbaugh, C.A., Camp, S. 2007. Verticillium wilt in commercial sugar beet cultivars in Cassia County, ID, 2006. *Plant Disease Management Reports*. 1:V112.
- Strausbaugh, C.A., Camp, S. 2007. Verticillium wilt in experimental sugar beet cultivars in Cassia County, ID, 2006. *Plant Disease Management Reports*. 1:V113.
- Strausbaugh, C.A., Camp, S. 2007. Verticillium wilt in transgenic sugar beet cultivars in Cassia County, ID, 2006. *Plant Disease Management Reports*. 1:V114.
- Strausbaugh, C.A., Gillen, A.M. 2007. Sugarbeet root rot in the Intermountain West [abstract]. *Journal of Sugarbeet Research*. 44:154.

- Strausbaugh, C.A., Gillen, A.M., Gallian, J.J., Tindall, K., Camp, S., Stander, J.R. 2007. Influence of host resistance and insecticide seed treatments on curly top in sugarbeets [abstract]. *Journal of Sugarbeet Research*. 44:162.
- Strausbaugh, C.A., Gillen, A.M., Camp, S., Shock, C.C., Eldredge, E.P., Gallian, J.J. 2007. Relationship of beet curly top foliar ratings to sugar beet yield. *Plant Disease*. 91:1459-1463.
- Strausbaugh, C.A., Rearick, E., Camp, S., Gallian, J.J., Dyer, A.T. 2007. Sugar beet storability and the influence of beet necrotic yellow vein virus. *American Phytopathology Society*. 97:S111.
- Strausbaugh, C.A., Gillen, A.M., Gallian, J.J., Camp, S., Stander, J.R. 2006. Influence of host resistance and insecticide seed treatments on curly top in sugar beets. *Plant Disease*. 90(12):1539-1544.
- Strausbaugh, C.A., Gillen, A.M., Gallian, J.J., Camp, S., Stander, J.R. 2006. Influence of host resistance and insecticide seed treatments on curly top in sugar beets [abstract]. *Phytopathology*. 96:S111.
- Strausbaugh, C.A., Windes, J.M. 2006. Influence of subsoiling and direct seeded cereals in southeastern Idaho. *Canadian Journal of Plant Pathology*. 28:596-608.

5402-21000-007-00D

PRESERVATION AND QUALITY ASSESSMENT OF PLANT GENETIC RESOURCES – David Ellis (P), David Dierig, and Maria Jenderek; Fort Collins, Colorado

- Ellis, D.D., Campbell, K., Grotenhuis, J.A., Jenderek, M.M., Pedersen, J.F. 2010. Crop Registration: The Pathway to Public Access of Plant Genetic Materials to Build Crops for the Future. *Crop Science*. 50:1151-1160.
- Jenderek, M.M., Hannan, R.M. 2009. Diversity in seed production characteristics within the USDA-ARS *Limnanthes alba* germplasm collection. *Crop Science*. 49:1387-1394.
- Park, J., Canam, T., Kang, K., Ellis, D.D., Mansfield, S.D. 2008. Over-expression of an arabidopsis family A sucrose phosphate synthase (SPS) gene alters plant growth and fibre development. *Transgenic Research* 17:181-192.
- Donnelly, L.M., M.M. Jenderek J.P. Prince, P.A. Reeves A. Brown and R.M. Hannan. 2008. Genetic diversity in the USDA *Limnanthes* germplasm collection assessed by simple sequence repeats. *Plant Genetic Resources: Characterization and Utilization* 1:33-41.
- Nass, L.L., Pereira, P.A., Ellis, D.D. 2007. Biofuels in Brazil: An overview. *Crop Science* 47:2228-2237
- Canam, T., Park, J., Yu, K.Y., Campbell, M.M., Ellis, D.D., Mansfield, S.D. 2006. Varied growth, biomass and cellulose content in tobacco expressing yeast-derived invertases. *Planta* 224:1315-1327.

5402-21000-012-00D

RESEARCH TO DEVELOP STRATEGIES AND TECHNOLOGIES FOR PRESERVING PLANT GENETIC DIVERSITY IN EX SITU GENE BANKS – Christina Walters (P), David Dierig, Gayle Volk, and Christopher Richards; Fort Collins, Colorado

- Walters, C.T., Ballesteros, D., Vertucci, V. 2011. Structural mechanics of seed deterioration: Standing the test of time. *Plant Science*. 179:565-573.
- Gross, B.L. 2011. MADS-box out of the black box. *Molecular Ecology*. 20:25-26.
- Mira, S., Elena Gonzalez-Benit, M., Hill, L.M., Walters, C.T. 2010. Characterization of Volatile Production during Storage of Lettuce (*Lactuca sativa*) Seed. *Journal of Experimental Botany*. 61:3915-3924.
- Niedzielski, M., Walters, C.T., Luczak, W., Hill, L.M., Wheeler, L.J., Puchalski, J. Assessment of Variation in Seed Longevity within Rye, Wheat and the Intergeneric Hybrid Triticale. *Seed Science Research*. 19:213-224.
- Reeves, P.A., Richards, C.M. 2010. Species delimitation under the general lineage concept: An empirical example using wild North American hops (cannabaceae: *Humulus lupulus*). *Systematic Biology*. 60:45-59.
- Richards, C.M., Lockwood, D.R., Volk, G.M., Walters, C.T. 2010. Modeling demographics and genetics in ex situ collections during seed storage and regeneration. *Crop Science*. 50:2440-2447.
- Richards, C.M., Volk, G.M. 2010. New Challenges for Data Management in Genebanks. *Acta Horticulturae*. 859:333-336.
- Volk, G.M. 2010. Advantages for the use of standardized phenotyping in databases. *HortScience* 45:1310-1313.
- Volk, G.M. 2010. Application of functional genomics and proteomics to plant cryopreservation. *Current Genomics*. 11:24-29.
- Volk, G.M., and Bonnard, R.M. 2010. Increased efficiency using the encapsulation-dehydration cryopreservation technique for *Arabidopsis thaliana*. *CryoLetters*. 31:95-100.
- Volk, G.M., Richards, C.M., Forsline, P.L. 2010. A comprehensive approach toward conserving *Malus* germplasm. *Acta Horticulturae*. 859:177-182.
- Volk, G.M., Bonnard, R.M., Waddell, J.W., Widrechner, M.P. 2009. Cryopreservation Of Dormant Buds From Diverse *Fraxinus* Species. *CryoLetters*. 30:262-267.

- Richards, C.M., Volk, G.M., Reeves, P.A., Reilley, A., Henk, A.D., Forsline, P.L., Aldwinckle, H. 2009. Selection of stratified core sets representing wild apple (*Malus sieversii*). *Journal of the American Society for Horticultural Science*. 134:228-235.
- Routson, K., Reilley, A., Henk, A.D., Volk, G.M. 2009. Identification of historic apple trees in the Southwestern United States and implications for conservation. *HortScience* 44:589-594.
- Reeves, P.A., Richards, C.M. 2009. Accurate inference of subtle population structure (and other genetic discontinuities) using principal coordinates. *Plos one*. 4(1) e4269.
- Richards C.M., G. Volk, A.A. Reilley, A.D. Henk, D. Lockwood, P.A. Reeves, and P.L. Forsline. 2009. Genetic diversity and population structure in *Malus sieversii*, a wild progenitor species of domesticated apple. *Tree Genetics and Genomics* 5:339-347.
- Volk, G.M., Richards, C.M., Henk, A.D., Reilley, A., Miller, D.D., Forsline, P.L. 2009. Novel diversity identified in a wild apple population from Kyrgyz Republic. *HortScience* 44:516-519.
- Volk, G.M., Richards, C.M., Henk, A.D., Reilley, A.A., Reeves, P.A., Forsline, P.L., Aldwinckle, H.S. 2009. Capturing The Diversity Of Wild *Malus orientalis* From Georgia, Armenia, Russia And Turkey. *Journal of the American Society for Horticultural Science*. 134:453-459.
- Volk, G.M., Stern, D. 2009. Phenotypic Characteristics Of Ten Garlic Cultivars Grown At Different North American Locations. *HortScience*. 44:1238-1247.
- Volk, G.M., Waddell, J.W., Towill, L., Grauke, L.J. 2009. Variation in low temperature exotherms of pecan cultivar dormant twigs. *HortScience*. 44:317-321.
- Kothera, L., Zimmerman, E.M., Richards, C.M., Savage, H.M. 2009. Microsatellite characterization of subspecies and their hybrids in *Culex pipiens* complex mosquitoes along a north-south transect in the central United States of America. *Journal of Medical Entomology* 46:236-248.
- Rai, H.S., Reeves, P.A., Peakal, R., Olmstead, R.G., Graham, S.W. 2008. Inference of higher-order conifer relationships from a multi-locus plastid data set. *Botany* 86:658-669.
- Volk, G.M., Richards, C.M. 2008. Availability of genotypic data for USDA-ARS National Plant Germplasm System accessions using the Genetic Resources Information Network (GRIN) database. *HortScience* 43:1365-1366.
- Volk, G.M., Richards, C.M., Reilley, A., Henk, A.D., Reeves, P.A., Forsline, P.L., Aldwinckle, H. 2008. Genetic diversity and disease resistance of wild *Malus orientalis* from Turkey and southern Russia. *Journal of the American Society for Horticultural Science* 133:383-389.
- Volk, G.M., Waddell, J.W., Bonnart, R.M., Towill, L.E., Ellis, D.D., Luffman, M. 2008. High viability of dormant *Malus* buds after 10 years of storage in liquid nitrogen vapor. *CryoLetters* 29:89-94.
- Walters, C.T., Volk, G.M., Richards, C.M. 2008. Genebanks in the post-genomic age: emerging roles and anticipated uses. *Biodiversity Issues* 9:68-71.
- Ballesteros, D., Walters, C.T. 2007. Calorimetric properties of water and triacylglycerols in fern spores relating to storage at cryogenic temperatures. *Cryobiology* 55:1-9.
- Ballesteros, D., Walters, C.T. 2007. Water properties in fern spores: sorption characteristics relating to water affinity, glassy states and storage stability. *Journal of Experimental Botany* 58:1185-1196.
- Kothera, L., Richards, C.M., Carney, S.E. 2007. Genetic diversity and structure in the rare Colorado endemic plant *Physaria bellii* Mulligan (Brassicaceae). *Conservation Genetics* 8:1043-1050.
- Volk, G.M., Crane, J., Caspersen, A.M., Kovach, D.A., Gardner, C.A., Walters, C.T. 2007. Hydration of *Cuphea* seeds containing crystallized triacylglycerols. *Functional Plant Biology* 34:360-367.
- Walters, C. 2007. Materials used for seed storage containers: Response. *Seed Science Research* 17:233-242.
- Eira, M.T.S., E.A. da Silva, R.D. de Castro, S. Dussert, C. Walters, J.D. Bewley and H.W.M. Hilhorst. 2006. Coffee seed physiology. *Brazilian Journal of Plant Physiology* 18:149-163.
- Volk, G.M., Richards, C.M., Henk, A.D., Reilley, A., Bassil, N.V., Postman, J.D. 2006. Diversity of wild *pyrus communis* based on microsatellite analyses. *Journal of the American Society for Horticultural Science*. Vol. 131:408-417.

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NONCHEMICAL PEST CONTROL AND ENHANCED SUGAR BEET GERmplasm VIA TRADITIONAL AND MOLECULAR TECHNOLOGIES – Leonard Panella (P) and Kimberly Webb; Fort Collins, Colorado

- Webb, K.M., Hill, A.L., Laufman, J., Hanson, L.E., Panella, L.W. 2011. Long-term Preservation of a Collection of *Rhizoctonia solani*, using Cryogenic Storage. *Annals of Applied Biology*. 158(3):297-304.
- Panella, L.W., Strausbaugh, C.A. 2011. Beet Curly Top Resistance of USDA-ARS National Plant Germplasm System Plant Introductions, 2009. *Plant Disease Management Reports*. 5:FC066. Online publication doi:10.1094/PDMR05.
- Panella, L.W., Strausbaugh, C.A. 2011. Beet Curly Top Resistance of USDA-ARS National Plant Germplasm System Plant Introductions, 2010. *Plant Disease Management Reports*. 5:FC066. Online publication doi:10.1094/PDMR05.

- Panella, L.W., Vagher, T.O., Fenwick, A.L., Webb, K.M. 2011. Rhizoctonia Crown and Root Rot Resistance of Beta Plant Introductions from the USDA, Agricultural Research Service's National Plant Germplasm System, 2010. Plant Disease Management Reports. 5:FC067. Online publication doi:10.1094/PDMR05.
- Stevanato, P., Zavalloni, C., Marchetti, R., Bertaggia, M., Saccomani, M., Mcgrath, J.M., Panella, L.W., Biancardi, E. 2010. Relationship Between Subsoil Nitrogen Availability and Sugar Beet Processing Quality. Agronomy Journal: 102 (1)17-22.
- Panella, L.W., Fenwick, A.L., Hill, A.L., Vagher, T.O., Webb, K.M. 2010. Rhizoctonia Crown and Root Rot Resistance of Beta PI's from the USDA-ARS NPGS, 2009. Plant Disease Management Reports. 4:FC004
- Panella, L.W., Strausbaugh, C.A. 2010. Beet Curly Top Resistance of Released Germplasm from the USDA-ARS Program at Fort Collins, CO, 2009. Plant Disease Management Reports. 4:FC071 DOI:10.1094/PDMR04.
- Webb, K.M., Garcia, E., Vera Cruz, C.M., Leach, J.E. 2010. Influence of Rice Development on the Function of Bacterial Blight Resistance Genes. European Journal of Plant Pathology. 128:399-407 DOI 10.1007/s10658-010-9668-z.
- Webb, K.M., Ona, I., Bai, J., Garrett, K.A., Mew, T., Vera Cruz, C.M., Leach, J.E. 2010. A Benefit of High Temperature: Increased Effectiveness of a Rice Bacterial Blight Disease Resistance Gene. New Phytologist. 185: 568-576
- Hanson, L.E., Hill, A.L., Jacobsen, B.J., Panella, L.W. 2009. Varying Response of Sugar Beet Lines to Different Fusarium Oxysporum F. sp. Betae Isolates from the United States. Journal of Sugarbeet Research. 46(1):11-26.
- Panella, L.W., Wheeler, L.J., McClintock, M.E. 2009. Long-term Survival of Cryopreserved Sugar Beet Pollen. Journal of Sugar Beet Research. Vol.46 No.1 pp 1-9.
- Larson, R.L., Wintermantel, W.M., Hill, A.L., Fortis, L.L., Nunez, A. 2008. Proteome changes in sugar beet in response to Beet necrotic yellow vein virus. Physiological and Molecular Plant Pathology. doi:10.1016/j.pmpp.2008.04.003
- Panella, L.W., Fenwick, A.L., Hill, A.L., McClintock, M.E., Vagher, T.O. 2008. Rhizoctonia root rot resistance of Beta PIs from the USDA-ARS NPGS, 2007. Plant Disease Management Reports. (online) 2:V057.DOI:10.1094/PDMR02. The American Phytopathological Society. St. Paul, MN.
- Panella, L.W., Hanson, L.E., Lewellen, R.T. 2008. Breeding for Multiple Disease Resistance in Sugarbeet: Registration of FC220 and FC221. Journal of Plant Registrations.
- Larson, R.L., Hill, A.L., Nunez, A. 2007. Characterization of Protein Changes Associated with Sugar Beet (*Beta vulgaris*) Resistance and Susceptibility to *Fusarium oxysporum*. Journal of Agricultural and Food Chemistry. 55:7905-7915.
- Hanson, L.E., Lewellen, R.T. 2007. Stalk rot of sugar beet caused by *Fusarium solani* on the Pacific coast. Plant Disease. 91:1204.
- Hanson, L.E., Panella, L.W. 2007. Rhizoctonia root rot resistance of Beta PIs from the USDA-ARS NPGS, 2006. Plant Disease Management Reports. (online) 1:V023. DOI: 10.1094/PDMR01. The American Phytopathological Society. St. Paul, MN.
- Larson, R.L., Hill, A.L., Fenwick, A.L., Kniss, A., Hanson, L.E., Miller, S. 2006. Influence of glyphosate on rhizoctonia and fusarium root rot in sugar beet. Pest Management Science 62:1182-1192 (2006); DOI: 10.1002/ps.1297
- Hanson, L.E. 2006. First report of *Fusarium* yellows of sugar beet caused by *F. oxysporum* in Michigan. Plant Disease. DOI:10.1094
- Panella, L.W., Hanson, L.E. 2006. Registration of FC723 and FC723 CMS Monogerm Sugarbeet Germplasm Resistant to Rhizoctonia Root Rot and moderately Resistant to *Cercospora* Leaf Spot. Journal of Plant Registrations. 1:66-67 (2007) doi: 10.3198/jpr2006.12.0771crg

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GENETIC ENHANCEMENT FOR RESISTANCE TO BIOTIC AND ABIOTIC STRESSES IN HARD WINTER WHEAT – Robert Bowden (P), Ming Shum, Guihua Bai, and John Fellers; Manhattan, Kansas

- Bernardo, A., Ma, H., Zhang, D., Bai, G. 2011. Single nucleotide polymorphism in wheat chromosome region harboring *Fhb1* for fusarium head blight resistance. Molecular Breeding. DOI.10.1007/s11032-011-9565-Y.
- Jayatilake, D.V., Bai, G., Dong, Y. 2011. A novel quantitative trait locus for fusarium head blight resistance in chromosome 7A of wheat. Journal of Theoretical and Applied Genetics. 122:1189-1198.
- Khajuria, C., Buschman, L.L., Chen, M., Zurek, L., Zhu, K. 2011. Characterization of six antibacterial response genes from the European corn borer (*Ostrinia nubilalis*) larval gut and their expression in response to bacterial challenge. Journal of Insect Physiology. 57:345-355.
- Li, T., Bai, G., Wu, S. 2011. Quantitative trait loci for resistance to fusarium head blight in a chinese wheat landrace haiyanzhong. Theoretical and Applied Genetics. DOI10.1007/S00122-011-1549-0.

- Li, X., Zhu, C.L., Lin, Z., Wu, Y., Zhang, D., Bai, G., Song, W., Ma, J., Muehlbauer, G.J., Scanlon, M.J., Zhang, M., Yu, J. 2011. Chromosome size in diploid eukaryotic species centers on the average length with a conserved boundry. *Molecular Biology and Evolution*. DOI 10.1093/molbev/msr011.
- Seifers, D.L., Martin, J.T., Fellers, J.P. 2011. Occurrence and yield effects of wheat infected with Triticum mosaic virus. *Plant Disease*. 95:183-188.
- Yu, X., Bai, G., Luo, N., Cheng, Z., Liu, S., Liu, J., Warnke, S.E., Jiang, Y. 2011. Association of simple sequence repeat (SSR) markers with submergence tolerance in diverse population of perennial ryegrass. *Plant Science*. 180:391-398.
- Liu, S., Bai, G. 2010. Dissection and Fine Mapping of a Major QTL for Preharvest Sprouting Resistance in White Wheat Rio Blanco. *Theoretical and Applied Genetics*. DOI 10.1007/s00122-010-1396-4.
- Liu, S., Bai, G., Cai, S., Chen, C. 2010. Dissection of Genetic Components of Preharvest Sprouting Resistance in White Wheat. *Molecular Breeding*. DOI:10.1007/s11032-010-9448-7.
- Cainong, J., Chen, M., Johnson, J., Friebe, B., Gill, B., Lukaszewski, A., Zavatsky, L. 2010. Wheat-rye recombinants T2BS.2BL-2RL conferring resistance to Hessian fly (H21). *Crop Science*. 50:920-925.
- Chen, J., Yu, J., Ge, L., Wang, H., Berbel, A., Liu, Y., Chen, Y., Li, G., Tadege, M., Wen, J., Cosson, V., Mysore, K.S., Ratet, P., Madueno, F., Bai, G., Chen, R. 2010. Control of Dissected Leaf Morphology by a Cys(2)His(2) Zinc Finger Transcription Factor in the Model Legume *Medicago Truncatula*. *Proceedings of the National Academy of Sciences*. 107:10754-10759.
- Chen, M., Liu, X., Zhao, H., Stuart, J.J., Shukle, R.H., Yang, Z., Hulbert, S. 2010. Unconventional Conservation Among Genes Encoding Small Secreted Salivary Gland Proteins From a Gall Midge. *BMC Evolutionary Biology*. 10:296.
- Liu, X., Williams C. E., Nemacheck, J. A., Wang, H., Subramanyam, S., Zheng, C., Chen, M.-S. 2010. Reactive Oxygen Species are Involved in Plant Defense Against a Gall Midge. *Plant Physiology*. 152:985-999.
- Khajuria, C., Buschman, L.L., Chen, M., Muthukrishnan, S., Zhu, K. 2010. A New Gut-Specific Chitinase Gene Essential for Regulation of Chitin Content of Peritrophic Matrix and Growth of *Ostrinia Nubilalis* Larvae. *Insect Biochemistry and Molecular Biology*. 40:621-629.
- Price, J.A., Smith, J.T., Simmons, A., Fellers, J.P., Rush, C.M. 2010. Multiplex Real Time PCR For Detection of Wheat Streak Mosaic Virus and Triticum Mosaic Virus. *Journal of Virological Methods*. 165: 198-201.
- Seifers, D., Martin, J.T., Fellers, J.P. 2010. An Experimental Host Range of Triticum Mosaic Virus. *Plant Disease*. 94:1125-1131.
- Singh, S., Bowden, R.L. 2010. Molecular Mapping of Adult-Plant Race-Specific Leaf Rust Resistance Gene Lr12 in Bread Wheat. *Molecular Breeding*. DOI 10.1007/s11032-010-9467-4.
- Smith, C., Liu, X., Wang, L., Liu, X., Chen, M., Starkey, S., Bai, J. 2010. Aphid Feeding Activates Expression of a Transcriptome of Oxylipin-Based Defense Signals in Wheat Involved in Resistance to Herbivory. *Journal of Chemical Ecology*. 36:260-276.
- Sun, X., Bai, G., Carver, B.F., Bowden, R.L. 2010. Molecular Mapping of Wheat Leaf Rust Resistance Gene Lr42. *Crop Science*. 50:59-66.
- Voss, H., Bowden, R.L., Leslie, J.F., Miedaner, T. 2010. Variation and Transgression of Aggressiveness Among Two *Gibberella Zeae* Crosses Developed from Highly Aggressive Parental Isolates. *American Phytopathology Society*. 10:1094/PHYTO-100-9-0904.
- Yu, J., Bai, G. 2010. Mapping Quantitative Trait Loci for Long Coleoptile in Chinese Wheat Landrace Wangshuibai. *Crop Science*. 50:59-66.
- Zhang, D., Bai, G., Zhu, C., Yu, J., Carver, B. 2010. Genetic Diversity, Population Structure and Linkage Disequilibrium in U.S. Elite Winter Wheat (*Triticum Aestivum* L.). *The Plant Genome*. 3:117-127.
- Zhang, S., Shukle, R. H., Mittapalli, O., Zhu, Y. C., Reese, J. C., Wang, H., Hua, B.Z., Chen, M.S. 2010. The gut transcriptome of a gall midge, *Mayetiola destructor*. *Journal of Insect Physiology*. 56:1198-1206.
- Zhu, L., Liu, X., Chen, M. 2010. Differential Accumulation of Phytohormones in Wheat Seedlings Attacked by Avirulent and Virulent Hessian Fly (Diptera: Cecidomyiidae) Larvae. *Journal of Economic Entomology*. 103:178-185.
- Aggarwal, R., Benatti, T., Gill, N., Chen, M., Schemerhorn, B.J., Fellers, J.P., Stuart, J.J. 2009. A BAC-based physical map of the Hessian fly (*Mayetiola destructor*) genome anchored to polytene chromosomes. *Biomed Central (BMC) Genomics*. 10:293.
- Bohssini, M., Chen, M., Lhaloui, S., Zharmukamedoua, G., Rihawi, F. 2009. Virulence of Hessian Fly (Diptera: Cecidomyiidae) in the Fertile Crescent. *Journal of Applied Entomology*. 133(5):381-385.
- Chen, M., Echegaray, E., Whitworth, J., Wang, H., Sloderbeck, P.E., Knutson, A., Giles, K.L., Royer, T.A. 2009. Virulence Analysis of Hessian Fly (*Mayetiola destructor*) Populations from Texas, Oklahoma, and Kansas. *Journal of Economic Entomology*. 102:774-780.
- Chen, M., Liu, X., Wang, H., El Bouhssini, M. 2009. Hessian fly (*Mayetiola destructor*) interactions with barley, rice, and wheat seedlings. *Journal of Economic Entomology*. 102:1663-1672.
- Fellers, J.P., Seifers, D., Ryba-White, M., Martin, J. 2009. The Complete Sequence of Triticum Mosaic Virus, a New Wheat Infecting Virus of the Great Plains. *Archives of Virology*. 154:1511-1515.

- Li, T., Bai, G. 2009. Lesion Mimic Associates with Adult Plant Resistance to Leaf Rust Infection in Wheat. *Theoretical and Applied Genetics*. 119:13-21.
- Prasad, B., Babar, M.A., Xu, X.Y., Bai, G., Klatt, A.R. 2009. Genetic Diversity in the U.S. Hard Red Winter Wheat Cultivars as Revealed by Microsatellite Markers. *Crop and Pasture Science*. 60:16-24.
- Pumphrey, M.O., Bai, J., Chingcuanco, D.L., Anderson, O.D., Gill, B. 2009. Non-Additive Expression of Homoeologous Genes is Established Upon Polyploidization in Hexaploid Wheat. *Genetics*. 181(3):1147-1157.
- Ristic, Z., Momcilovic, I., Bukovnik, U., Prasad, P., Fu, J., De Ridder, B., Elthon, E., Mladenov, N. 2009. Rubisco activase and wheat productivity under heat stress conditions. *Journal of Experimental Botany*. 60:4003-4014.
- Seifers, D., Martin, J., Harvey, T., Fellers, J.P., Michaud, J.P. 2009. Identification of the Wheat Curl Mite as the Vector of Triticum Mosaic Virus. *Plant Disease*. 93:25-29.
- Sood, S., Kuruparth, V., Bai, G., Gill, B.S. 2009. The Major Threshability Genes Soft Glume (sog) and Tenacious Glume (Tg), of Diploid and Polyploid Wheat, Trace Their Origin to Independent Mutations at Non-Orthogous Loci. *Theoretical and Applied Genetics*. 119:341-351.
- Sood, S., Kuruparth, V., Bai, G., Gill, B.S. 2009. The Major Threshability Genes Soft Glume (sog) and Tenacious Glume (Tg), of Diploid and Polyploid Wheat, Trace Their Origin to Independent Mutations at Non-Orthogous Loci. *Theoretical and Applied Genetics*. 119:341-351.
- Sun, X., Marza, F., Ma, H., Carver, B., Bai, G. 2009. Mapping Quantitative Trait Loci for Quality Factors in an Inter-Class Cross of US and Chinese Wheat. *Theoretical and Applied Genetics*. 120:1041-1051.
- Wu, S., Pumphrey, M.O., Bai, G. 2009. Molecular Mapping of Stem Rust Resistance Gene Sr40 in Wheat. *Crop Science*. 49:1681-1686.
- Huang, L., Brooks, S.A., Li, W., Fellers, J.P., Nelson, J., Gill, B.S. 2009. Evolution of New Disease Specificity at a Single Resistance Locus in a Crop-Weed Complex: Reconstitution of the Lr21 Gene in Wheat. *Genetics* 182:595-602.
- Bowden, R.L., Fuentes-Bueno, I., Leslie, J.F., Lee, J., Lee, Y. 2008. Methods for Detecting Chromosome Rearrangements in *Gibberella Zeae*. *Cereal Research Communications*. 36:609-615.
- Bukovnik, U., Fu, J., Bennett, M., Prasad, P., Ristic, Z. Heat tolerance and expression of protein synthesis elongation factors, EF-Tu and EF-1a, in spring wheat. *Functional Plant Biology*. 36: 234-241.
- Barrett, B., Faris, J.D., Fellers, J.P. 2008. Molecular Mapping of the leaf rust resistance gene Lr17a in Wheat. *Crop Science*. 48:1124-1128.
- Bernardo, A., Bai, G., Guo, P., Xiao, K., Guenzi, A., Ayoubi, P. *Fusarium graminearum* - induced differential gene expression between fusarium head blight-resistant and susceptible wheat cultivars. *Functional and Integrative Genomics* 7:69-77.
- Cai, S., Bai, G., Zhang, D. 2008. Quantitative Trait Loci for Aluminum Resistance in Chinese Wheat Landrace FSW. *Theoretical and Applied Genetics*. 117:49-56.
- Chen, M. 2008. Inducible direct plant defense against insect herbivores - a review. *Insect Science*. 15:101-114.
- Chen, M., Huixian, Z., Zhu, Y., Scheffler, B.E., Liu, X., Liu, X., Hulbert, S., Stuart, J. 2008. Analysis of Transcripts and Proteins Expressed in the Salivary Glands of Hessian Fly (*Mayetiola destructor*) Larvae. *Journal of Insect Physiology*. 54:1-16.
- Hu, S., Bai, G., Carver, B., Zhang, D. 2008. Diverse Origins of Aluminum-resistant Sources in Wheat. *Theoretical and Applied Genetics*. DOI 10.1007/500122-008-0874-4.
- Liu, S., Cai, S., Graybosch, R.A., Chen, C., Bai, G. 2008. Quantitative trait loci for resistance to pre-harvest sprouting in U.S. hard white winter wheat. *Journal of Theoretical and Applied Genetics*. 117:691-699.
- Mutti, N.S., Louis, J., Pappan, L.K., Pappan, K., Begum, K., Chen, M., Park, Y., Dittmer, N., Marshall, J., Reese, J.C., Reeck, G.R. 2008. A protein from the salivary glands of the pea aphid, *Acyrtosiphon pisum*, is essential in feeding on a host plant. *Proceedings of the National Academy of Sciences*. 105:9965-9969.
- Qi, L., Pumphrey, M.O., Friebe, B., Chen, P., Gill, B. 2008. Molecular cytogenetic characterization of alien introgressions with gene *Fhb3* for resistance to *Fusarium* head blight disease of wheat. *Journal of Theoretical and Applied Genetics*. 117:1155-1166.
- Seifers, D.L., Martin, T., Harvey, T.J., Fellers, J.P., Stack, J., Ryba-White, M., Haber, S., Krokhn, O., Spicer, V., Lovat, N., Yamchuk, A., Standing, K.G. 2008. Triticum Mosaic Virus: A New Virus Isolated From Wheat in Kansas. *Plant Disease*. 92:808-817.
- Singh, S., Bockus, W., Sharma, I., Bowden, R.L. 2008. A novel source of resistance in wheat to *pyrenophora tritici-repentis* race 1. *Plant Disease*. 92:91-95.
- Wu, J., Liu, X., Zhang, S., Zhu, Y., Whitworth, R., Chen, M. 2008. Differential Responses of Wheat Inhibitor-Like Genes to Hessian Fly (*Mayetiola destructor*) Attacks during Compatible and Incompatible Interactions. *Journal of Chemical Ecology*. 34:1005-1012.
- Yu, J., Bai, G., Cai, S., Dong, Y., Ban, T. 2008. New FHB-resistant Sources from Asian Wheat Germplasm. *Crop Science*. 48:1090-1097.
- Ristic, Z., Bukovnik, U., Prasad, P., West, M.S. 2008. A model for prediction of heat stability of photosynthetic membranes. *Crop Science*. 48:1513-1522.

- Lee, J., Jurgenson, J.E., Leslie, J.F., Bowden, R.L. 2008. Alignment Between Genetic and Physical Maps of *Gibberella zeae*. *Applied and Environmental Microbiology*. 74:2349-2359.
- Lee, J., Leslie, J.F., Bowden, R.L. 2008. Expression and Function of Sex Pheromones and Receptors in the Homothallic Ascomycete *Gibberella zeae*. *Eukaryotic Cell*. 1211-1221.
- Prasad, V., Pisipati, S., Ristic, Z., Bukovnik, U., Fritz, A.K. 2008. Impact of Nighttime Temperature on Physiology and Growth of Spring Wheat. *Crop Science*. 48:2372-2380.
- Fellers, J.P. 2008. Genome filtering using methylation-sensitive restriction enzymes with six-base pair recognition sites. *The Plant Genome*. 1:2.
- Fu, J., Momcilovic, I., Clemente, T., Nersesian, N., Trick, H., Ristic, Z. 2008. Heterologous expression of a plastid EF-Tu reduces protein thermal aggregation and enhances CO₂ fixation in wheat following heat stress. *Plant Molecular Biology*. 68:277-288.
- Das, M.K., Bai, G., Kazi, M. 2007. Genetic diversity in conventional and synthetic wheats with drought and salinity tolerance based on AFLP. *Canadian Journal of Plant Science*. 87:691-702.
- Zhou, L., Bai, G., Carver, B. 2007. Identification of New Sources of Aluminum Resistance in Wheat. *Plant and Soil Journal*. DOI: 10.1007/s11104-007-9324-3.
- Zhu, L., Liu, X., Liu, X., Jeannotte, R., Reese, J., Harris, M., Stuart, J., Chen, M. 2007. Hessian Fly (*Mayetiola Destructor*) Attack Causes Dramatic Shift in Carbon/Nitrogen Metabolism in Wheat. *Molecular Plant-Microbe Interactions*. 21:70-78.
- Hulbert, S.H., Bai, J., Fellers, J.P., M. G., P., Bowden, R.L. 2007. Gene expression patterns of wheat rust resistance gene *Lr34/Yr18* indicate novel mode of action. *Phytopathology* 97:1083-1093.
- Yu, J., Bai, G., Zhou, W., Dong, Y., Kolb, F. 2007. Quantitative Trait Loci for *Fusarium* Head Blight Resistance in Chinese Wheat Landrace Wangshuibai. *Phytopathology*. 98:87-94
- Ristic, Z., Bukovnik, U., Momcilovic, I., Fu, J., Prasad, V. 2007. Heat-induced Accumulation of Chloroplast Protein Synthesis Elongation Factor, EF-TU, in Winter Wheat. *Journal of Plant Pathology*. DOI:10.1016/j.jplph.2007.03.003.
- Ristic, Z., Bukovnik, U., Prasad, V. 2007. Correlation between heat stability of thylakoid membranes and loss of chlorophyll in winter wheat under heat stress. *Crop Science*. 47:2067-2073.
- Ristic, Z., Momcilovic, I., Fu, J., Callegari, E., Deridder, B.P. 2007. Chloroplast EF-Tu and thermal aggregation of Rubisco activase. *Journal of Physiology*. 164:1564-1571.
- Liu, X., Bai, J., Zhu, L., Liu, X., Weng, N., Reese, J.C., Harris, M., Stuart, J.J., Chen, M. 2007. Differential Gene Expression of H9 and H13 Wheat Genotypes during Attack by Virulent and Avirulent Hessian Fly (*Mayetiola destructor*) Larvae. *Journal of Chemical Ecology*. 33:2171-2194.
- Guo, P., Bai, G., Ronghua, L., Carver, B., Baum, M. 2007. Molecular characterization of Atlas 66 derived wheat near-isogenic lines contrasting in Aluminum (Al) tolerance. *Science in China*.
- Kuraparthi, V., Chhuneja, P., Dhaliwal, H., Kaur, S., Bowden, R.L., Gill, B. 2007. Characterization and mapping of cryptic alien introgression from *Aegilops geniculata* with new leaf rust and stripe rust resistance genes *Lr57* and *Yr40* in wheat. *Theoretical and Applied Genetics*. 114:1379-1389.
- Kuraparthi, V., Sood, S., Chhuneja, P., Dhaliwal, H., Kaur, S., Bowden, R.L., Gill, B. 2007. A cryptic wheat-*Aegilops triuncialis* translocation with leaf rust resistance gene *Lr58*. *Crop Science*. 47:1995-2003 (2007).
- Chen, C., Cai, S., Bai, G. 2007. A major QTL controlling seed dormancy and pre-harvest sprouting resistance on chromosome 4A in a Chinese wheat landrace. *Molecular Breeding*. 21:351-358.
- Chen, M., Liu, X., Zhu, Y., Reese, J.E., Wilde, G.E. 2006. Genes encoding a group of related small secreted proteins from the gut of hessian fly larvae [*mayetiola destructor* (say)]. *Journal of Insect Science* 13, 339-348.
- Guo, P., Bai, G., Carver, B., Li, R., Bernardo, A., Baum, M. 2006. Transcriptional analysis between two wheat near-isogenic lines contrasting in aluminum (al) tolerance under al stress. *Molecular Genetics and Genomics*. 277:1-12.
- Lobo, N.F., Behura, S.K., Aggarwal, R., Chen, M., Collins, F.H., Stuart, J.J. 2006. Genomic analysis of a 1 mb region near the telomere of hessian fly chromosome x2 and avirulence gene *vh13*. *Biomed Central (BMC) Genomics* 7:7.
- Ma, H., Bai, G., Gill, G., Hart, L. 2006. Deletion of a chromosome arm altered wheat resistance to *fusarium* head blight and deoxynivalenol accumulation in chinese spring. *Plant Disease* 90:1545-1549.
- Ma, H., Bai, G., Zhang, X., Lu, W. 2006. Main effects, epistasis and enviromental interactions of qtls for *fusarium* head blight resistance in a recombinant inbred population. *Phytopathology* 96:534-541
- Ma, H., Zhang, K., Gao, L., Bai, G., Chen, H., Cai, Z., Lu, W. 2006. Qtl for the resistance to wheat *fusarium* head blight and deoxynivalenol accumulation in wangshuibai under field conditions. *Plant Pathology*. DOI:10.1111/J.1365-3059.2006.01447.x
- Zhao, H., Liu, X., Chen, M. 2006. H22, a major resistance gene to the hessian fly (*mayetiola destructor*), is mapped to the distal region of wheat chromosome 1ds. *Theoretical and Applied Genetics*.
- Wu, Y., Taliaferro, C., Bai, G., Martin, D., Anderson, J., Anderson, M., Edwards, R. 2006. Genetic analyses of chinese cynodon accessions by flow cytometry and aflp markers. *Crop Science* 46:917-926
- Yu, J., Bai, G., Cai, S., Ban, T. 2006. Marker-assisted characterization of asian wheat lines for resistance to *fusarium* head blight. *Theoretical and Applied Genetics*. 113:308-320.

Webb, C.A., Fellers, J.P. 2006. Cereal rust genomics and the pursuit of virulence and avirulence factors. *FEMS Immunology and Medical Microbiology*.

5440-21000-029-00D

GENETIC IMPROVEMENT AND EVALUATION OF HARD WINTER AND SPRING WHEATS – Robert Graybosch (P); Lincoln, Nebraska

- Baenziger, P.S., Graybosch, R.A., Nelson, L.A., Regassa, T., Klein, R.N., Baltensperger, D.D., Santra, D., Ibrahim, A., Chen, M., Bai, G. 2011. Registration of 'NH03614 CL' Wheat. *Journal of Plant Registrations*. 5:75-80.
- Hansen, L.E., Jackson, D.S., Wehling, R.L., Graybosch, R.A. 2010. Functionality of Chemically Modified Wild-Type, Partial Waxy and Waxy Starches from Tetraploid Wheats. *Journal of Cereal Science*. Volume 51: 409-414.
- Hansen, L.E., Jackson, D.S., Wehling, R.L., Wilson, J.D., Graybosch, R.A. 2010. Functionality of Native Tetraploid Wheat Starches: Effects of Waxy Loci Alleles and Amylose Concentration in Blends. *Journal of Cereal Science*. 52:39-45.
- Graybosch, R.A., Seabourn, B.W., Chen, Y.R., Blechl, A.E. 2010. Quality and agronomic effects of three high-molecular-weight glutenin subunit transgenic events in winter wheat. *Cereal Chemistry*. 88(1):95-102.
- Graybosch, R.A., Peterson, C.J. 2010. Genetic Improvement in Winter Wheat Yields in the Great Plains of North America, 1959-2008. *Crop Science* Volume 50:1882-1890.
- Graybosch, R.A., Peterson, C.J., Baenziger, P.S., Baltensperger, D.D., Nelson, L.A., Jin, Y., Kolmer, J.A., Seabourn, B.W., French, R.C., Hein, G.L. 2009. Registration of 'Mace' Hard Red Winter Wheat. *Journal of Plant Registrations* 3:51-56.
- Guan, L. P. A. Seib, R. A. Graybosch, S. Bean and Yong-Cheng Shi. 2009. Dough rheology and wet milling of hard waxy wheat flours. *J. Agric. Food Chem.*, Publication Date (Web): July 13, 2009 (web version) DOI: 10.1021/jf900438v.
- Graybosch, R.A., Baltensperger, D. 2009. Evaluation of the waxy endosperm trait in proso millet (*Panicum mileaceum* L.). *Plant Breeding* 128, 70-73 (2009).
- Baenziger, P.S., Graybosch, R.A., Nelson, L.A., Klein, R.N., Baltensperger, D.D., Xu, L., Wegulo, S.N., Watkins, J.E., Jin, Y., Kolmer, J.A., Hatchett, J.H., Chen, M., Bai, G. 2009. Registration of 'Camelot' Wheat. *Journal of Plant Registrations*. 3:256-263.
- Baenziger, P., Beecher, B., Graybosch, R.A., Ibrahim, A., Nelson, L., Jin, Y., Wegulo, S., Watkins, J., Chen, M., Bai, G. 2008. Registration of 'NE01643' Wheat. *Journal of Plant Registrations* 2:36-42.
- Heyduck, R., Baltensperger, D., Nelson, L., Graybosch, R.A. 2008. Yield and Agronomic Traits of Waxy Proso in the Central Great Plains. *Plant Breeding* 48:741-748.
- Saito, M., Vrinten, P., Ishikawa, G., Graybosch, R.A., Nakamura, T. 2008. A novel codominant marker for selection of the null Wx-B1 allele in wheat breeding programs. *Molecular Breeding* 23:209-217 (2009).
- Mishra, R., Baenziger, P.S., Russell, W.K., Graybosch, R.A., Baltensperger, D.D., Eskridge, S.M. 2006. Crossover interactions for grain yield in multienvironmental trials of winter wheat.
- Nash, D., Lanning, S., Fox, P., Martin, J., Blake, N., Souza, E., Graybosch, R.A., Giroux, M., Talbert, L. 2006. Relationship of Dough Extensibility to Dough Strength in a Spring Wheat Cross. *Cereal Chemistry* 83:255-258.
- Sahlstrom, S., Baevre, A., Graybosch, R.A. 2006. Impact of Waxy, Partial Waxy, and Wildtype Wheat Starch Fraction Properties on Hearth Bread Characteristics. *Cereal Chemistry* 83: 647-654.

5440-21220-027-00D

ENHANCEMENT OF SORGHUM FOR BIOENERGY, FEED, AND FOOD VALUE – Jeffrey Pedersen (P), Scott Sadler, and Deanna Funnell-Harris; Lincoln, Nebraska

- Funnell-Harris, D.L., Pedersen, J.F., Sattler, S.E. 2010. Alteration in Lignin Biosynthesis Restricts Growth of Fusarium Species in Brown Midrib Sorghum. *Phytopathology* 100:671-681.
- Funnell-Harris, D.L., Pedersen, J.F., Sattler, S.E. 2010. Soil and Root Populations of Fluorescent *Pseudomonas* spp. Associated with Seedlings and Field-grown Plants are Affected by Sorghum Genotype. *Plant and Soil Journal*. Volume 335: 439-455.
- Griess, J., Mason, S., Jackson, D., Galusha, T., Yaseen, M., Pedersen, J.F. 2010. Environment and Hybrid Influences on Food-Grade Sorghum Grain Yield and Hardness. *Crop Science*. 50: 1480-1489.
- Pfeiffer, T.W., Bitzer, M.J., Toy, J.J., Pedersen, J.F. 2010. Heterosis in Sweet Sorghum and Selection of a New Sweet Sorghum Hybrid for Use in Syrup. *Crop Science*. doi: 10.2135/cropsci2009.09.0475.

- Sahoo, L., Schmidt, J.J., Pedersen, J.F., Lee, D.L., Lindquist, J.L. 2010. Growth and fitness components of wild X cultivated *Sorghum bicolor* (Poaceae) hybrids in Nebraska. *American Journal of Botany*. 97(10): 1610-1617.
- Sattler, S.E., Funnell-Harris, D.L., Pedersen, J.F. 2010. Brown Midrib Mutations and Their Importance to the Utilization of Grasses. *Plant Science*. 178:229-238.
- Sattler, S.E., Funnell-Harris, D.L., Pedersen, J.F. 2010. Efficacy of Singular and Stacked Brown Midrib 6 and 12 in Modification of Lignocellulose and Grain Chemistry. *Journal of Agricultural and Food Chemistry*. 58:3611-3616.
- Wong, J.H., Marx, D.B., Wilson, J.D., Lemaux, P.G., Buchanan, B.B., Pedersen, J.F. 2010. Principal component analysis and biochemical characterization of protein and starch reveal primary targets for improving sorghum grain. *Plant Science*. Volume 179 (2010) 598-611.
- Sattler, S.E., Saathoff, A.J., Haas, E.J., Palmer, N.A., Funnell-Harris, D.L., Sarath, G., Pedersen, J.F. 2009. A nonsense mutation in a cinnamyl alcohol dehydrogenase gene is responsible for the sorghum brown midrib-6 phenotype. *Plant Physiology* 150 (2):584-95.
- Hooks, T., Marx, D.B., Kachman, S.D., Pedersen, J.F. 2009. *Revista Colombiana de Estadística*. Colombian Journal of Statistics 32:17-31.
- Funnell-Harris, D.L., Pedersen, J.F. 2008. Inoculation strategies to assess biological interactions between fusarium and alternaria species infecting sorghum. *Canadian Journal of Plant Pathology* 30: 404-413.
- Funnell-Harris, D.L., Pedersen, J.F., Marx, D.B. 2008. Effect of sorghum seedlings and previous crop, on soil fluorescent pseudomonas spp.. *Plant and Soil Journal* 311:173-187.
- Pedersen, J.F., Toy, J.J., Funnell-Harris, D.L., Sattler, S.E., Oliver, A.L. 2008. Registration of BN611, A/BN612, RN613 Sorghum Genetic Stocks with Stacked bmr-6 and bmr-12 Genes. *Journal of Plant Registrations* 2:258-262.
- Hooks, T., Marx, D.B., Kachman, S.D., Pedersen, J.F., Eigenberg, R.A. 2008. Analysis of covariance with spatially correlated secondary variables. *Revista Colombiana de Estadística* 31:95-109. 2008.
- Tesso, T., Kapran, I., Grenier, C., Snow, A., Sweeney, P., Pedersen, J.F., Marx, D., Bothma, G., Ejeta, G. 2008. The Potential for Crop-to-Wild Gene Flow in Sorghum in Ethiopia and Niger: A Geographical Survey. *Crop Science* 48: 1425-1431.
- Wong, J.H., Lau, T., Cai, N., Singh, J., Pedersen, J.F., Vensel, W.H., Hurkman II, W.J., Wilson, J.D., Lemaux, P.G., Buchanan, B.B. 2008. Digestibility of Protein and Starch from Sorghum (*Sorghum bicolor*) is Linked to Biochemical and Structural Features of Grain Endosperm. *Journal of Cereal Science*. 49(1):73-82.
- Sattler, Scott E., Jaswinder Singh, Eric J. Haas, Lining Guo, Gautam Sarath, and Jeffrey F. Pedersen Two distinct waxy alleles impact the granule-bound starch synthase in sorghum. *Molecular Breeding* 24:349-359
- Hooks, T., Pedersen, J.F., Marx, D.B., Gaussoin, R.E. 2007. Changing the support of a spatial covariate: a simulation study. *Crop Science* 47:622-626.

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GENOMIC RELATIONSHIPS AND CYTOGENETIC MANIPULATION OF DURUM WHEAT BY CLASSICAL AND MOLECULAR TECHNIQUES – Prem Jauhar (P); Fargo, North Dakota

- Jauhar, P.P., Peterson, T.S. 2009. Chromosome Engineering of Durum Wheat with Alien Chromatin of Diploid Wheatgrass. *Journal of Crop Improvement*. 23(4):319-331.
- Jauhar, P.P., Peterson, T.S., Xu, S.S. 2009. Cytogenetic and Molecular Characterization of Durum Alien Disomic Addition Line with Enhanced Tolerance to Fusarium Head Blight Resistance. *Genome* 52:467-483.
- Dogramaci, M., Jauhar, P.P. 2008. Synthesis of trigenic hybrids of hexaploid wheat with diploid wheatgrasses: Specificity of chromosome pairing. *The Nucleus*. 50:491-500.
- Jauhar, P.P., Dogramaci, M. 2007. Chromosome pairing in durum wheat haploids with and without ph1b of bread wheat. *Euphytica*. Published on-line 9-15-07 published 12-23-07 paper form 12-23-07 Volume 159 #3: 353-358
- Jauhar, P.P., Peterson, T.S. 2007. Dge-1, a durum alien disomic addition line with resistance to fusarium head blight. *Journal of Plant Registration* 2:167-168.

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IMPROVEMENT OF HARD RED SPRING AND DURUM WHEAT FOR DISEASE RESISTANCE AND QUALITY USING GENETICS AND GENOMICS – Justin Faris (P), Steven Xu, Shunwen Lu, and Shiaoman Chao; Fargo, North Dakota

- Niu, Z., Klindworth, D.L., Wang, R., Jauhar, P.P., Larkin, P.J., Xu, S.S. 2011. Characterization of high molecular weight glutenin subunits in *Thinopyrum intermedium*, *Th. bessarabicum*, *Lophopyrum elongatum*, *Aegilops markgrafii*, and their addition lines in wheat. *Crop Science*. 51:667-677.

- Zhang, Z., Friesen, T.L., Xu, S.S., Shi, G., Liu, Z., Rasmussen, J., Faris, J.D. 2011. Two Putatively Homoeologous Wheat Genes Mediate Recognition of SnTox3 to Confer Effector-triggered Susceptibility to *Stagonospora nodorum*. *Plant Journal*. 65:27-38.
- Zitzewitz, J., Condon, F., Corey, A., Cuesta-Marcos, A., Filichkin, T., Haggard, K., Fisk, S.P., Smith, K.P., Muehlbauer, G.J., Karsai, I., Veisz, O., Chao, S., Castro, A.J., Gutierrez, L., Hayes, P.M. 2011. The genetics of winterhardiness in barley: Perspectives from genome-wide association mapping. *The Plant Genome*. 4:76-91.
- Cistué, L., Chao, S., Chutimanitsakun, Y., Corey, A., Cuesta-Marcos, A., Echávarri, B., Filichkina, T., Garcia-Mariño, N., Romagosa, I., Hayes, P.M. 2011. Comparative Mapping of the Oregon Wolfe Barley Using Doubled Haploid Lines Derived from Female and Male Gametes. *Theoretical and Applied Genetics*. 122:1399-1410.
- Niu, Z., Klindworth, D.L., Friesen, T.L., Chao, S., Jin, Y., Cai, X., Xu, S.S. 2011. Targeted introgression of a wheat stem rust resistance gene by DNA marker-assisted chromosome engineering genetics. *Genetics*. 187(4):1011-1021.
- Simons, K.J., Abate, Z., Chao, S., Zhang, W., Rouse, M., Jin, Y., Elias, E., Dubcovsky, J. 2011. Genetic Mapping of Stem Rust Resistance Gene Sr13 in Tetraploid Wheat (*Triticum turgidum* subsp. *durum* L.). *Theoretical and Applied Genetics*. 122:649-658.
- Cai, X., Xu, S.S., Zhu, X. 2010. Ploidy-Dependent Unreductional Meiotic Cell Division in Polyploid Wheat. *Chromosoma*. 119:275-285.
- Chao, S., Dubcovsky, J., Dvorak, J., Luo, M., Baenziger, S.P., Matnyazov, R., Clark, D.R., Talbert, L., Anderson, J.A., Dreisigacker, S., Glover, K., Chen, J., Campbell, K., Bruckner, P.L., Rudd, J.C., Haley, S., Carver, B.F., Sims, R., Sorrells, M.E., Akhunov, E.D. 2010. Population-Specific Patterns of Linkage Disequilibrium and SNP Variation in Spring and Winter Polyploid Wheat. *Biomed Central (BMC) Genomics*. 11:727.
- Chao, S., Xu, S.S., Elias, E., Faris, J.D., Sorrells, M. 2010. Identification of Chromosome Locations of Genes Affecting pre-Harvest Sprouting and Seed Dormancy using Chromosome Substitution Lines in Tetraploid Wheat (*Triticum turgidum* L.). *Crop Science*. 50:1180-1187.
- Faris, J.D., Zhang, Z., Lu, H., Lu, S., Cloutier, S., Fellers, J.P., Meinhardt, S.W., Rasmussen, J.B., Xu, S.S., Oliver, R.P., Simons, K.J., Friesen, T.L. 2010. A Unique Wheat Disease Resistance-like Gene Governs Effector-Induced Susceptibility to Necrotrophic pathogens. *Proceedings of the National Academy of Sciences*. 107:13544-13549.
- Gu, X., Zhang, L., Glover, K., Chu, C., Xu, S.S., Faris, J.D., Friesen, T.L., Ibrahim, A. 2010. Genetic Variation of Seed Dormancy in Synthetic Hexaploid Wheat-Derived Populations. *Crop Science*. 50:1318-1324.
- Lu, S., Platz, G., Edwards, M.C., Friesen, T.L. 2010. Mating Type (MAT) Locus -Specific PCR Markers for Differentiation of *Pyrenophora teres* f. *teres* and *P. teres* f. *maculata*, the Causal Agents of Barley Net Blotch. *Phytopathology*. 100(12):1298-1306.
- Martin, T., Lu, S., Tilbeurgh, H., Ripoll, D., Dixelius, C., Turgeon, B.G., Debuchy, R. 2010. Tracing the Origin of the fungal Sex a1 domain places its ancestor in the HMG-box superfamily: implication for fungal mating-type evolution. *Proceedings of the National Academy of Sciences*. 5(12):e15199.
- Roy, J.K., Smith, K.P., Muehlbauer, G.J., Chao, S., Close, T.J., Steffenson, B.J. 2010. Association Mapping of Spot Blotch Resistance in Wild Barley. *Molecular Breeding*. 26:243-256.
- Xu, S.S., Chu, C.G., Chao, S., Klindworth, D.L., Faris, J.D., Elias, E.M. 2010. Marker-assisted Characterization of Durum Wheat Langdon-Golden Ball Disomic Substitution Lines. *Theoretical and Applied Genetics*. 120:1575-1585.
- Xu, S.S., Chu, C.G., Harris, M., Williams, C.E. 2010. Comparative analysis of genetic background in eight near-isogenic wheat lines with different H genes conferring resistance to Hessian fly. *Genome*. 54:81-89.
- Xu, S.S., Khan, K., Klindworth, D.L., Nygard, G. 2010. Evaluation and characterization of high-molecular weight 1D glutenin subunits from *Aegilops tauschii* in synthetic hexaploid wheats. *Journal of Cereal Science* 52:333-336.
- Yu, G., Zhang, Q., Klindworth, D.L., Friesen, T.L., Knox, R., Jin, Y., Zhong, S., Cai, X., Xu, S.S. 2010. Molecular and Cytogenetic Characterization of *Aegilops Speltoide*s Chromosome Segments Carrying Sr39 in the Original Translocation Stock and Derived Wheat Lines. *Crop Science*. 50:1393-1400.
- Abeysekara, N., Friesen, T.L., Keller, B., Faris, J.D. 2009. Identification and Characterization of a Novel Host-Toxin Interaction in the Wheat - *Stagonospora Nodorum* Pathosystem. *Theoretical and Applied Genetics*. 120:117-126.
- Chu, C., Chao, S., Friesen, T.L., Faris, J.D., Zhong, S., Xu, S.S. 2009. Novel Tan Spot Resistance QTLs Detected Using an SSR-Based Linkage Map of Tetraploid Wheat. *Molecular Breeding*. 25:327-338.
- Close, T.J., Bhat, P., Lonardi, S., Wu, Y., Rostoks, N., Ramsay, L., Druka, A., Stein, N., Svensson, J., Wanamaker, S., Bozdog, S., Roose, M., Moscou, M., Chao, S., Varshney, R., Szucs, P., Sato, K., Hayes, P., Matthews, D.E., Kleinhofs, A., Muehlbauer, G., Deyoung, J., Marshall, D.F., Madishetty, K., Fenton, R.D., Condamine, P., Graner, A., Waugh, R. 2009. Development and Implementation of High-Throughput SNP Genotyping in Barley. *Biomed Central (BMC) Genomics*. 10:582.

- Faris, J.D., Friesen, T.L. 2009. Re-Evaluation of a Tetraploid Wheat Population Indicates That the Tsn1-ToxA Interaction is the Only Factor Governing Stagonospora Nodorum Blotch Susceptibility. *Phytopathology*. 99:(8) 906-912.
- Feng, J., Primomo, V., Zhang, Y., Jan, C., Tulsieram, L., Xu, S.S. 2009. Physical Localization and Genetic Mapping of Fertility Restoration Gene Rfo in Canola (*Brassica napus* L.). *Genome*. 52:401-407.
- Klindworth, D.L., Hareland, G.A., Elias, E.M., Faris, J.D., Chao, S., Xu, S.S. 2009. Agronomic and Quality Characteristics of Two New Sets of Langdon Durum – wWld Emmer Wheat Chromosome Substitution Lines. *Journal of Cereal Science* 50: 29-35.
- Szucs, P., Blake, V.C., Chao, S., Marcos-Cuesta, A., Muehlbauer, G.J., Ramsay, L., Waugh, R., Hayes, P.M. 2009. An integrated resource for barley linkage map and malting quality QTL alignment. *The Plant Genome*. 2(2):134-140, 2009.
- Tsilo, T., Chao, S., Jin, Y., Anderson, J. 2009. Identification and validation of SSR markers linked to the stem rust resistance gene Sr6 on the short arm of chromosome 2D in wheat. *Theoretical and Applied Genetics*. 118:515-524.
- Xu, S.S., Jin, Y., Klindworth, D.L., Wang, R., Cai, X. 2009. Evaluation and Characterization of Seedling Resistances to Stem Rust Ug99 Races in Wheat-Alien Species Derivatives. *Crop Science*. 49:2167–2175.
- Zhang, Z., Friesen, T.L., Simons, K.J., Xu, S.S., Faris, J.D. 2009. Development, Identification and Validation of Markers for Marker-Assisted Selection Against the Stagonospora nodorum Toxin Sensitivity Genes Tsn1 and Snn2 in Wheat. *Molecular Breeding*, 23:35-49.
- Zhang, W., Chao, S., Manthey, F., Chicaiza, O., Brevis, J.C., Echenique, J., Dubcovsky, J. 2008. Qtl analysis of pasta quality using a composite microsatellite - snp map of durum wheat. *Theoretical and Applied Genetics*. 117:1361-1377.
- Faris, J.D., Xu, S.S., Xiwen, C., Friesen, T.L., Jin, Y. 2008. Molecular and cytogenetic characterization of a durum wheat *Aegilops speltoides* chromosome translocation conferring resistance to stem rust. *Chromosome Research*. 16(8): 1097-1105.
- Liu, S., Chao, S., Anderson, J.A. 2008. New DNA Markers for High Molecular Weight Glutenin Subunits in Wheat. *Theoretical and Applied Genetics*. 118:177-183.
- Anderson, J.A., Chao, S., Liu, S. 2008. Molecular Breeding Using a Major QTL for Fusarium Head Blight Resistance in Wheat. *Crop Science*. 47(3):S112-S119.
- Boda Slotta, T.A., Brady, L., Chao, S. 2008. High throughput tissue preparation for large-scale genotyping experiments. *Molecular Ecology Resources*. 8:83-87.
- Chalupska, D., Lee, H.Y., Faris, J.D., Evrard, A., Chalhoub, B., Haselkorn, R., Gornicki, P. 2008. Acc homoeoloci and the evolution of wheat genomes. *Proceedings of the National Academy of Sciences*. 105:9691-9696.
- Chao, S., Zhang, W., Akhunov, E., Sherman, J., Ma, Y., Luo, M., Dubcovsky, J. 2008. Analysis of gene-derived SNP marker polymorphism in wheat (*Triticum aestivum* L.). *Molecular Breeding*. 23:23-33.
- Chu, C., Friesen, T.L., Xu, S.S., Faris, J.D. 2008. Identification of novel tan spot resistance loci beyond the known host-selective toxin insensitivity genes in wheat. *Theoretical and Applied Genetics* 117:873-881.
- Chu, C., Xu, S.S., Friesen, T.L., Faris, J.D. 2008. Whole Genome Mapping in a Wheat Doubled Haploid Population Using SSRs and TRAPS and the Identification of QTL for Agronomic Traits. *Molecular Breeding*. 22:251-266.
- Chu, C.G., Friesen, T.L., Faris, J.D., Xu, S.S. 2008. Evaluation of seedling resistance to tan spot and Stagonospora nodorum blotch in tetraploid wheat. *Crop Science*. 48:1107-1116.
- Faris, J.D., Zhang, Z., Fellers, J.P., Gill, B.S. 2008. Micro-colinearity between rice, Brachypodium, and Triticum monococcum at the wheat domestication locus Q. *Functional and Integrative Genomics*. 8:149-164.
- Oliver, R.E., Cai, X., Friesen, T.L., Halley, S., Stack, R.W., Xu, S.S. 2008. Evaluation of Fusarium head blight resistance in tetraploid wheat (*Triticum turgidum* L.). *Crop Science*. 48:213-222.
- Oliver, R.P., Lord, M., Rybak, K., Faris, J.D., Solomon, P.S. 2008. The emergence of tan spot disease caused by toxigenic *Pyrenophora tritici-repentis* in Australia is not associated with increased deployment of toxin-sensitive cultivars. *Phytopathology*. 98:488-491.
- Reedy, L., Friesen, T.L., Meinhardt, S.W., Chao, S., Faris, J.D. 2008. Genomic analysis of the Snn1 locus on wheat chromosome arm 1BS and the identification of candidate genes. *The Plant Genome*. 1:55-66.
- Tai, Y. 2007. Detection of reactive oxygen species can be used to distinguish ToxA-induced cell death from the hypersensitive response. *Research Journal of Botany* 2(1):1-12.
- Tai, Y. 2007. The Role of Wheat Q Gene and its Signaling Pathways in Response to Abiotic Stress. *American Journal of Plant Physiology*. 2:295-302.
- Tai, Y., Bragg, J. 2007. Dual applications of a virus vector for studies of wheat-fungal interactions. *Biotechnology* 6 (2):288-291.
- Tai, Y., Bragg, J., Meinhardt, S. 2007. Functional Characterization of ToxA and Molecular Identification of its Intracellular Targeting Protein in Wheat. *American Journal of Plant Physiology* 2:76-89.
- Klindworth, D.L., Miller, J., Jin, Y., Xu, S.S. 2007. Chromosomal locations of genes for stem rust resistance in monogenic lines derived from tetraploid wheat accession ST464. *Crop Science*. 47:1441-1450.

- Kumar, S., Gill, B.S., Faris, J.D. 2007. Identification and Characterization of Segregation Distortion Loci Along Chromosome 5B in Tetraploid Wheat. *Molecular Genetics and Genomics*. 278:187-196.
- Kumar, S., Stack, R.W., Friesen, T.L., Faris, J.D. 2007. Identification of a novel Fusarium head blight resistance QTL on chromosome 7A in tetraploid wheat. *Phytopathology*. 97:592-597.
- Li, J., Klindworth, D.L., Shireen, F., Cai, X., Hu, J., Xu, S.S. 2007. Molecular characterization and chromosome-specific TRAP-marker development for Langdon durum D-genome disomic substitution lines. *Genome*. 49:1545-1554.
- Chao, S., Zhang, W., Dubcovsky, J., Sorrells, M. 2007. Evaluation of genetic diversity and genome-wide linkage disequilibrium among U.S. wheat (*Triticum aestivum* L.) germplasm representing different market classes. *Crop Science*. 47:1018-1030.
- Chen, X., Faris, J.D., Hu, J., Stack, R.W., Adhikari, T., Elias, E.M., Kianian, S.F., Cai, X. 2006. Saturation and comparative mapping of a major Fusarium head blight resistance QTL in tetraploid wheat. *Molecular Breeding*. 19:113-124.

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SUNFLOWER GERMLASM DIVERSIFICATION AND CHARACTERIZATION UTILIZING WILD SUNFLOWER SPECIES, CYTOGENETICS, AND APPLIED GENOMICS – Brady Vick (P), Lili Qi, Chao-Chien Jan, and Gerald Seiler; Fargo, North Dakota

- Qi, L.-L., Gulya, T.J., Seiler, G.J., Hulke, B.S., Vick, B.A. 2011. Identification of resistance to new virulent races of rust in sunflowers and validation of DNA markers in the gene pool. *Phytopathology*. 101:241-249.
- Cantamutto, M., Presotto, A., Moroni, I.F., Alvarez, D., Poverene, M., Seiler, G. 2010. High Intraspecific Diversity of Wild Sunflowers (*Helianthus annuus* L.) Naturally Developed in Central Argentina. *Flora*. 205:306-312.
- Qi, L., Friebe, B., Wu, J., Gu, Y., Qian, C., Gill, B.S. 2010. The compact Brachypodium genome conserves centromeric regions of a common ancestor with wheat and rice. *Functional and Integrative Genomics*. 10:477-492.
- Seiler, G.J. 2010. Germination and viability of wild sunflower species achenes stored at room temperature for 20 years. *Seed Science and Technology*. 38:786-791.
- Seiler, G.J., Gulya, T.J., Kong, G. 2010. Oil Concentration and Fatty Acid Profile of Wild *Helianthus* Species from the Southeastern United States. *Industrial Crops and Products*. 31(3):527-533.
- Yue, B., Vick, B.A., Cai, X., Hu, J. 2010. Genetic Mapping for the Rf1 (Fertility Restoration) Gene in Sunflower (*Helianthus annuus* L.) by SSR and TRAP Markers. *Plant Breeding*. 129:24-28.
- Zhao, W., Qi, L., Gao, X., Zhang, G., Dong, J., Chen, Q., Friebe, B., Gill, B.S. 2010. Development and characterization of two new *Triticum aestivum*-*Dasypyrum villosum* Robertsonian translocation lines T1DS.1V#3L and T1DL.1V#3S and their effect on grain quality. *Euphytica*. 175:343-350.
- Zheljaskov, V.D., Vick, B.A., Baldwin, B.S., Buehring, N., Coker, C., Astatkie, T., Johnson, B. 2010. Oil productivity and composition of sunflower as a function of hybrid and planting date. *Industrial Crops and Products*. 33:537-543.
- Dechaine, J.M., Burger, J.C., Chapman, M.A., Seiler, G.J., Brunick, R., Knapp, S.J., Burke, J.M. 2009. Fitness Effects and Genetic Architecture of Plant-Herbivore Interactions in Sunflower Crop-Wild Hybrids. *New Phytologist*. 184:828-841.
- Mulpuri, S., Liu, Z., Feng, J., Gulya, T.J., Jan, C. 2009. Inheritance and Molecular Mapping of a Downy Mildew Resistance Gene, Pl13 in Cultivated Sunflower (*Helianthus annuus* L.). *Theoretical and Applied Genetics*. 119(5):795-803.
- Presotto, A., Cantamutto, M., Poverene, M., Seiler, G. 2009. Phenotypic Diversity in Wild *Helianthus annuus* from Argentina. *Helia*. 32(50):37-50.
- Qi, L., Friebe, B., Zhang, P., Gill, B.S. 2009. A Molecular-Cytogenetic Method for Locating Genes to Pericentromeric Regions Facilitates a Genomewide Comparison of Synteny Between the Centromeric Regions of Wheat and Rice. *Genetics*. 183:1235-1247.
- Xu, S., Huang, Q., Shu, Q., Chen, C., Vick, B.A. 2009. Reproductive Organography of *Bougainvillea spectabilis* Willd. *Scientia Horticulturae*. 120:399-405.
- Zheljaskov, V.D., Vick, B.A., Balwin, B.S., Buehring, N., Astatkie, T., Johnson, B. 2009. Oil Content and Saturated Fatty Acids in Sunflower as a Function of Planting Date, Nitrogen Rate, and Hybrid. *Agronomy Journal*. 101:1003-1011.
- Vick, B.A., Cook, L.W., Jan, C.C. 2009. Characterization of a Green Non-Dormant Sunflower (*Helianthus annuus*) Mutant NDG. *Plant Breeding*. 128:645-650.
- Yue, B., Cai, X., Vick, B.A., Hu, J. 2009. Genetic Characterization and Molecular Mapping of a Chlorophyll Deficiency Gene in Sunflower (*Helianthus annuus*). *Journal of Plant Physiology*. 166:644-651.
- Yue, B., Cai, X., Yuan, W., Vick, B.A., Hu, J. 2009. Mapping Quantitative Trait Loci (QTL) Controlling Seed Morphology and Disk Diameter in Sunflower (*Helianthus annuus* L.). *Helia*. 32(50):17-36.

- Zhang, J., Wang, L., Zhao, A., Liu, H., Jan, C.C., Qi, D., Liu, G. 2009. Morphological and Cytological Study in a New Type of Cytoplasmic Male Sterile Line CMS-GIG2 in Sunflower (*Helianthus annuus*). *Plant Breeding*. 129:19-23.
- Poverene, M., Cantamutto, M., Seiler, G.J. 2008. Ecological Characterization of Wild *Helianthus annuus* and *Helianthus petiolaris* Germplasm in Argentina. *Plant Genetic Resources: Characterization and Utilization*. 7(1):42-49.
- Feng, J., Jan, C.C. 2008. Introgression and molecular tagging of Rf4, a new male fertility restoration gene from wild sunflower *Helianthus maximiliani* L. *Theoretical and Applied Genetics*. 117:241-249.
- Hu, J., Yue, B., Yuan, W., Vick, B.A. 2008. Growing sunflower plants from seed to seed in small pots in greenhouse. *Helia*. 31(48):119-126.
- Rojas-Barros, P., Hu, J., Jan, C.C. 2008. Molecular mapping of an apical branching gene of cultivated sunflower (*Helianthus annuus* L.). *Theoretical and Applied Genetics*. 117:19-28.
- Seiler, G.J. 2008. Root Growth of Interspecific Sunflower Seedlings Derived from Wild Perennial Sunflower Species. *Canadian Journal of Plant Science*. 88:705-712.
- Yue, B., Radi, S.A., Vick, B.A., Cai, X., Tang, S., Knapp, S.J., Gulya, T.J., Miller, J.F., Hu, J. 2008. Identifying quantitative trait loci for resistance to *Sclerotinia* head rot in two USDA sunflower germplasms. *Phytopathology*. 98(8):926-931.
- Yue, B., Vick, B.A., Yuan, W., Hu, J. 2008. Mapping one of the 2 genes controlling lemon ray flower color in sunflower (*Helianthus annuus* L.). *Journal of Heredity*. 99(5):564-567.
- Zheljaskov, V.D., Vick, B.A., Ebelhar, M.W., Buehring, N., Baldwin, B.S., Astatkie, T., Miller, J.F. 2008. Yield, oil content, and composition of sunflower grown at multiple locations in Mississippi. *Agronomy Journal*. 100(3):635-642.
- Deng, Z., Goktepe, F., Harbaugh, B.K., Hu, J. 2007. Assessment of genetic diversity and relationships among caladium cultivars and species using molecular markers. *Journal of the American Society for Horticultural Science*. 132(2):219-229.
- Hu, J., Mou, B., Vick, B.A. 2007. Genetic diversity of 38 spinach (*Spinacia oleracea* L.) germplasm accessions and 10 commercial hybrids assessed by TRAP markers. *Genetic Resources and Crop Evolution*. 54:1667-1674.
- Hu, J., Yue, B., Vick, B.A. 2007. Integration of TRAP markers onto a sunflower SSR marker linkage map constructed from 92 recombinant inbred lines. *Helia*. 30(46):25-36.
- Seiler, G.J. 2007. The potential of wild sunflower species for industrial uses. *Helia*. 30(46):175-198.
- Seiler, G.J. 2007. Wild annual *Helianthus anomalus* and *Helianthus deserticola* as potential sources of improved oil concentration and quality in sunflower. *Industrial Crops and Products*. 25:95-100. Available: <http://dx.doi.org/10.1016/j.indcrop.2006.07.007>.
- Vick, B.A., Jan, C.C., Miller, J.F. 2007. Registration of a sunflower genetic stock RS3 with reduced levels of palmitic and stearic acids. *Journal of Plant Registrations*. 1:80.
- Maiti, R.K., Vidyasagar, P., Shahapur, S.C., Seiler, G.J. 2006. Studies on Genotypic Variability and Seed Dormancy in Sunflower Genotypes (*Helianthus annuus* L.). *Indian Journal of Crop Science*. 1(1-2):84-87.
- Seiler, G.J., Gulya, T.J., Marek, L.F. 2006. Exploration for wild *Helianthus* species from the desert southwestern USA for potential drought tolerance. *Helia*. 29(45):1-10.

5442-21220-035-00D

GENETIC IMPROVEMENT OF BARLEY – Lynn Dahleen (P); Fargo, North Dakota

- Tyagi, N., Dahleen, L.S. 2011. Timing of Ethylene Modification Is Critical For Regeneration In Barley. *Plant and Molecular Biology and Biotechnology*. 4:47-55.
- Tyagi, N., Dahleen, L.S., Bregitzer, P.P. 2010. Candidate Genes Within Tissue Culture Regeneration QTL Revisited with a Linkage Map Based on Transcript Derived Markers. *Crop Science*. 50:1697-1707.
- Mittal, S., Dahleen, L.S., Mornhinweg, D.W. 2009. Barley Germplasm STARS-9577B lacks a Russian Wheat Aphid Resistance Quantitative Trait Locus Present in STARS-9301B. *Crop Science*. 49:1999-2004

5442-21220-024-00D

SUNFLOWER GERMPLASM DEVELOPMENT FOR IMPROVED INSECT AND DISEASE RESISTANCE – Thomas Gulya Jr (P) and Brent Hulke; Fargo, North Dakota

- Mathew, F., Kirkeide, B., Gulya, T.J., Markell, S. 2010. First report of pathogenicity of *Fusarium sporotrichioides* and *Fusarium acuminatum* on sunflowers in the United States. *Plant Health Progress*. Available: <http://www.plantmanagementnetwork.org/sub/php/brief/2010/fusarium/>.

- Charlet, L.D., Seiler, G.J., Grady, K.A., Hulke, B.S., Chirumamilla, A. 2010. Resistance in Cultivated Sunflower Germplasm to the Red Sunflower Seed Weevil (Coleoptera: Curculionidae) in the Northern Great Plains. *Journal of the Kansas Entomological Society*. 83:51-57.
- Gulya, T.J., Mengistu, A., Kinzer, K., Balbyshev, N., Markell, S. 2010. First Report of Charcoal Rot of Sunflower in Minnesota, USA. *Plant Health Progress*. Available: <http://www.plantmanagementnetwork.org/sub/php/brief/2010/charcoal/>.
- Hulke, B.S., Miller, J.F., Gulya, T.J. 2010. Registration of the Restorer Oilseed Sunflower Germplasm RHA 464 Possessing Genes for Resistance to Downy Mildew and Sunflower Rust. *Journal of Plant Registrations*. 4:249-254.
- Hulke, B.S., Miller, J.F., Gulya, T.J., Vick, B.A. 2010. Registration of the Oilseed Sunflower Genetic Stocks HA 458, HA 459, and HA 460 Possessing Genes for Resistance to Downy Mildew. *Journal of Plant Registrations*. 4:93-97.
- Johnsen, A.R., Horgan, B.P., Hulke, B.S., Cline, V. 2009. Evaluation of Remote Sensing to Measure Plant Stress in Creeping Bentgrass (*Agrostis stolonifera* L.) Fairways. *Crop Science*. 49:2261-2274.
- Charlet, L.D., Seiler, G.J., Miller, J.F., Hulke, B.S., Knodel, J.J. 2009. Resistance Among Cultivated Sunflower Germplasm to the Banded Sunflower Moth (Lepidoptera: Tortricidae) in the Northern Great Plains. *Helia*. 32:1-10.
- Morris, B.D., Charlet, L.D., Foster, S.P. 2009. Isolation of Three Diterpenoid Acids from Sunflowers, as Oviposition Stimulants for the Banded Sunflower Moth, *Cochylis hospes*. *Journal of Chemical Ecology*. 35:50-57.
- Charlet, L.D., Aiken, R.M., Miller, J.F., Seiler, G.J. 2009. Resistance Among Cultivated Sunflower Germplasm to Stem-Infesting Pests in the Central Great Plains. *Journal of Economic Entomology*. 102(3):1281-1290.
- Charlet, L.D., Aiken, R.M., Seiler, G.J., Chirumamilla, A., Hulke, B.S., Knodel, J.J. 2008. Resistance in Cultivated Sunflower to the Sunflower Moth (Lepidoptera: Pyralidae). *Journal of Agricultural and Urban Entomology*. 25(4):245-257.
- Charlet, L.D., Aiken, R.M., Meyer, R.F., Gebre-Amlak, A. 2007. Impact of combining planting date and chemical control to reduce larval densities of stem-infesting pests of sunflower in the Central Plains. *Journal of Economic Entomology*. 100(4):1248-1257.
- Charlet, L.D., Aiken, R.M., Meyer, R.F., Gebre-Amlak, A. 2007. Impact of irrigation on larval density of stem-infesting pests of cultivated sunflower in Kansas. *Journal of Economic Entomology*. 100(5):1555-1559.
- Gulya, T.J., Seiler, G.J., Kong, G., Marek, L.F. 2007. Exploration and collection of rare *Helianthus* species from southeastern United States. *Helia*. 30(46):13-24.
- Miller, J.F., Gulya Jr, T.J., Vick, B.A. 2006. Registration of three maintainer (HA 456, HA 457, and HA 412 HO) high-oleic oilseed sunflower germplasms. *Crop Science*. 46:2728.
- Miller, J.F., Gulya Jr, T.J., Vick, B.A. 2006. Registration of two maintainer (HA 451 and HA 452) and three restorer (RHA 453-RHA 455) Sclerotinia-tolerant oilseed sunflower germplasms. *Crop Science*. 46:2727-2728.

5442-21430-007-00D

SUCROSE ACCUMULATION AND RETENTION IN SUGARBEETS – Karen Fugate (P) and Larry Campbell; Fargo, North Dakota

- Lafta, A.M., Fugate, K.K. 2011. Metabolic profile of wound-induced changes in primary carbon metabolism in sugarbeet root. *Phytochemistry*. 72:476-489.
- Campbell, L.G., Panella, L.W., Smigocki, A.C. 2011. Registration of F1024 sugarbeet germplasm with resistance to sugarbeet root maggot. *Journal of Plant Registrations*. 5(2):241-247.
- Fugate, K.K., Suttle, J.C., Campbell, L.G. 2010. Ethylene Production and Ethylene Effects on Respiration Rate of Postharvest Sugarbeet Roots. *Postharvest Biology and Technology*. 56:71-76

6202-21000-027-00D

GENETIC ANALYSES AND TRAIT MAPPING TO EXPLOIT UNTAPPED GENETIC DIVERSITY IN SORGHUM – Robert Klein (P); College Station, Texas

- Jordan, D.R., Mace, E.S., Henzell, R.G., Klein, P.E., Klein, R.R. 2010. Molecular mapping and candidate gene identification of the Rf2 gene for pollen fertility restoration in sorghum (*Sorghum bicolor* (L.) Moench). *Theoretical and Applied Genetics*. 120(7):1279-1287.
- Mace, E.S., Rami, J., Bouchet, S., Klein, P.E., Klein, R.R., Kilian, A., Wenzl, P., Zia, L., Halloran, K., Jordan, D. 2009. A consensus genetic map of sorghum that integrates multiple component maps and high-throughput diversity array technology (DArT) markers. *BMC Plant Biology*. 9:13.

- Klein, R.R., Mullet, J.E., Jordan, D.R., Miller, F.R., Rooney, W.L., Menz, M.A., Franks, C.D., Klein, P.E. 2008. The effect of tropical sorghum conversion and inbred development on genome diversity as revealed by high-resolution genotyping. *Crop Science*. 48(S1):S12-S26.
- Harris, K., Borrell, A., Jordan, D., Rosenow, D., Nguyen, H., Klein, P., Klein, R.R., Mullet, J. 2007. Sorghum stay-green QTL individually reduce post-flowering drought-induced leaf senescence. *Journal of Experimental Botany*. 58:327-338.

6202-21000-029-00D

PECAN GENETICS AND IMPROVEMENT – Tommy Thompson (P) and Larry Grauke; College Station, Texas

- Thompson, T.E., Grauke, L.J. 2010. 'Apalachee' Pecan. *HortScience*. 45(9):1386-1387.
- Thompson, T.E., Grauke, L.J. 2010. 'Mandan' Pecan. *HortScience*. 45(9):1392-1393.
- Venkatachalam, M., Kshirsagar, H.H., Seeram, N.P., Heber, D., Thompson, T.E., Roux, K.H., Sathe, S.K. 2007. Biochemical composition and immunological comparison of select pecan [*Carya illinoensis* (Wangenh.) K. Koch] cultivars. *Journal of Agricultural and Food Chemistry*. 55:9899-9907.

6202-21000-030-00D

COTTON GENOMICS AND GENETIC ANALYSIS – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

- Xu, Z., Yu, J., Cho, J., Yu, J., Kohel, R.J., Percy, R.G. 2010. Polyploidization altered gene functions in cotton (*Gossypium* spp.). *PLoS One*. 5(12):Article e14351. Available: <http://www.plosone.org/article/info%3Adoi%2F10.1371%2Fjournal.pone.0014351>.
- Yu, J., Kohel, R.J., Smith, C. 2010. The construction of a tetraploid cotton genome wide comprehensive reference map. *Genomics*. 95:230-240.
- Van Deynze, A., Stoffel, K., Lee, M., Wilkins, T.A., Kozik, A., Cantrell, R.G., Yu, J., Kohel, R.J., Stelly, D.M. 2009. Sampling nucleotide diversity in cotton. *BMC Plant Biology*. 9:Article 125.
- Abdurakhmonov, I.Y., Kohel, R.J., Yu, J., Pepper, A.E., Abdullaev, A.A., Kushanov, F.N., Salakhutdinov, I.B., Buriev, Z.T., Saha, S., Scheffler, B.E., Jenkins, J.N., Abdurkarimov, A. 2008. Molecular diversity and association mapping of fiber quality traits in exotic *G. hirsutum* L. germplasm. *Genomics*. 92:478-487.
- Xu, Z., Kohel, R.J., Song, G., Cho, J., Alabady, M., Yu, J., Poo, K., Chu, J., Yu, S., Wilkins, T.A., Zhu, Y., Yu, J. 2008. Gene-rich islands for fiber development in the cotton genome. *Genomics*. 92:173-183.
- Xu, Z., Kohel, R.J., Song, G., Cho, J., Yu, J., Yu, S., Tomkins, J., Yu, J. 2008. An integrated genetic and physical map of homoeologous chromosomes 12 and 26 in Upland cotton (*G. hirsutum* L.). *BMC Genomics*. 9:108.
- Hoffman, S.M., Yu, J., Grum, D.S., Xiao, J., Kohel, R.J., Pepper, A.E. 2007. Identification of 700 new microsatellite loci from cotton (*G. hirsutum*). *Journal of Cotton Science*. 11:208-241.
- Wang, K., Song, X., Han, Z., Guo, W., Yu, J., Sun, J., Pan, J., Kohel, R.J., Zhang, T. 2006. Complete assignment of the *G. hirsutum* L. chromosomes by BAC-FISH and translocation analyses. *Journal of Theoretical and Applied Genetics*. 113:73-80.

6202-21000-031-00D

COTTON GENETIC RESOURCE MANAGEMENT – Richard Percy (P), James Frelichowski, Lori Hinze, and John Yu; College Station, Texas

- Hinze, L.L., Kohel, R.J., Campbell, B.T., Percy, R.G. 2011. Variability in four diverse cotton (*Gossypium hirsutum* L.) germplasm populations. *Genetic Resources and Crop Evolution*. 58(4):561-570.
- Hinze, L.L., Percy, R.G., Jones, D. 2010. Fiber quality of cultivars and breeding lines in the Cotton Winter Nursery and U.S. environments. *Journal of Cotton Science*. 14(3):138-144.
- Percy, R.G., Campbell, B.T., Chee, P.W., Jones, D.C., Lubbers, E.L., Myers, G.O., Ulloa, M. 2010. Registration of CRB 252, an upland cotton germplasm line possessing superior fiber quality traits. *Journal of Plant Registrations*. 4(3):1-4.
- Percy, R.G., Ulloa, M., Zhang, J. 2009. Registration of PSI 113 and PSI 425 germplasm lines of Pima cotton possessing superior fiber quality traits. *Journal of Plant Registrations*. 3(3):297-299.
- Lu, Y., Curtiss, J., Percy, R.G., Hughs, S.E., Yu, S., Zhang, J. 2009. DNA polymorphisms of genes involved in fiber development in a selected set of cultivated tetraploid cotton. *Crop Science*. 49:1695-1704.

6202-21000-032-00D

CHARACTERIZING AND CONSERVING THE GENETIC DIVERSITY OF CARYA POPULATIONS – Larry Grauke (P) and Tommy Thompson; College Station, Texas

- Sagaram, M., Lombardini, L., Grauke, L.J. 2011. Variation in anatomical characteristics in leaves of pecan seedstocks from Mexico and the United States. *Journal of the American Society for Horticultural Science*. 136(2):103-108.
- Grauke, L.J., Thompson, T.E. 2010. Pecan: Register of new fruit and nut cultivars - list 45. *HortScience*. 45(5):739-740.
- Grauke, L.J., Thompson, T.E. 2008. Pecan. *HortScience*. 43(5):1334.
- Sagaram, M., Lombardini, L., Grauke, L.J. 2007. Variation in leaf anatomy of pecan cultivars from three ecogeographic locations. *Journal of the American Society for Horticultural Science*. 132:1-5.

6203-21000-001-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Shannon Pinson, and Ming Hsuan Chen; Stuttgart, Arkansas

- Min, B., McClung, A.M., Chen, M.H. 2011. Phytochemicals and antioxidant capacity in rice brans of different color. *Journal of Food Science*. 76:C117-126.
- Chen, M., Fjellstrom, R.G., Christensen, E.F., Bergman, C.J. 2010. Development of three allele-specific codominant rice Waxy gene PCR markers suitable for marker assisted selection of amylose content and paste viscosity. *Molecular Breeding*. 26:513-523.
- Heuberger, A.L., Lewis, M.R., Chen, M., Brick, M.A., Leach, J.E., Ryan, E.P. 2010. Metabolomic and functional genomic analyses reveal varietal differences in bioactive compounds of cooked rice. *PLoS One* 5:e12915.
- Pinson, S.R., Shahjahan, A., Rush, M.C., Groth, D.E. 2010. Bacterial panicle blight resistance QTL in rice (*Oryza sativa* L.) and their association with resistance to other diseases. *Crop Science*. 50:1287-1297.
- Fitzgerald, M.A., Bergman, C.J., Ressurreccion, A.P., Moller, J., Jimenez, R., Reinke, R.F., Martin, M., Blanco, P., Molina, F., Chen, M., Kuri, V., Romero, M.V., Habibi, F., Umamoto, T., Jongdee, S., Graterol, E., Reddy, K.R., Bassinello, P.Z., Sivakami, R., Rani, N.S., Das, S., Wang, Y., Indrassi, S.D., Ramli, A., Rauf, A., Dipti, S.S., Xie, L., Lang, N.T., Singh, P., Toro, D.C., Tavasoli, F. 2009. Addressing the dilemmas of measuring amylose in rice. *Cereal Chemistry*. 86(5):492-498.
- Sharma, A., McClung, A.M., Pinson, S.R., Kepiro, J.L., Shank, A.R., Tabien, R.E., Fjellstrom, R.G. 2009. Genetic mapping of sheath blight resistance to QTLs within tropical Japonica rice cultivars. *Crop Science* 49:256-264.
- Chen, M., Bergman, C.J., Pinson, S.R., Fjellstrom, R.G. 2008. Waxy gene haplotypes: Associations with amylose content and the effect by the environment in an international rice germplasm collection. *Journal of Cereal Science*. 47:536-545.
- Chen, M., Bergman, C.J., Pinson, S.R., Fjellstrom, R.G. 2008. Waxy gene haplotypes: Associations with pasting properties in an international rice germplasm collection. *Journal of Cereal Science*. 48:781-788.
- Kepiro, J.L., McClung, A.M., Chen, M., Yeater, K.M., Fjellstrom, R.G. 2008. Mapping QTL for milling yield and grain characteristics in a tropical japonica long grain cross. *Journal of Cereal Science* 48:477-485.
- Samonte, S.B., Wilson, L.T., Tabien, R.E., McClung, A.M. 2008. Use of gross income as a measure of productivity in rice breeding. *Canadian Journal of Plant Science* 88:1015-1022.
- Chen, M.H., Bergman, C.J. 2007. A method to determine the content, molecular weights, and weight- and molar-based distributions of degree of polymerization of amylose and fine-structure of amylopectin. *Carbohydrate Polymers* 69:562-578.
- Samonte, S.O., Wilson, L.T., Pinson, S.R., McClung, A.M., Lales, J.S. 2006. Nitrogen utilization efficiency: Relationships with grain yield, grain protein, and yield-related traits in rice. *Agronomy Journal*. 98:168-176.
- Aboubacar, A., Moldenhauer, K.A., McClung, A.M., Beighley, D.H., Hamaker, Bruce R. 2006. Effect of growth location in the United States on amylose content, amylopectin fine structure, and thermal properties of starches of long grain rice cultivars. *Cereal Chemistry*. 83(1):93-98.

6208-21000-017-00D

MOLECULAR AND GENETIC ENHANCEMENT OF ABIOTIC STRESS TOLERANCE IN SORGHUM – Zhanguo Xin, Gloria Burow, and John Burke; Lubbock, Texas

- Mahan, J.R., Conaty, W., Neilsen, J., Gitz, D.C. 2010. Field performance in an agricultural setting of a wireless temperature monitoring system based on a low-cost infrared sensor. *Computers and Electronics in Agriculture*. 71(2):176-181.

- Burke, J.J., Franks, C.D., Burow, G.B., Xin, Z. 2010. Selection system for the Stay-green" drought tolerance trait in sorghum germplasm. *Agronomy Journal*. 102(4):1118-1122."
- Lu, C., Xin, Z., Ren, Z., Miquel, M., Browse, J. 2009. An enzyme regulating triacylglycerol composition is encoded by the ROD1 gene of Arabidopsis. *Proceedings of the National Academy of Sciences*. 106(44):18837-18842.
- Xin, Z., Aiken, R., Burke, J.J. 2009. Genetic diversity in sorghum transpiration efficiency. *Field Crops Research*. 11(1-2): 74-80.
- Xin, Z., Wang, M.L., Burow, G.B., Burke, J.J. 2009. An induced sorghum mutant population suitable for bioenergy research. *BioEnergy Research*. 2(1-2):10-16.
- Burow, G.B., Franks, C.D., Xin, Z., Acosta Martinez, V. 2009. Molecular mapping and characterization of BLMC, a locus for profuse wax (bloom) and enhanced cuticular features of sorghum (*Sorghum bicolor* (L.) Moench.). *Theoretical and Applied Genetics*. 118:423-431.
- Burow, G.B., Franks, C.D., Xin, Z. 2008. Genetic and physiological analysis of an irradiated bloomless mutant (epicuticular wax mutant) of sorghum. *Crop Science*. 48:41-48.
- Niu, C., Kebede, H., Auld, D., Woodward, J., Burow, G.B., Wright, R. 2008. A safe inexpensive method to isolate high quality plant and fungal DNA in an open laboratory environment. *African Journal of Biotechnology*. 7(16):2818-2822.
- Xin, Z., Wang, M.L., Barkley, N.L., Burow, G., Franks, C., Pederson, G., Burke, J. 2008. Applying genotyping (TILLING) and phenotyping analyses to elucidate gene function in a chemically induced sorghum mutant population. *Biomed Central (BMC) Plant Biology*. 8:article 103.
- Casa, A., Pressoir, G., Brown, P., Mitchell, S., Rooney, W., Tuinstra, M., Franks, C.D., Kresovich, S. 2008. Community resources and strategies for association mapping in sorghum. *Crop Science*. 48(1):30-40.
- Gomez, M., Burow, G.B., Burke, J.J., Burow, M., Denwar, N., Simpson, C., Ramasubramanian, T., Puppala, N. 2008. Identification of peanut hybrids using microsatellite markers and horizontal polyacrylamide gel electrophoresis. *Peanut Science*. 35(2):123-129.
- Mahan, J.R., Yeater, K.M. 2008. Development of a low cost infrared thermometry system for agricultural applications. *Computers and Electronics in Agriculture*. 64(2): 262-267.
- Kottapalli, K., Burow, M., Burow, G.B., Burke, J.J., Puppala, N. 2007. Molecular characterization of the US peanut mini core collection using microsatellite markers. *Crop Science*. 47(4):1718-1727.
- Burow, G.B., Franks, C.D., Burke, J.J. 2006. Evaluation of anthocyanin pigmentation as an indicator of chilling sensitivity of sorghum seedlings. *International Sorghum and Millets Newsletter*. 47:39-42.

6208-21000-018-00D

CHARACTERIZATION AND ENHANCEMENT OF PLANT RESISTANCE TO WATER-DEFICIT AND THERMAL STRESSES – John Burke (P), James Mahan, Jeffrey Velten, Zhanguo, Paxton Payton, and Junping Chen; Lubbock, Texas

- Chen, J., Xu, W., Burke, J.J., Xin, Z. 2010. Role of phosphatidic acid in high temperature tolerance in maize [abstract]. *Crop Science*. 50:2506-2515.
- Pasapula, V., Shen, G., Kuppu, S., Paez-Valencia, J., Mendoza, M., Hou, P., Chen, J., Qui, X., Zhu, L., Zhang, X., Auld, D., Blumwald, E., Zhang, H., Gaxiola, R., Payton, P.R. 2010. Expression of an arabidopsis vacuolar H⁺-pyrophosphatase gene (AVP1) in cotton improves drought- and salt tolerance and increases fibre yield in the field conditions. *Plant Biotechnology Journal*. 8(5):1-12.
- Burke, J.J., Franks, C., Burow, G.B., Xin, Z. 2010. Selection System for the Stay-Green Drought Tolerance Trait in Sorghum Germplasm. *Agronomy Journal*. 102(4):1118-1122.
- Cakir, C., Scofield, S.R., Gillespie, M.E. 2010. Rapid determination of gene function by virus-induced gene silencing in wheat and barley. *Crop Science*. 50(2, Suppl. S):S77-S84.
- Velten, J.P., Cakir, C., Cazzonelli, C. 2010. Gene dosage induction of silencing directed against an Arabidopsis Myb transgene in tobacco[abstract]. *Keystone Symposia: RNA Silencing Mechanisms in Plants*, February 21-26, 2010, Santa Fe, New Mexico.

6217-21000-007-00D

GENETIC AND GENOMIC APPROACHES TO IMPROVE INSECT RESISTANCE AND OTHER VALUE-ADDED TRAITS IN WHEAT, BARLEY, AND SORGHUM – Yinghua Huang (P), Gary Puterka, and Dolores Mornhinweg; Stillwater, Oklahoma

- Dunn, B.L., Porter, D.R., Baker, C.A., Carver, B.F. 2011. Screening USDA-ARS wheat germplasm for bird cherry-oat aphid tolerance. *Journal of Crop Improvement*. 25(2):176-182.
- Chou, J., Huang, Y. 2010. Differential expression of thaumatin-like proteins in sorghum infested with greenbugs. *Zeitschrift Fur Naturforschung*. 65c(3-4):271-276.

- Huss, J., Baker, C.A. 2010. The farmer in the lab. *Science and Children*. 47(6):40-43.
- Mornhinweg, D.W., Bregitzer, P.P., Porter, D.R., Peairs, F.B., Baltensperger, D.D., Hein, G.L., Randolph, T.A., Koch, M., Walker, T. 2009. Registration of 'Sidney' spring feed barley resistant to Russian wheat aphid. *Journal of Plant Registrations*. 3(3):214-218.
- Wu, Y.Q., Huang, Y. 2008. Molecular mapping of QTLs for resistance to the greenbug *Schizaphis graminum* (Rondani) in *Sorghum bicolor* (Moench). *Theoretical and Applied Genetics*. 117(1):117-124.
- Carver, B.F., Hunger, B.M., Edwards, J.T., Rayas-Buarte, P., Klatt, A.R., Porter, D.R., Seabourn, B.W., Bai, G., Dowell, F.E., Yan, L., Martin, B.C. 2008. Registration of 'Guymon' wheat. *Journal of Plant Registrations*. 2(1):33-35. Available: <http://jpr.sciijournals.org/cgi/reprint/2/1/33>.
- Mornhinweg, D.W., Bregitzer, P.P., Porter, D.R. 2008. Registration of seven spring two-rowed barley germplasm lines resistant to Russian wheat aphid. *Journal of Plant Registrations*. 2(3):230-234.
- Carver, B.F., Hunger, R.M., Edwards, J.T., Porter, D.R., Peeper, T.F., Seabourn, B.W., Rayas-Durate, P., Klatt, A.R., Martin, B.C. 2007. Registration of 'Okfield' wheat. *Journal of Plant Registrations*. 1(2):102-103.
- Mornhinweg, D.W., Bregitzer, P.P., Porter, D.R. 2007. Registration of nineteen spring six-rowed barley germplasm lines resistant to Russian wheat aphid. *Journal of Plant Registrations*. 1(2):137-138.
- Mornhinweg, D.W., Bregitzer, P.P., Porter, D.R. 2007. Registration of seventeen spring two-rowed barley germplasm lines resistant to Russian wheat aphid. *Journal of Plant Registrations*. 1(2):135-136.
- Porter, D.R., Baker, C.A., Carver, B.F., Marais, G.F. 2007. Registration of STARS 0601W wheat germplasm. *Journal of Plant Registrations*. 1(2):170-171.
- Wu, Y., Huang, Y. 2007. An SSR genetic map of *Sorghum bicolor* (L.) Moench and its comparison to a published genetic map. *Genome*. 50:84-89.
- Wu, Y.Q., Huang, Y., Porter, D.R., Tauer, C.G., Holloway, L. 2007. Identification of a major quantitative trait locus conditioning resistance to greenbug biotype E in sorghum PI 550610 using simple sequence repeat markers. *Journal of Economic Entomology*. 100(5):1672-1678.
- Voothuluru, P., Meng, J., Khajuria, C., Louis, J., Zhu, L., Starkey, S., Wilde, G.E., Baker, C.A., Smith, C. 2006. Categories and inheritance of resistance to Russian wheat aphid (Homoptera: Aphididae) biotype 2 in a selection from wheat cereal introduction 2401. *Journal of Economic Entomology*. 99(5):1854-1861.
- Wang, X., Huang, Y., Mort, A.J., Zeng, Y., Tauer, C.G., Cochran, K.D. 2006. Variation of taxane content in needles of *Taxus x media* cultivars with different growth characteristics. *Zeitschrift Fur Naturforschung*. 61c:619-624.

6217-21220-006-00D

ENHANCING DISEASE RESISTANCE AND OIL QUALITY ATTRIBUTES OF PEANUT – Hassan Melouk (P) and Kelly Chamberlin; Stillwater, Oklahoma

- Chamberlin, K.D., Melouk, H.A., Payton, M.E. 2010. Evaluation of the U.S. peanut mini core collection using a molecular marker for resistance to *Sclerotinia minor* Jagger. *Euphytica*. 172:109-115.
- Damicone, J.P., Holbrook Jr, C.C., Smith, D.L., Melouk, H.A., Chamberlin, K.D. 2010. Reaction of the core collection of peanut germplasm to *Sclerotinia* blight and pepper spot. *Peanut Science*. 37(1):1-11.
- Sreedharan, A., Hunger, R.M., Singleton, L.L., Payton, M.E., Melouk, H.A. 2010. Pathogenicity of three isolates of *Rhizoctonia* sp. from wheat and peanut on hard red winter wheat. *International Journal of Agricultural Research*. 5(3):132-147.
- Al-Saleh, M.A., Chamberlin, K.D., Melouk, H.A. 2009. Characterization of TSWV - isoates infecting peanut in Oklahoma and Texas, USA. *Egyptian Journal of Virology*. 6:157-180.
- Chenault, K.D., Maas, A.L., Damicone, J.P., Payton, M.E., Melouk, H.A. 2009. Discovery and characterization of a molecular marker for *Sclerotinia minor* (Jagger) resistance in peanut. *Euphytica*. 166:357-365.
- Bannore, Y.C., Chenault, K.D., Melouk, H.A., El Rassi, Z. 2008. Capillary electrophoresis of some free fatty acids using partially aqueous electrolyte systems and indirect UV detection: Application to the analysis of oleic and linoleic acids in peanut breeding lines. *Journal of Separation Science*. 31:2667-2676.
- Ng, E., Dunford, N.T., Chenault, K.D. 2008. Chemical characteristics and volatile profile of genetically modified peanut cultivars. *Journal of Bioscience and Bioengineering*. 106(4):350-356.
- Baring, M.R., Simpson, C.E., Burow, M.D., Black, M.C., Cason, J.M., Ayers, J., Lopez, Y., Melouk, H.A. 2006. Registration of 'Tamrun OL07' peanut. *Crop Science*. 46:2721.
- Choppakatla, V., Hunger, R.M., Melouk, H.A. 2006. First report of seedling blight caused by *Sclerotium rolfsii* on wheat in Oklahoma. *Plant Disease*. 90:686.
- Isleib, T.G., Rice, P.W., Mazingo II, R.W., Copeland, S.C., Graeber, J.B., Shew, B.B., Smith, D.L., Melouk, H.A., Stalker, H.T. 2006. Registration of N96076L peanut germplasm line. *Crop Science*. 46:2329-2330.

6225-21000-009-00D

DEVELOPMENT AND CHARACTERIZATION OF GENETIC RESOURCES FOR AGRONOMIC AND QUALITY TRAITS USING GENOMIC TOOLS – Anna McClung (P), Rolfe Bryant, Robert Fjellstrom, Helen Miller, and Wengui Yan; Stuttgart, Arkansas

- Agrama, H.A., McClung, A.M., Yan, W. 2011. Population Classification Using Minimum DNA Marker Loci for Maximum Accuracy in Rice (*Oryza sativa* L.). *Molecular Breeding*. DOI:10.1007/s11032-011-9558-x.
- Bryant, R.J., McClung, A.M. 2011. Volatile profile of aromatic and non-aromatic rice cultivars using SPME/GC-MS. *Food Chemistry*. 124:501-513.
- Li, X., Yan, W., Agrama, H., Jia, L., Shen, X., Jackson, A.K., Moldenhauer, K., Yeater, K.M., McClung, A.M., Wu, D. 2011. Mapping QTLs for improving grain yield using the USDA rice mini-core collection. *Planta*. DOI 10.1007/s00425-011-1405-0.
- Nelson, J.C., McClung, A.M., Fjellstrom, R.G., Moldenhauer, K.K., Boza, E., Jodari, F., Oard, J.H., Linscombe, S., Scheffler, B.E., Yeater, K.M. 2011. Mapping QTL main and interaction influences on milling quality in elite U.S. rice germplasm. *Theoretical and Applied Genetics*. 122(2):291-309.
- Somenahally, A., Hollister, E.B., Loeppert, R.H., Yan, W., Gentry, T.J. 2011. Microbial communities in rice rhizosphere altered by intermittent and continuous flooding in fields with long-term arsenic application. *Soil Biology and Biochemistry*. 43(2011):1220-1228.
- Traore, K., McClung, A.M., Chen, M., Fjellstrom, R.G. 2011. Inheritance of Starch Paste Viscosity is Directly Associated with a Rice Waxy Gene Exon 10 SNP Marker. *Journal of Cereal Science*. 53: 37-44.
- Agrama, H.A., Yan, W. 2010. Genetic diversity and relatedness of rice cultivars resistant to straighthead disorder. *Plant Breeding*. 129:304-312.
- Agrama, H.A., Yan, W., Jia, M.H., Fjellstrom, R.G., McClung, A.M. 2010 Genetic structure associated with diversity and geographic distribution in the USDA rice world collection. *Natural Science*. 2(4):247-291. DOI: 10.4236/ns.2010.24036.
- Li, X., Yan, W., Agrama, H., Hu, B., Jia, L., Jia, M.H., Jackson, A.K., Moldenhauer, K., McClung, A.M., Wu, D. 2010. Genotypic and phenotypic characterization of genetic differentiation and diversity in the USDA rice mini-core collection. *Genetica*. 138:1221-1230.
- Jia, Y., Moldenhauer, K.A. 2010. Development of morogenic and digenic rice lines for blast resistance genes of Pi-ta, Pi-kh/Pi-ks. *Journal of Plant Registrations*. 4:163-166.
- Oard, J.H., Moldenhauer, K.K., Fjellstrom, R.G., Nelson, J., Linscombe, S.D., Silva, J., May, G.D. 2010. Registration of the MY2 Cypress/LaGrue rice recombinant inbred line mapping population. *Journal of Plant Registrations*. 4(3):1-5.
- Pillai, T.R., Yan, W., Agrama, H.A., James, W.D., Ibrahim, A.M., Gentry, T.J., McClung, A.M., Loeppert, R.H. 2010. Total grain-arsenic and arsenic-species concentrations in diverse rice cultivars under flooded conditions. *Crop Science*. 50(1):2065-2075.
- Yan, W., Agrama, H., Jia, M.H., Fjellstrom, R.G., McClung, A.M. 2010. Geographic description of genetic diversity and genetic relationships in the USDA Rice World Collection. *Crop Science*. 50:2406-2417. DOI: 10.2135/cropsci2010.02.0096.
- Yan, W., McClung, A.M. 2010. 'Rondo', a long-grain indica rice with resistances to multiple diseases. *Journal of Plant Registrations*. 4:131-136.
- Agrama, H.A., Yan, W. 2009. Association mapping of straighthead disorder induced by arsenic in rice. *Plant Breeding*. 128:551-558.
- Agrama, H.A., Yan, W., Lee, F.N., Fjellstrom, R.G., Chen, M., Jia, M.H., McClung, A.M. 2009. Genetic assessment of a mini-core subset developed from the USDA rice genebank. *Crop Science*. 49:1336-1346.
- Johnson, D.T., Lewis, B.A., Bryant, R.J., Liyanage, R., Lay, J.O., Pszczolkowski, M.A. 2009. Attractants for the Green June Beetle (Coleoptera: Scarabaeidae). *Journal of Economic Entomology*. 102(6):2224-2232.
- Xie, J., Agrama, H.A., Kong, D., Zhuang, J., Hu, B., Wan, Y., Yan, W. 2009. Genetic diversity associated with conservation of endangered Dongxiang wild rice (*Oryza rufipogon*). *Genetic Resources and Crop Evolution*. 57:597-609.
- Yan, W., Li, Y., Agrama, H.A., Luo, D., Gao, F., Ren, G. 2009. Association mapping of stigma and spikelet characteristics in rice (*Oryza sativa* L.). *Molecular Breeding*. 24(3):277-292.
- Bryant, R.J., Anders, M.M., McClung, A.M. 2009. Effect of cultural management practices on the grain quality of two rice cultivars. *Cereal Chemistry*. 86(4):405-409.
- Gao, F.Y., Ren, G.J., Lu, X.J., Sun, S.X., Li, H.J., Gao, Y.M., Luo, H., Yan, W., Zhang, Y.Z. 2008. QTL analysis for resistance to pre-harvest sprouting in rice (*Oryza sativa*). *Plant Breeding*. 127:268-273.
- Prasad, B., Eizenga, G.C. 2008. Sheath blight disease screening methods to identify resistant *Oryza* spp. accessions. *Plant Disease*. 92:1503-1509.
- Tabien, R.E., Samonte, O.B., McClung, A.M. 2008. Forty eight years of rice improvement in Texas since the release of cultivar Bluebonnet in 1944. *Crop Science*. 48:2097-2106.
- Yan, W., Agrama, H.A., Slaton, N.A., Gibbons, J.W. 2008. Soil and plant minerals associated with rice straighthead disorder induced by arsenic. *Agronomy Journal*. 100:1655-1661.

- Agrama, H.A., Eizenga, G.C. 2007. Molecular diversity and genome-wide linkage disequilibrium patterns in a worldwide collection of *Oryza sativa* and its wild relatives. *Euphytica*. 160(3):339-355.
- Agrama, H.A., Eizenga, G.C., Yan, W. 2007. Association mapping of yield and its components in rice cultivars. *Molecular Breeding* 19:341-356.
- Luan, L., Tu, S.B., Long, W.B., Wang, W., Liu, Y.H., Kong, F.L., He, T., Yan, W., Yu, M.Q. 2007. Cytogenetic studies on two F1 hybrids of autotetraploid rice varieties showing extremely high level of heterosis. *Plant Systematics and Evolution*. 267:205-213.
- Rutger, J.N., Bernhardt, L. 2007. Registration of four indica rice genetic stocks. *Crop Science*. 47:461-462.
- Tu, Shengbin, Luan, L., Liu, Y., Long, W., Kong, F., He, T., Xu, Q., Yan, W., Yu, M. 2007. Production and heterosis analysis of rice (*Oryza sativa* L.) autotetraploid hybrids. *Crop Science*. 47:2356-2363.
- Yan, W., Rutger, J.N., Bockelman, H.E., Fjellstrom, R.G., Chen, M.H., Tai, T., McClung, A.M. 2007. Development and evaluation of a core subset of the USDA rice (*Oryza sativa* L.) germplasm collection. *Crop Science*. 47(2):869-878.
- Gibbons, J.W., Moldenhauer, K.K., Gravois, K., Lee, F.N., Bernhardt, J.L., Meullenet, J., Bryant, R.J., Norman, R.J., Cartwright, R., Anders, M., Taylor, K., Bulloch, J., Blocker, M.M. 2006. Registration of 'Medark' Rice. *Crop Science*. 46:2316-2317.
- Gibbons, J.W., Moldenhauer, K.K., Gravois, K., Lee, F.N., Bernhardt, J.L., Meullenet, J.F., Bryant, R.J., Anders, M., Norman, R.J., Cartwright, K., Taylor, K., Bullock, J., Blocker, M.M. 2006. Registration of 'Cybonnet' Rice. *Crop Science*. 46:2317-2318.

6225-21220-004-00D

RESPONSE OF DIVERSE RICE GERMPLASM TO BIOTIC AND ABIOTIC STRESSES – David Gealy (P), Georgia Eizenga, Helen Miller, Anna McClung, and Yulin Jia; Stuttgart, Arkansas

- Brooks, S.A., Anders, M.M., Yeater, K.M. 2010. Effect of furrow irrigation on the severity of false smut in susceptible rice varieties. *Plant Disease*. 94(5):570-574.
- Costanzo, S., Jia, Y. 2010. Sequence variation at the rice blast resistance gene Pi-km locus: Implications for the development of allele specific markers. *Plant Science*. 178-523-530.
- Dai, Y., Jia, Y., Correll, J.C., Wang, X., Wang, Y. 2010. Diversification and evolution of the avirulence gene AVR-Pita1 in field isolates of *Magnaporthe oryzae*. *Fungal Genetics and Biology*. 47:973-980.
- Gealy, D.R., Fischer, A.J. 2010. 13C discrimination: A stable isotope method to quantify root interactions between C3 rice (*Oryza sativa*) and C4 barnyardgrass (*Echinochloa crus-galli*) in flooded fields. *Weed Science*. 58(3):359-368.
- Goss, B.L., Reagon, M., Hsu, S., Caicedo, A.L., Jia, Y., Olsen, K.M. 2010. Seeing red: The origin of grain pigmentation in US weedy rice. *Molecular Ecology*. 19:3380-3393.
- Reagon, M., Thurber, C.S., Gross, B.L., Olsen, K.M., Jia, Y., Caicedo, A.L. 2010. Genomic patterns of nucleotide diversity in divergent populations of U.S. weedy rice. *BMC Evolutionary Biology*. 10(180):1-16.
- Shivrain, V.K., Burgos, N.R., Agrama, H.A., Lawton-Rauh, A., Lu, B., Sales, M.A., Boyett, V.A., Gealy, D.R., Moldenhauer, K.K. 2010. Genetic diversity of weedy red rice (*Oryza sativa* L.) in Arkansas, USA. *Weed Research*. 50:289-302.
- Thurber, C.S., Reagon, M., Gross, B.L., Olsen, K.M., Jia, Y., Caicedo, A.L. 2010. Molecular evolution of the sh4 shattering locus in U.S. weedy rice. *Molecular Ecology*. 19:3271-3284.
- Wang, X., Fjellstrom, R.G., Jia, Y., Yan, W., Jia, M.H., Scheffler, B.E., Wu, D., Shu, Q., McClung, A.M. 2010. Characterization of Pi-ta Blast resistance gene in an international rice core collection. *Plant Breeding*. 129(5):491-501.
- Zhao, K., Wright, M.K., Kimball, J.A., Eizenga, G.C., McClung, A.M., Kovach, M.J., Tyagi, W., Ali, M., Tung, C., Reynolds, A., Bustamante, C.D., McCouch, S.R. 2010. Genomic diversity and introgression in *O. sativa* reveal the impact of domestication and breeding on the rice genome. *PLoS One*. 5(5):e10780. doi:10.1371/journal.pone.0010780.
- Brooks, S.A., Anders, M.A., Yeater, K.M. 2009. Effect of cultural management practices on the severity of false smut and kernel smut of rice. *Plant Disease*. 93: 1202 – 1208.
- Costanzo, S., Jia, Y. 2009. Alternatively spliced transcripts of Pi-ta blast resistance gene in *Oryza sativa*. *Plant Science*. 177:468-478.
- Gealy, D.R., Agrama, H., Eizenga, G.C. 2009. Exploring genetic and spatial structure of U.S. weedy red rice (*Oryza sativa* L.) populations in relation to rice relatives worldwide. *Weed Science*. 57:627-643.
- Han, Z., Wu, F., Deng, G., Qian, G., Yu, M., Jia, Y. 2009. Structural and expressional analysis of the B-hordein genes in Tibetan hull-less barley. *Genetica*. 138-227-239.
- Jia, Y. 2009. A user friendly method to isolate and single spore the fungi *Magnaporthe oryzae* and *Magnaporthe grisea* obtained from diseased field samples. *Plant Health Progress*. Page 1-4. doi:10.1094/PHP-2009-1215-01-BR, Online.

- Jia, Y. 2009. Artificial introgression of a large fragment around the Pi-ta rice blast resistance gene in backcross progenies and several elite rice cultivars. *Heredity*. 103(4):333-339.
- Jia, Y., Liu, G., Costanzo, S., Lee, S., Dai, Y. 2009. Current progress on genetic interactions of rice with rice blast and sheath blight fungi. *Frontiers of Agriculture in China*. 3(3):231-239.
- Lee, S., Costanzo, S., Jia, Y., Olsen, K.M., Caicedo, A.L. 2009. Evolutionary dynamics of the genomic region around the blast resistance gene Pi-ta in AA genome *Oryza* species. *Genetics*. 183:1315-1325.
- Liu, G., Jia, Y., Correa-Victoria, F.J., Prado, G.A., Yeater, K.M., McClung, A.M., Correll, J.C. 2009. Mapping quantitative trait loci responsible for resistance to rice sheath blight disease using greenhouse assays. *Phytopathology*. 99(9):1078-1084.
- Shivrain, V.K., Burgos, N.R., Gealy, D.R., Salesa, M.A., Smith, K.L. 2009. Gene flow from weedy red rice (*Oryza sativa* L.) to cultivated rice and fitness of hybrids. *Pest Management Science*. 65(10):1124-1129.
- Shivrain, V.K., Burgos, N.R., Gealy, D.R., Smith, K.L., Scott, R.C., Mauromoustakos, A., Black, H.L. 2009. Red rice (*Oryza sativa* L.) emergence characteristics and influence on rice (*O. sativa*) yield at different planting dates. *Weed Science*. 57(1):94-102.
- Shivrain, V.K., Burgos, N.R., Salesa, M.A., Mauromoustakos, A., Gealy, D.R., Smith, K.L., Black, H.L., Jia, M.H. 2009. Factors affecting the outcrossing rate between Clearfield rice and red rice (*Oryza sativa*). *Weed Science*. 57:394-403.
- Jia, Y., Lee, F.N., McClung, A.M. 2009. Determination of resistance spectra of the Pi-ta and Pi-k genes to U.S. races of *Magnaporthe oryzae* causing rice blast in a recombinant inbred line population. *Plant Disease*. 93(6):639-644.
- Lee, S., Wasmishe, Y., Jia, Y., Liu, G., Jia, M.H. 2009. Identification of two major resistance genes against race IE-1k of *Magnaporthe oryzae* in the indica rice cultivar ZHE733. *Molecular Breeding*. 24:127-134.
- Shivrain, V.K., Burgos, N.R., Gealy, D.R., Baquiereza, C.J. 2008. Maximum outcrossing rate and genetic compatibility between red rice (*Oryza sativa*) biotypes and Clearfield rice. *Weed Science*. 56(6):807-813.
- Brooks, S.A., Yan, W., Jackson, A.K., Deren, C.W. 2008. A natural mutation in *Rc* reverts white-rice-pericarp to red and results in a new, dominant, wild-type allele: *Rc-g*. *Theoretical and Applied Genetics*. 117(4):575-580.
- Jia, Y., Marin, R. 2008. Identification of a new locus *Ptr(t)* required for rice blast resistance gene Pi-ta-mediated resistance. *Molecular Plant-Microbe Interactions*. 21(4):396-403.
- Li, W., Lei, C., Cheng, Z., Jia, Y., Huang, D., Liu, Z., Wang, J., Shi, K., Zhang, X., Su, N., Guo, X., Zhai, H., Wan, J. 2008. Identification of SSR markers for a broad-spectrum blast resistance gene Pi-20(t) for marker-assisted breeding. *Molecular Breeding*. 22:141-149.
- Wang, X., Jia, Y., Shu, Q., Wu, D. 2008. Haplotype diversity at the Pi-ta locus in cultivated rice and its wild relatives. *Phytopathology*. 98:1305-1311.
- Brooks, S.A. 2007. Sensitivity to a phytotoxin from *Rhizoctonia solani* correlates with sheath blight susceptibility in rice. *Phytopathology*. 97:1207-1212.
- Chen, C., Yun, K., Ransom, H., Mohanty, B., Bajic, V.B., Jia, Y., Yun, S., De Los Reyes, B.G. 2007. Reactive oxygen species trigger a regulatory module involved in the early responses of rice seedlings to cold stress. *Biomed Central (BMC) Genomics*. 18:175.
- Olsen, K.M., Caicedo, A.L., Jia, Y. 2007. The evolutionary genomics of weedy red rice in the U.S.A. *Journal of Integrative Plant Biology*. 49:811-816.
- Venu, R.C., Jia, Y., Gowda, M., Jia, M.H., Jantasuriyarat, C., Stahlbert, E., Li, H., Rhineheart, A., Reddy, P., Singh, P., Rutger, J.N., Kudrna, D., Wing, R., Nelson, J.C., Wang, G. 2007. RL-sage and microarray analysis of the rice defense transcriptome after *Rhizoctonia solani* infection. *Molecular Genetics and Genomics*. 278:421-431.
- Zhou, E., Jia, Y., Singh, P., Correll, J.C., Lee, F.N. 2007. Instability of the *Magnaporthe oryzae* Avirulence gene AVR-Pita alters virulence. *Fungal Genetics and Biology*. 44:1024-1034.
- Jia, Y., Coarrea-Victoria, F., McClung, A.M., Zhu, L., Liu, G., Wasmishe, Y., Xie, J., Marchetti, M.A., Pinson, S.R., Rutger, J.N., Correll, J.C. 2006. Rapid determination of rice cultivar responses to the sheath blight pathogen *Rhizoctonia solani* using a micro-chamber screening method. *Plant Disease*. 91:485-489.

6402-21000-031-00D

COMBINING CONVENTIONAL AND MOLECULAR TECHNIQUES TO IMPROVE SEED COTTON VALUE – Jodi Scheffler (P), Lawrence Young, and Rickie Turley; Stoneville, Mississippi

- Romano, G.B., Sacks, E.J., Stetina, S.R., Robinson, A.F., Fang, D.D., Gutierrez, O., Scheffler, J.A. 2009. Identification and Genomic Location of a Reniform Nematode (*Rotylenchulus reniformis*) Resistance locus (*Ren_ari*) Introgressed from *Gossypium aridum* into Upland Cotton (*G. hirsutum*). *Theoretical and Applied Genetics*. 120:139-150.
- Romano, G.B., Scheffler, J.A. 2009. Lowering Seed Gossypol Content in Glanded Cotton (*Gossypium hirsutum* L.) Lines. *Plant Breeding*. 127(6):619-624.

- Benbiuza, H., Lognay, G., Scheffler, J.A., Baudoin, J., Mergeai, G.G. 2009. Expression of the Glanded-Plant and Glandless-Seed" Trait of Australian Diploid Cottons in Different Genetic Backgrounds. *Journal of Agricultural and Food Chemistry*. 165:211-221"
- Scheffler, J.A., Romano, G.B. 2008. Modifying Gossypol in Cotton (*Gossypium hirsutum* L.): Cost Effective Gossypol Analysis Method for Small Samples. *Journal of Cotton Science*. 12(3):202-209
- Turley, R.B. 2008. Expression of a Phenylcoumaran Benzylic Ether Reductase - Like Protein in the Ovules of *Gossypium hirsutum*. *Biologia Plantarum*. 52:759-762.
- Turley, R.B., Taliercio, E.W. 2008. Cotton Benzoquinone Reductase: Up-Regulation During Early Fiber Development and Heterologous Expression and Characterization in *Pichia Pastoris*. *Plant Physiology and Biochemistry*. 46:780-785
- Zhang, J., Turley, R.B., Stewart, J.M. 2007. Comparative Analysis of Gene Expression Between CMS-D8 Restored Plants and Normal Non-restoring Fertile Plants in Cotton by Differential Display. *Plant Cell Reports*. 27:553-561
- Taliercio, E.W., Boykin, D.L. 2007. Analysis of Gene Expression in Cotton Fiber Initials. *Biomed Central (BMC) Plant Biology*. 7:22.
- Blenda, A.V., Scheffler, J.A., Scheffler, B.E., Palmer, M., Lacape, J., Yu, J., Jesudraui, C., Jung, S., Muthukumar, S., Yellambalase, P., Ficklin, S., Staton, M., Eschelman, R., Ulloa, M., Saha, S., Burr, B., Lui, S., Zhang, T., Fang, D., Pepper, A.E., Kumpatla, S.P., Jacobs, J., Tomkins, J.P., Cantrell, R.G., Main, D. 2006. CMD: A Cotton Microsatellite Database Resource for *Gossypium* Genomics. *Biomed Central (BMC) Genomics*. 7:132.

6402-21000-033-00D

GENETIC-PHYSIOLOGICAL TEAM RESEARCH TO IMPROVE PRODUCTION, FIBER QUALITY AND COMPETITIVE ABILITY OF COTTON – Lawrence Young (P), Efrem Bechere, and Linghe Zeng; Stoneville, Mississippi

- Bechere, E., Auld, D.L., Krifa, M., Smith, C.W., Cantrell, R.G. 2011. Registration of TTU-0782 Upland Cotton Germplasm Line with Superior Fiber Quality. *Journal of Plant Registrations*. 5 (2):207-2010.
- Bechere, E., Boykin Jr, J.C., Meredith Jr, W.R. 2011. Evaluation of Cotton Genotypes for Ginning Energy and Ginning Rate. *Journal of Cotton Science*. 15:11-21.
- Pettigrew, W.T. 2010. Effects of Foliar Fertilizer and Mepiquat Pentaborate on Early Planted Cotton Growth and Lint Production. *Crop Management*. doi:10.1094/CM-2010-0215-01-RS.
- Bechere, E., Auld, D.L., Dotary, P., Kebede, H.A. 2010. Registration of Four Upland Cotton (*Gossypium hirsutum* L.) Genetic Stock Mutants with Tolerance to Imazamox. *Journal of Plant Registrations*. 4:155-158
- Zeng, L., Meredith Jr, W.R. 2010. Neppiness in an Introgressed Population of Cotton: Genotypic Variation and Genotypic Correlation. *Journal of Cotton Science*. 14:17-25.
- Bechere, E., Auld, D.L., Dotrux, P.A., Gilbert, L.V., Kebede, H.A. 2009. Imazamox Tolerance in Mutation Derived Lines of Upland Cotton. *Crop Science*. 49:1586-1592.
- Bechere, E., Auld, D.L., Hequet, E. 2009. Development of Naked-Tufted" Seed Coat Mutants for Potential Use in Cotton Production. *Euphytica*. 167:333-339".
- Zeng, L., Meredith Jr, W.R., Gutierrez, O.A., Boykin, D.L. 2009. Identification of Association Between SSR Markers and Fiber Related Traits in Exotic Germplasm Derived from Multiple Crosses among *Gossypium* Tetraploid Species. *Theoretical and Applied Genetics*. 119:93-103.
- Pettigrew, W.T., Meredith Jr, W.R. 2009. Seed Quality and Plant Date Effects on Cotton Lint Yield Components, and Fiber Quality. *Journal of Cotton Science*. 13:11-24.
- Pettigrew, W.T., Molin, W.T., Stetina, S.R. 2009. Impact of Varying Planting Dates and Tillage Systems on Cotton Growth and Lint Yield Production. *Agronomy Journal*. 101:1131-1138.
- Zeng, L., Meredith Jr, W.R. 2009. Associations Among Lint Yield, Yield Components, and Fiber Properties in an Introgressed Population of Cotton. *Crop Science*. 49:1647-1654.
- Zeng, L., Meredith Jr, W.R. 2009. Registration of Five Exotic Germplasm Lines of Cotton Derived From Multiple Crosses Among *Gossypium* Tetraploid Species. *Journal of Plant Registrations*. 3:77-80.
- Pettigrew, W.T. 2008. Potassium Influences on Yield and Quality Production for Maize, Wheat, Soybean and Cotton. *Physiologia Plantarum*. 133:670-681.
- Pettigrew, W.T. 2008. The Effect of Higher Temperatures on Cotton Lint Yield Production and Fiber Quality. *Crop Science*. 48:278-285.
- Pettigrew, W.T., Meredith Jr, W.R., Bruns, H.A., Stetina, S.R. 2007. Effects of a Short-term Corn Rotation on Cotton Dry Matter Partitioning, Lint Yield, and Fiber Quality Production. *Journal of Cotton Science*. 10:244-251.
- Zeng, L., Meredith Jr, W.R., Boykin, D.L., Taliercio, E. 2007. Evaluation of an Exotic Germplasm Population Derived From Multiple Crosses Among *Gossypium* Tetraploid Species. *Journal of Cotton Science*. 11:118-127.

6402-21000-034-00D

DEVELOPMENT OF SOYBEAN GENEOTYPES AND MANAGEMENT SYSTEMS FOR EARLY SEASON AND STRESS ENVIRONMENTS – Lawrence Young (P), Anne Gillen, and Nacer Bellaloui; Stoneville, Mississippi

- Bellaloui, N., Smith, J.R., Gillen, A.M., Ray, J.D. 2010. Effect of maturity on seed sugars in the early soybean production system as measured on near-isogenic soybean (Glycine max). *Crop Science*. 49:608-620.
- Bruns, H.A., Abbas, H.K. 2010. Additional Potassium Did Not Decrease Aflatoxin or Fumonisin nor Increase Corn Yields. *Plant Management Network*. doi: 10.1094/CM-2010-0216-01-RS.
- Bruns, H.A., Abbas, H.K. 2010. Comparisons of Herbicide Treated and Cultivated Herbicide-Resistant Corn. *Field Crops Research*. doi:10.1155/2010/798127.
- Zeng, L., Meredith Jr, W.R. 2010. Germplasm Potential for Continuing Improvement of Fiber Quality in Upland Cotton: Combining Ability for Lint Yield and Fiber Quality. *Crop Science*. 51:60-68.
- Bellaloui, N., Reddy, K.N., Abel, C.A., Gillen, A.M. 2010. Nitrogen metabolism and seed composition as influenced by foliar boron application in soybean. *Plant and Soil Journal*. doi 10.1007/s11104-010-0455-6.
- Bellaloui, N., Smith, J.R., Ray, J.D., Gillen, A.M. 2009. Effect of Maturity on Seed Composition in the Early Soybean Production System as Measured on Near-isogenic Soybean Lines. *Crop Science*. 49:608-620
- Bellaloui, N., Abbas, H.K., Gillen, A.M., Abel, C.A. 2009. Effect of Glyphosate-boron Application on Seed Composition and Nitrogen Metabolism in Glyphosate-resistant Soybean. *Journal of Agriculture and Food Chemistry*. DOI:10.1021/jf901801z.
- Bellaloui, N., Reddy, K.N., Zablutowicz, R.M., Abbas, H.K., Abel, C.A. 2009. Effects of Glyphosate Application on Seed Iron and Root Ferric (III) Reductase in Soybean Cultivars. *Journal of Agriculture and Food Chemistry*. 57:9569-9574.
- Bruns, H.A., Abbas, H.K. 2009. Clipping Corn Plants at the 3 and 5 Leaf Growth Stage Fail to Simulate Uneven Emergence. *Crop Management*. doi:10.1094/CM-2009-1016-01-RS.
- Bellaloui, N., Hanks, J.E., Fisher, D.K., Mengistu, A. 2009. Soybean seed composition is influenced by with-in field variability in soil nutrients. *Crop Management*. doi:10.1094/CM-2009-1203-01-RS.
- Gillen, A.M., Strausbaugh, C.A., Tindall, K.V. 2008. Evaluations of Beta corolliflora for Resistance to Curly Top in Idaho. *Journal of Sugar Beet Research*. 45(3&4):99-118.
- Bellaloui, N., Mengistu, A., Paris, R.L. 2008. Soybean Seed Composition in Cultivars Differing in Resistance to Charcoal Rot (Macrophomina phaseolina). *Journal of Agricultural Science*. vol. 146(6):667-675.
- Bellaloui, N., Zablutowicz, R.M., Reddy, K.N., Abel, C.A. 2008. Nitrogen Metabolism and Seed Composition as Influenced by Glyphosate Application in Glyphosate-Resistant Soybean. *Journal of Agriculture and Food Chemistry*. 56: 2765-2772.

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REDUCING SOYBEAN YIELD LOSSES THROUGH GENETIC IMPROVEMENT – Jeffery Ray (P), Lawrence Young, Shuxian Li, and James Smith; Stoneville, Mississippi

- Charlson, D.V., Bhatnagar, S., King, C.A., Ray, J.D., Sneller, C.H., Carter Jr, T.E., Purcell, L.C. 2009. Polygenic Inheritance of Canopy Wilting in Soybean [Glycine max (L.) Merr.]. *Theoretical and Applied Genetics*. DOI 10.1007/s11033-009-9496-4.
- Lee, J., Woolard, M., Sleper, D.A., Smith, J.R., Pantalone, V.R., Nyinyi, C.N., Cardinal, A., Shannon, G.J. 2009. Environmental Effects on Oleic Acid in Soybean Seed Oil of Plant Introductions with Elevated Oleic Concentration. *Crop Science*. 49:1762-1768.
- Li, S., Boykin, D.L., Sciombato, G., Wrather, A., Shannon, G., Sleper, D. 2009. Reaction of Soybean Cultivars to Phomopsis Seed Decay in the Mississippi Delta, 2007. *Plant Disease Management Reports*. DOI:10.1094/PDMR03:FC096.
- Li, S., Chen, P., Rupe, J., Wrather, A. 2010. Reaction of Maturity Group V Soybean Plant Introductions to Phomopsis Seed Decay in Arkansas, Mississippi, and Missouri, 2009. *Plant Disease Management Reports*. 4:ST034.
- Li, S., Hartman, G.L., Boykin, D.L. 2010. Aggressiveness of Phomopsis longicolla and other Phomopsis species on soybean. *Plant Disease*. 94:1035-1040.
- Li, S., Moon, J.S., Lee, S.H., Domier, L.L. 2009. First Report of Soybean Yellow Mottle Mosaic Virus in Soybean in North America. *Plant Disease*. 93 (11):1214.
- Li, S., Rupe, J., Chen, P., Wrather, A. 2010. Reaction of maturity group IV soybean plant introductions to Phomopsis Seed Decay in Arkansas, Mississippi, and Missouri, 2009. *Plant Disease Management Reports*. 4:ST035
- Li, S., Wrather, A., Rupe, J., Chen, P. 2010. Reaction of maturity group III soybean plant introductions to Phomopsis seed decay in Arkansas, Mississippi and Missouri, 2009. *Plant Disease Management Reports*. 4:ST036.

- Li, S., Young, L.D. 2009. Evaluation of Selected Genotypes of Soybean for Resistance to *Phakopsora pachyrhizi*. Plant Health Progress. doi:1094/PHP-2009-0615-01-RS
- Lygin, A., Li, S., Vittal, R., Hartman, G.L., Widholm, J., Lozovaya, V. 2009. The Importance of Phenolic Metabolism to Limit the Growth of *Phakopsora pachyrhizi*. Phytopathology. 99:1412-1420.
- Meng, J., Xu, Y., Li, S., Li, C., Zhang, X., Chen, P. 2010. Effects of Conventional and Conservation Tillage Systems on Soybean Growth and Soil Microbial Populations. Journal of Crop Improvement. 337-348.
- Qian, H., Li, S., Xu, Y., Li, C., Sun, Y. 2011. Assessment of the effects of *Hirsutella minnesotensis* on Soybean Cyst Nematode and growth of soybean. Soybean Science. 30(2):266-271.
- Ray, J.D., Sinclair, T.R., Glaz, B.S. 2010. Sugarcane Response to High Water Tables and Intermittent Flooding. Journal of Crop Improvement. 24:12-27.
- Ray, J.D., Wilfrid, M., Smith, J.R., Frederick, R.D., Miles, M.R. 2009. Genetics and mapping of adult plant rust resistance in soybean PI 587886 and PI 587880A. Theoretical and Applied Genetics. 119:271-280.
- Li, S. 2009. Reaction of Soybean Rust Resistant Lines Identified in Paraguay to Mississippi Isolates of *Phakopsora pachyrhizi*, the Causal Agent of Soybean Rust. Crop Science. 49(3):887-894
- Li, S., Hartman, G.L., Chen, Y. 2009. Evaluation of Aggressiveness of *Fusarium virguliforme* Isolates That Cause Soybean Sudden Death Syndrome. Journal of Plant Pathology. 91(1):77-86
- Li, S., Hartman, G.L., Domier, L.L., Boykin, D.L. 2008. Quantification of *Fusarium solani* f. sp. *glycines* Isolates in Soybean Roots by Colony-forming Unit Assays and Real-time Quantitative PCR. Journal of Theoretical and Applied Genetics. 117:343-352.
- Li, S., Lygin, A., Zernova, O., Lozovaya, V., Hartman, G.L., Widholm, J.M. 2008. Genotype Response of Soybean (*Glycine max*) Whole Plants and Hairy Roots to *Fusarium solani* f. sp. *glycines* Infection. Soybean Science. 27:275-282.
- Seversike, T.M., Ray, J.D., Shultz, J.L., Purcell, L.C. 2008. Soybean Molecular Linkage Group B1 Cooresponds to Classical Linkage Group 16 Based on Map Location of the 1f2 Gene. Journal of Theoretical and Applied Genetics. 117:143-147
- Shultz, J.L., Ray, J.D., Smith, J.R. 2008. Mapping Two Genes in the Purine Metabolism Pathway of Soybean.. DNA Research. 19(3):264-269
- Smith, J.R., Mengistu, A., Nelson, R.L., Paris, R.L. 2008. Identification of Soybean Accessions with High Germinability in High-temperature Environments. Crop Science. 48:2279-2288
- Li, S., Moore, W.F., Spinks, B.L., Wells, B.C., Sciumbato, G.L., Robinson, S.J., Libous Bailey, L.M. 2007. Occurrence of Asian Soybean Rust Caused by *Phakopsora pachyrhizi* in Mississippi. Plant Health Progress. doi:10.1094/PHP-2007-0917-02-BR
- Shultz, J.L., Ray, J.D., Lightfoot, D.A. 2007. A Sequence Based Synteny Map Between Soybean and Arabidopsis Thaliana. Genome. DOI:10.1186/1471-2164-8-8
- Shultz, J.L., Ray, J.D., Smith, J.R., Mengistu, A. 2007. A soybean mapping population specific to the early soybean production system. DNA Sequence. 18(2):104-111.

6402-21310-003-00D

Genomics and Bioinformatics Research in Agriculturally Important Organisms – Brian Scheffler (P); Stoneville, Mississippi

- Arias De Ares, R.S., Ray, J.D., Mengistu, A., Scheffler, B.E. 2011. Discriminating microsatellites from *Macropmina phaseolina* and their potential association to biological functions.. Plant Pathology. DOI: 10.1111/j.1365-3059.2010.02421.x.
- Narina, S.S., Buyyarapu, R., Kottapalli, R., Sartie, A.M., Moahmed, A., Asiedu, R., Mignouna, H., Sayre, B., Scheffler, B.E. 2011. Generation and analysis of expressed sequence tags(ESTs) for marker development in yam (*Dioscorea alata* L.). Biomed Central (BMC) Genomics. 1471-2164.
- Rinehart, T., Trigiano, R., Wadl, P., Hadziabdic, D., Pooler, M., Scheffler, B. 2010. Characterization of Twelve Microsatellite Markers for the Native Redbud Tree (*Cercis canadensis*). Molecular Ecology Resources. 10(4), 751-754. <http://tomato.biol.trinity.edu/manuscripts/10-4/mer-09-0487.pdf>.
- Blacklock, B.J., Scheffler, B.E., Shepard, M.R., Jayasuriya, N., Minto, R.E. 2010. Functional Diversity in Fungal Fatty Acid Synthesis. The First Acetylase from the Pacific Golden Chanterelle, *Cantharellus formosus*. Journal of Biological Chemistry. 287(37):28442-28449.
- Arias De Ares, R.S., Techen, N., Rinehart, T.A., Olsen, R.T., Kirkbride, J.H., Scheffler, B.E. 2010. Development of SSR markers of *Chionanthus retusus* (Oleaceae) and cross amplification of closely related taxa. HortScience. 46(1):23-29.
- Fang, X., Dong, W., Thornton, C., Scheffler, B.E., Willet, K. 2010. Benzo(A)Pyrene Induced Glycine N-Methyltransferase Messenger Rna Expression in *Fundulus heteroclitus* Embryos. Marine Environmental Research. 69:274-276

- Han, K.M., Strauss, S.H., Dharmawardhana, P., Arias De Ares, R.S., Ma, C., Busov, V. 2010. Gibberellic acid cisgenes modify growth rate and stature in transgenic poplars. *Plant Biotechnology Journal* pp. 1–17
- Techen, N., Arias De Ares, R.S., Glynn, N.C., Pan, Z., Khan, I.A., Scheffler, B.E. 2010. Optimized Construction of SSR-enriched Libraries. *Molecular Ecology Resources*. 10(3):508-515
- Wadl, P.A., Dean, D., Li, Y., Vito, L.M., Scheffler, B.E., Hadziabdic, D., Windham, M.T., Trigiano, R.N. 2010. Development and characterization of microsatellites for switchgrass rust fungus (*Puccinia emaculata*). *Conservation Genetics*. (2011) 3:185–188. Published on line
- Wang, X., Dean, D., Wadl, P., Hadziabdic, D., Scheffler, B.E., Rinehart, T.A., Trigiano, R. 2010. Development of microsatellite markers from Crape Myrtle (*Lagerstroemia* L.). *HortScience*. VOL. 45(5).
- Yuan, J.S., Abercrombie, L., Cao, Y., Halfhill, M.D., Zhou, X., Peng, Y., Hu, J., Rao, M.R., Heck, G.R., Larosa, T.J., Sammons, R.D., Wang, X., Ranjan, P., Johnson, D.H., Wadl, P.A., Scheffler, B.E., Rinehart, T.A., Trigiano, R.N., Stewart, Jr, C.N. 2010. Functional Genomics Analysis of Horseweed (*Conyza canadensis*) with Special Reference to the Evolution of Non-Target-Site Glyphosate Resistance. *Weed Science*. 58:109-117.
- Arias De Ares, R.S., Stetina, S.R., Tonos, J.L., Scheffler, J.A., Scheffler, B.E. 2009. Microsatellites Reveal Genetic Diversity in *Rotylenchulus reniformis* Populations. *Journal of Nematology*. 41(2): 146-156.
- Arias De Ares, R.S., Ballard, L.L., Scheffler, B.E. 2009. UPIC: How Many SSR Markers to Run. *Bioinformatics* 3(8):352-360.
- Bowers, S., Gandy, S., Anderson, B., Ryan, P., Willard, S. 2009. Assessment Of Pregnancy in the Late Gestation Mare Using Digital Infrared Thermography. *Theriogenology*. 72(3):372-377.
- Hu, Y., Willett, K.L., Khan, I.A., Scheffler, B.E., Dasmahapatra, A.K. 2009. Ethanol Disrupts Chondrification of the Neurocranial Cartilages in Medaka Embryos without Affecting Aldehyde Dehydrogenase 1A2 (*Aldh1A2*) Promoter Methylation. *Comparative Biochemistry and Physiology*. 150(4):495-502.
- Jung, S.Y., Bowers, S.D., Willard, S.T. 2009. Simulated Microgravity Influences Bovine Oocyte In Vitro Fertilization and Preimplantation Embryo Development. *Journal of Animal and Veterinary Advances* 8(9):1807-1814.
- Jung, S.Y., Willard, S.T. 2009. The Effect of Culture Methods and Serum Supplementation on Developmental Competence of Bovine Embryos Cultured In Vitro. *Journal of Animal and Veterinary Advances*. 8(1):67-74.
- Moulton, K., Ryan, P., Christiansen, D., Hopper, R., Klauser, C., Bennett, W., Rodts-Palenik, S., Willard, S. Hormonal Profiles of Late Gestation Ewes Following Intra-Uterine Inoculation With and Without Lux-Modified *Escherichia Coli*. *Journal of Reproduction and Development* 55(1):55-62.
- Moulton, K., Ryan, P., Lay Jr, D.C., Willard, S. 2009. Postmortem Photonic Imaging of Lux-Modified *Salmonella Typhimurium* Within the Gastrointestinal Tract of Swine Following Oral Inoculation In Vivo. *Journal of Animal Science* 87:2239-2244.
- Ryan, P.L., Christiansen, D.L., Hopper, R.M., Bagnell, C.A., Vaala, W.E., Leblanc, M.M. 2009. Evaluation of Relaxin Blood Profiles of Horses as A Means of Assessing Placental Function in High-Risk Pregnancies And Responsiveness to Therapeutic Strategies. *Annals of the New York Academy of Sciences*. 1160:169-178.
- Techen, N., Pan, Z., Scheffler, B.E., Kahn, I.A. 2009. Detection of *Illicium anisatum* as Adulterant of *Illicium verum*. *Planta Medica*. 75:392–395.
- Joshi, V., Techen, N., Scheffler, B.E., Khan, I.A. 2009. Identification and Differentiation Between *Hoodia gordonii* Masson) Sweet ex Decne., *Opuntia ficus indica* (L.) P. Miller and Other Related *Hoodia* Species by Microscopy and PCR Methods. *Journal of Herbs, Spices and Medicinal Plants*. 15(3):253-264.
- Schlueter, J.A., Scheffler, B.E., Jackson, S., Shoemaker, R.C. 2008. Fractionation of Synteny in a Genomic Region Containing Tandemly Duplicated Genes Across *Glycine max*, *Medicago truncatula* and *Arabidopsis thaliana*. *Journal of Heredity*.
- Dong, W., Wang, L., Thornton, C., Scheffler, B.E., Willett, K.L. 2008. Benzo(A)pyrene Decreases Brain and Ovarian Aromatase mRNA Expression. *Aquatic Toxicology*.

6404-21000-008-00D

SMALL FRUIT AND ORNAMENTAL GENETIC RESEARCH FOR THE MID-SOUTH – Timothy Rinehart (P), Cecil Pounders, James Spiers, Stephen Stringer, and Hamidou Sakhanokho; Poplarville, Mississippi

- Stringer, S.J., Marshall, D.A., Gray, D. 2011. Eudora muscadine grape. *HortScience*. 46(1).
- Wang, X., Wadl, P.A., Cabrera, R.I., Pounders Jr, C.T., Scheffler, B.E., Pooler, M.R., Rinehart, T.A., Trigiano, R.N. 2011. Evaluation of genetic diversity and pedigree within crapemyrtle (*Lagerstroemia* spp.) cultivars using simple sequence repeat (SSR) markers. *Journal of the American Society for Horticultural Science*. 136(2):116-128.
- Baird, R., Mcneil, D., Wadl, P.A., Trigiano, R.N., Allen, T., Shier, T., Wang, X., Rinehart, T.A., Abbas, H.K. 2010. Variability of United States Isolates of *Macrophomina phaseolina* Based on Simple Sequence Repeats and Cross Genus Transferability to Related *Botryosphaeraceae*. *Mycopathologia*. 170.3(2010): 169-80. Epub 2010 March 30.

- Johnson, D., Fitzpatrick, B., Wang, X., Wadl, P., Rinehart, T.A., Ownley, B., Windham, M., Trigiano, R. 2010. Analysis of genetic diversity of flowering dogwood natural stands using microsatellites: the effects of dogwood anthracnose. *Genetica*. 138:1047-1057.
- Pounders Jr, C.T., Blythe, E.K., Fare, D.C., Knox, G.W., Sibely, J.L. 2010. Crapemyrtle genotype × environment interactions and trait stability for plant height leaf-out and flowering. *HortScience*. 45(2):198-207.
- Rinehart, T.A., Scheffler, B.E., Reed, S.M. 2010. Ploidy Variation and Genetic Diversity in *Dichroa*. *HortScience*. 45:208-213.
- Stringer, S.J., Draper, A., Marshall, D.A., Spiers, J.M. 2010. Prince Rabbiteye Blueberry. *HortScience*. Vol.45(1):144-145.2010.
- Sylvere Sie, R., Gilbert, C., Sakhanokho, H.F., Toueix, Y., Dje, Y., Sangare, A., Branchard, M. 2010. Protocols for Callus and Somatic Embryo Initiation for *Hibiscus sabdariffa* L. (Malvaceae): Influence of Explant Type, Sugar, and Plant Growth Regulators. *Australian Journal of Crop Science*. 4(2):98-105.
- Wadl, P., Wang, X., Moulton, J.K., Hokanson, S.C., Skinner, J.A., Rinehart, T.A., Reed, S.M., Pantalone, V., Trigiano, R. 2010. Transfer of Flowering and Kousa Dogwood Simple Sequence Repeats (SSRs) to Selected *Cornus* (Cornaceae) Species. *Journal of the American Society for Horticultural Science*. 135:279-288.
- Kardos, J.H., Rodacker, C.D., Dirr, M.A., Rinehart, T.A. 2009. Production and Verification of *Hydrangea macrophylla* × *angustipetala* hybrids. *HortScience*. 44: 1534-1537.
- Li, Y., Mmbaga, M.T., Windham, A.S., Windham, M.T., Trigiano, R.N. 2009. Powdery Mildew of Dogwoods: Current Status and Future Prospects. *Plant Disease*. <http://apsjournals.apsnet.org/doi/pdf/10.1094/PDIS-93-11-1084>.
- Li, Y., Windham, M., Trigiano, M., Rinehart, T.A., Reed, S.M., Spiers, J.M. 2009. Assesment of Resistance Components of Bigleaf Hydrangeas (*Hydrangea macrophylla*) to *Erysiphe polygoni* In Vitro. *Canadian Journal of Plant Pathology*. 31:348-355.
- Sakhanokho, H.F., Rajasekaran, K. 2009. Pollen Biology of Ornamental Ginger (*Hedychium* spp. J. Koenig). *Scientia Horticulturae* pgs 129-135.
- Sakhanokho, H.F., Rajasekaran, K., Kelley, R.Y. 2009. Induced Polyploidy in Diploid Ornamental Ginger (*Hedychium muluense*) Using Colchicine and Oryzalin. *HortScience*. 44(7):1809-1814.
- Smith, B.J. 2009. Influence of Nitrogen, Phosphorus, and Potassium on the Severity of Strawberry Anthracnose Crown Rot. *Acta Horticulturae*. (ISHS) 842:235-238.
- Marshall, D.A., Spiers, J.M. 2009. Establishment of Rabbiteye Blueberries in Poultry Litter. *Acta Horticulturae*. 810:723-728.
- Sampson, B.J., Cane, J.H., Stringer, S.J., Spiers, J.M., Kirker, G.T. 2009. Biology and Management Potential for Three Orchard Bee Species (Hymenoptera: megachilidae): *Osmia Ribifloris* Cockerell, *O. Lignaria* (say) and *O. Chalybea* Smith with Emphasis on the Former. *Acta Horticulturae* (ISHS) 810:549-556.
- Stringer, S.J., Perkins Veazie, P.M., Marshall, D.A. 2009. Nutraceutical Compound Concentrations of Muscadine (*Vitis rotundifolia* Michx) Grape Cultivars and Breeding Lines. *Acta Horticulturae* 841 p. 553-556.
- Stringer, S.J., Spiers, J.M., Draper, A., Marshall, D.A. 2009. Potential New Blueberry Cultivars for the Gulf Coast Region of the U.S. *Acta Horticulturae*. 810:87-92.
- Marshall, D.A., Spiers, J.M. 2008. Establishment of Rabbiteye Blueberries in Poultry Litter. *Acta Horticulturae* (ISHS). 810:723-728.
- Marshall, D.A., Spiers, J.M., Stringer, S.J. 2008. Blueberry Splitting Tendencies as Predicted by Fruit Firmness. *HortScience* 43(2):567-570.
- Smith, B.J. 2008. Cultural Practices and Chemical Treatments Affect Phytophthora Root Rot Severity of Blueberries Grown in Southern Mississippi. *International Journal of Fruit Science* 8:173-181.
- Stringer, S.J., Marshall, D.A., Sampson, B.J., Spiers, J.M. 2008. Performance of Muscadine Grape Cultivars in South Mississippi. *HortTechnology* 18:549-745.
- Mmbaga, M.T., Oliver, J.B. 2007. Effects of Biopesticides on Foliar Diseases and Japanese Beetle (*Popillia japonica*) Adults in Roses (*Rosa* spp.), Oakleaf Hydrangea (*Hydrangea quercifolia*), and Crapemyrtle (*Lagerstroemia indica*). *Journal of Arboriculture*. 33:210-219.
- Chang, P., and Smith, B. J. Response of 'Treasure' and 22 Florida strawberry genotypes to the anthracnose pathogens *Colletotrichum acutatum*, *C. fragariae*, and *C. gloeosporioides*. Pages 69-73. In: Takeda, F., D. T. Handley, and E. B. Poling (ed.). Proc. 2007 N. American Strawberry Symposium. North American Strawberry Growers Association, Kemptville, ON Canada. p. 69-73. 2007.
- Smith, B. J. Developmental Stage and Temperature Affect Strawberry Flower and Fruit Susceptibility to Anthracnose. Pages 55-57. In: Takeda, F., D. T. Handley, and E. B. Poling (ed.). Proc. 2007 North American Strawberry Symposium. North American Strawberry Growers Association, Kemptville, ON Canada. 2007.
- Sampson, B.J., Rinehart, T.A., Liburd, O.E., Stringer, S.J., Spiers, J.M. 2006. Biology of Parasitoids (Hymenoptera) Attacking *Dasineura oxycoccana* and *Prodiplosis vaccinii* (Diptera: Cecidomyiidae) in Cultivated Blueberries. *Annals of the Entomological Society of America*. 99(1):113-120.

6406-21000-011-00D

ENHANCING CORN WITH RESISTANCE TO AFLATOXIN CONTAMINATION AND INSECT DAMAGE – William Williams (P), Marilyn Warburton, William Henry, and Gary Windham; Mississippi State, Mississippi

- Gill, T., Sandoya, G., Williams, W.P., Luthe, D.S. 2011. Belowground resistance to western corn rootworm in Lepidopteran-resistant maize genotypes. *Journal of Economic Entomology*. 104:299-307.
- Ankala, A., Bolla, B.K., Shivaji, R., Williams, W.P., Wilkinson, J. 2011. Comparative analysis of the performance of *Aspergillus flavus* on resistant and susceptible maize genotypes during infection. *Fungal Ecology*. 4:32-41.
- Pechanova, O., Techan, T., Williams, W.P., Luthe, D.S. 2011. Proteomic analysis of the maize rachis: Potential roles of constitutive and induced proteins in resistance to *Aspergillus flavus* and aflatoxin. *Proteomics*. 11:114-127.
- Williams, W.P., Ozkan, S., Ankala, A., Windham, G.L. 2011. Ear rot, aflatoxin accumulation, and fungal biomass in maize after inoculation with *Aspergillus flavus*. *Field Crops Research*. 120:196-200.
- Henry, W.B., Krakowsky, M.D., Windham, G.L., Williams, W.P., Scully, B.T., Rowe, D.E., Hawkins, L.K. 2010. Comparison of the side-needle and knife techniques for inducing *Aspergillus flavus* infection and aflatoxin accumulation in corn hybrids. *Toxin Reviews*. 29:123-129.
- Fu, Z.Y., Yan, J.B., Zheng, Y.P., Warburton, M.L., Crouch, J.H., Li, J.S. 2010. Nucleotide Diversity and Molecular Evolution of the PSY1 Gene in Zea mays Compared to some Other Grass Species. *Theoretical and Applied Genetics*. 120:709-720.
- Jianbing, Y., Xiaohong, Y., Shah, T., Sanchez-Viellda, H., Li, J., Warburton, M.L., Zhou, Y., Jonathan, J., Xu, Y. 2010. High-throughput SNP Genotyping with the GoldenGate Assay in Maize. *Molecular Breeding*. 25:441-451.
- Kelley, R.Y., Gresham, C., Harper, J., Bridges, S., Warburton, M.L., Hawkins, L.K., Pechanova, O., Peethambaran, B., Pechan, T., Luthe, D., Mylroie, J.E., Ankala, A., Ozkan, S., Henry, W.B., Williams, W.P. 2010. Integrated database for identifying candidate genes for *Aspergillus flavus* resistance in maize. *BMC Bioinformatics*. 11(Suppl 6):S25.
- Li, Q., Li, L., Yang, X., Warburton, M.L., Bai, G., Dai, J., Li, J., Yan, J. 2010. Relationship, Evolutionary Fate and Function of Two Maize Co-orthologs of Rice GW2 Associated with Kernel Size and Weight. *Biomed Central (BMC) Plant Biology*. 10:143-157.
- Li, Q., Yang, X., Bai, G., Warburton, M.L., Mahuku, G., Gore, M., Dai, J., Li, J. 2010. Cloning and Characterization of a Putative GS3 Ortholog Involved in Maize Kernel Development. *Theoretical and Applied Genetics*. 120:753-763.
- Pechanova, O., Pechan, T., Ozkan, S., Adams, J.P., McCarthy, F.M., Williams, W.P., Luthe, D.S. 2010. Proteome Profile of the Developing Maize (*Zea mays* L.) Rachis. *Proteomics*. 10:3051-3055.
- Prasanna, B.M., Pixley, K., Warburton, M.L., Xie, C. 2010. Molecular marker-assisted breeding for maize improvement in Asia. *Molecular Breeding*. 26:339-356.
- Setter, T.L., Yan, J., Warburton, M.L., Ribaut, J., Xu, Y., Sawkins, M., Buckler, E.S., Zhang, Z., Gore, M.A. 2010. Genetic association mapping identifies single nucleotide polymorphisms in genes that affect abscisic acid levels in maize floral tissues during drought. *Journal of Experimental Botany*. 62:701-716.
- Shivaji, R., Camas, A., Ankala, A., Engelberth, J., Tumlinson, J.H., Williams, W.P., Wilkinson, J.R., Luthe, D.S. 2010. Plants on Constant Alert: Elevated Levels of Jasmonic Acid and Jasmonate-Induced Transcripts in Caterpillar Resistant Maize. *Journal of Chemical Ecology*. 36:179-191.
- Warburton, M.L., Setimela, P., Franco, J., Cordova, H., Pixley, K., Banziger, M., Dreisigacker, S., Bedoya, C., Macrobert, J. 2010. Toward Cost-Effective Fingerprinting Methodology to Distinguish Maize Open-Pollinated Varieties. *Crop Science*. 50:467-477.
- Williams, W.P., Windham, G.L., Krakowsky, M.D., Scully, B.T., Ni, X. 2010. Aflatoxin accumulation in BT and non-BT maize testcrosses. *Journal of Crop Improvement*. 24:392-399.
- Windham, G.L., Hawkins, L.K., Williams, W.P. 2010. Aflatoxin Accumulation and Kernel Infection of Maize Hybrids Inoculated with *Aspergillus flavus* and *A. parasiticus*. *World Mycotoxin Journal*. 3:89-93.
- Yan, J., Kandianis, C.B., Haries, C.E., Bai, L., Kim, E., Yang, X., Skinner, D., Fu, Z., Mitchell, S., Li, Q., Fernandez, M., Zaharieva, M., Babu, R., Fu, Y., Palacios, N., Li, J., DellaPenna, D., Brutnell, T., Buckler, E.S., Warburton, M.L., Rocheford, T. 2010. Rare Genetic Variation at Zea mays crtRB1 Increases B-carotene in Maize Grain. *Nature Genetics*. 42:322-329.
- Yang, X., Yan, J., Shah, T., Warburton, M.L., Li, Q., Li, L., Gao, Y., Chai, Y., Fu, Z., Zhou, Y., Xu, S., Bai, G., Yijiang, M., Zheng, Y., Li, J. 2010. Genetic Analysis and Characterization of a New Maize Association Mapping Panel for Quantitative Trait Loci Dissection. *Theoretical and Applied Genetics*. 121:417-431.
- Ankala, A., Luthe, D.S., Williams, W.P., Wilkinson, J.R. 2009. Integration of Ethylene and Jasmonic Acid Signaling Pathways in the Expression of Novel Maize Defense Protein Mir1-CP. *Molecular Plant-Microbe Interactions*. 22:1555-1564.
- Henry, W.B., Williams, W.P., Windham, G.L., Hawkins, L.K. 2009. Evaluation of Maize Inbred Lines for Resistance to *Aspergillus* and *Fusarium* Ear Rot and Mycotoxin Accumulation. *Agronomy Journal*. 101:1219-1226.

- Jianbing, Y., Shah, T., Warburton, M.L., Buckler IV, E.S., McMullen, M.D., Crouch, J. 2009. Genetic Characterization and Linkage Disequilibrium Estimation of a Global Maize Collection Using SNP Markers. *PLoS One*. 4:1-14.
- Mideros, S.X., Windham, G.L., Williams, W.P., Nelson, R.J. 2009. *Aspergillus flavus* Biomass in Maize Estimated by Quantitative Real-Time Polymerase Chain Reaction is Strongly Correlated with Aflatoxin Concentration. *Plant Disease*. 93:1163-1170.
- Warburton, M.L., Brooks, T.D., Krakowsky, M.D., Shan, X., Windham, G.L., Williams, W.P. 2009. Identification and mapping of new sources of resistance to aflatoxin accumulation in maize. *Crop Science*. 49:1403-1408.
- Williams, W.P., Windham, G.L. 2009. Diallel Analysis of Fumonisin Accumulation in Maize. *Field Crops Research*. 114:324-326.
- Matthews Jr., G.A., Williams, W.P., Daves, C.A. 2009. Diallel Analysis of Corn Earworm (Lepidoptera: Noctuidae) Resistance in Maize. *Journal of Agricultural and Urban Entomology*. 24:59-66.
- Petthambaran, B., Hawkins, L.K., Windham, G.L., Williams, W.P., Luthe, D.S. 2009. Anti-fungal Activity of Maize Silk Proteins and Role of Chitinases in *Aspergillus flavus* Resistance. *Journal of Toxicology Toxins Reviews*. 29:27-39.
- Mohan, S., Ma, P., Williams, W.P., Luthe, D.L. 2008. A naturally occurring plant cysteine protease possesses remarkable toxicity against insect pests and synergizes *Bacillus thuringiensis*. *PLoS One*. Available: <http://www.plosone.org/article/info%3Adoi%2F10.1371%2Fjournal.pone.0001786>.
- Williams, W.P., Buckley, P.M. 2008. Fall Armyworm (Lepidoptera: Noctuidae) and Southwestern Corn Borer (Lepidoptera: Noctuidae) Leaf Feeding Damage and Its Effect on Larval Growth on Diets Prepared from Lyophilized Corn. *Journal of Agricultural and Urban Entomology*. 25:1-11.
- Williams, W.P., Krakowsky, M.D., Windham, G.L., Balint-Kurti, P.J., Hawkins, L.K., Henry, W.B. 2008. Identifying maize germplasm with resistance to aflatoxin accumulation. *Toxin Reviews*. 27:319-345.
- Alwala, S., Kimbeng, C.A., Williams, W.P., Kang, M.S. 2008. Molecular markers associated with resistance to *Aspergillus flavus* in maize: QTL and discriminant analyses. *Journal of New Seeds*. 9:1-18.
- Bridges, M., Henry, W.B., Shaner, D.L., Khosla, R., Westra, P., Reich, R. 2008. Spatial variability of atrazine and metolachlor dissipation on dryland no-tillage crop fields in Colorado. *Journal of Environmental Quality*. 37:1-9.
- Gaines, T.A., Henry, W.B., Byrne, P.F., Westra, P., Nissen, S.J., Shaner, D.L. 2008. Jointed goatgrass (*Aegilops cylindrica*) by imidazolinone-resistant wheat hybridization under field conditions. *Weed Science*. 56:32-36.
- Hawkins, L.K., Windham, G.L., Williams, W.P. 2008. Occurrence of aflatoxin in three maize (*Zea mays* L.) hybrids over 5 years in Northern Mississippi. *Mycopathologia*. 165:165-171.
- Henry, W.B., Nielsen, D.C., Vigil, M.F., Calderon, F.J., West, M.S. 2008. Proso millet yield and residue mass following direct harvest with a stripper-header. *Agronomy Journal*. 100:580-584.
- Williams, W.P., Windham, G.L., Buckley, P.M. 2008. Diallel analysis of aflatoxin accumulation in maize. *Crop Science*. 48:134-138.
- Windham, G.L., Williams, W.P., Hawkins, L.K., Brooks, T.D. 2008. Effect of *Aspergillus flavus* inoculation methods and environmental conditions on aflatoxin accumulation in corn hybrids. *Toxin Reviews*. 27:1-20.
- Bridges, S.M., Magee, G.B., Wang, N., Williams, W.P., Burgess, S., Nanduri, B. 2007. ProtQuant: A tool for the label-free quantification of MudPIT proteomics data. *BMC Bioinformatics*. 8(Suppl 7):S24:1-9.
- Brooks, T.D., Bushman, B.S., Williams, W.P., McMullen, M.D., Buckley, P.M. 2007. Genetic basis of resistance to fall armyworm (Lepidoptera: Noctuidae) and southwestern corn borer (Lepidoptera: Crambidae) leaf-feeding damage in maize. *Journal of Economic Entomology*. 100:1470-1475.
- Daves, C.A., Williams, W.P., Davis, F.M. 2007. Impact of plant resistance on southwestern corn borer (Lepidoptera: Crambidae) biology and plant damage. *Journal of Economic Entomology*. 100:969-975.
- Daves, C.A., Williams, W.P., Davis, F.M., Baker, G.T., Ma, P.W.K., Monroe, W.A., Mohan, S. 2007. Plant resistance and its effect on the peritrophic membrane of southwestern corn borer (Lepidoptera: Crambidae) larvae. *Journal of Economic Entomology*. 100:976-983.
- Gaines, T., Preston, C., Byrne, P., Henry, W.B., Westra, P. 2007. Adventitious presence of herbicide resistant wheat in certified and farm-saved seed lots. *Crop Science*. 47:749-754.
- Gaines, T.A., Byrne, P.F., Westra, P., Nissen, S.J., Henry, W.B., Shaner, D.L., Chapman, P.L. 2007. An empirically derived model of field-scale gene flow in winter wheat. *Crop Science*. 47:2308-2316.
- Lopez, L., Camas, A., Sshivaji, R., Ankala, A., Williams, W.P., Luthe, D.S. 2007. Mirl-CP, a novel defense proteinase accumulates in maize vascular tissues in response to herbivory. *Planta*. 226:517-527.
- Magbanua, Z.V., Demoraes, C.M., Brooks, T.D., Williams, W.P., Luthe, D.S. 2007. Is catalase activity one of the factors associated with maize resistance to *Aspergillus flavus*. *Molecular Plant-Microbe Interactions*. 20:697-706.
- McCarthy, F.M., Bridges, S.M., Wang, N., Magee, G.B., Williams, W.P., Luthe, D.S., Burgess, S.C. 2007. AgBase: a unified resource for functional analysis in agriculture. *Nucleic Acids Research*. 35:D599-D603.
- Williams, W.P., Buckley, P.M., Daves, C.A. 2007. Identifying resistance in corn to southwestern corn borer (Lepidoptera: Crambidae), fall armyworm (Lepidoptera: Noctuidae), and corn earworm (Lepidoptera: Noctuidae). *Journal of Agricultural and Urban Entomology*. 23:87-95.

- Williams, W.P., Windham, G.L., Buckley, P.M., Daves, C.A. 2007. Aflatoxin accumulation in corn hybrids infested at different growth stages with southwestern corn borer (Lepidoptera: Crambidae). *Journal of Agricultural and Urban Entomology*. 23:97-103.
- Windham, G.L., Williams, W.P. 2007. Systemic infection of stalks and ears of corn hybrids by *Aspergillus parasiticus*. *Mycopathologia*. 164:249-254.
- McCarthy, F.M., Wang, N., Magee, G.B., Nanduri, B., Lawrence, M.L., Camon, E.B., Barell, D.G., Hill, D.P., Dolan, M.E., Williams, W.P., Luthe, D.S., Bridges, S.M., Burgess, S.C. 2006. AgBase: a functional genomics resource for agriculture. *Biomed Central (BMC) Genomics*. 7:229.

6406-21000-012-00D

GERMPLASM ENHANCEMENT AND GENETIC IMPROVEMENT OF COTTON – Johnie Jenkins (P), Sukumar Saha, Jack McCarthy Jr., and Martin Wubben; Mississippi State, Mississippi

- Gutierrez, O., Robinson, A.F., Jenkins, J.N., McCarty Jr., J.C., Wubben, M., Callahan, F.E., Nichols, R.L. 2011. Identification of QTL regions and SSR markers associated with resistance to reniform nematode in *Gossypium barbadense* L. accession GB713. *Theoretical and Applied Genetics*. 122:271-280.
- Saha, S., Wu, J., Jenkins, J.N., McCarty, J.C., Hayes, R.W., Stelly, D.M. 2011. Delineation of interspecific epistasis on fiber quality traits in *Gossypium hirsutum* by ADAA analysis of intermated *G. barbadense* chromosome substitution lines. *Theoretical and Applied Genetics*. 122:1351-1361.
- Wu, J., Jenkins, J.N., McCarty Jr., J.C., Saha, S. 2010. Genetic effects of individual chromosomes in cotton cultivars detected by using chromosome substitution lines as genetic probes. *Genetica*. 138:1171-1179.
- An, C., Jenkins, J.N., McCarty Jr., J.C., Saha, S. 2010. Atypical Ligon Lintless-2 Phenotype in Cotton. *Journal of Cotton Science*. 14:13-16.
- Buriev, Z.T., Saha, S., Abdurakhmonov, I.Y., Jenkins, J.N., Abdurakarimov, A., Scheffler, B.E., Stelly, D.M. 2010. Clustering, haplotype diversity and locations of MIC-3: a unique root-specific defense-related gene family in upland cotton (*Gossypium hirsutum* L.). *Theoretical and Applied Genetics*. 120:587-606.
- Gutierrez, O.A., Jenkins, J.N., McCarty Jr., J.C., Wubben, M., Hayes, R.W., Callahan, F.E. 2010. SSR markers closely associated with genes for resistance to root-knot nematode on chromosomes 11 and 14 of upland cotton. *Theoretical and Applied Genetics*. 121:1323-1337.
- Ho, M., Saha, S., Jenkins, J.N., Ma, D. 2010. Characterization and Promoter Analysis of a Cotton Ring-Type Ubiquitin Ligase (E3) Gene. *Molecular Biotechnology*. 46:140-148.
- Saha, S., Wu, J., Jenkins, J.N., McCarty Jr., J.C., Hayes, R.W., Stelly, D. 2010. Genetic Dissection of Chromosome Substitution Lines of Cotton to Discover Novel *Gossypium barbadense* L. Alleles for Improvement of Agronomic Traits. *Journal of Theoretical and Applied Genetics*. 120:1193-1205.
- Wu, J., McCarty Jr., J.C., Jenkins, J.N., Meredith Jr., W.R. 2010. Breeding potential of introgressions into cotton: genetic effects and heterosis. *Plant Breeding*. 129:526-532.
- Wu, J., McCarty Jr., J.C., Jenkins, J.N. 2010. Cotton Chromosome Substitution Lines Crossed with Cultivars: Genetic Model Evaluation and Seed Trait Analyses. *Theoretical and Applied Genetics*. 120:1473-1483.
- Wubben, M., Callahan, F.E., Scheffler, B.E. 2010. Transcript Analysis of Parasitic Females of the Sedentary Semi-Endoparasitic Nematode *Rotylenchulus reniformis*. *Molecular and Biochemical Parasitology*. 172:31-40.
- An, C., Jenkins, J.N., Wu, J., Guo, Y., McCarty Jr., J.C. 2010. Use of Fiber and Fuzz Mutants to Detect QTL for Yield Components, Seed, and Fiber Traits of Upland Cotton. *Euphytica*. 172:21-34.
- Jenkins, J.N., McCarty Jr., J.C., Wu, J., Gutierrez, O. 2009. Genetic variance components and genetic effects among eleven diverse upland cotton lines and their F2 hybrids. *Euphytica*. 167:397-408.
- Myers, G.M., Jiang, B., Akash, M.W., Badigannavar, A., Saha, S. 2009. Chromosomal assignment of ALFP markers in upland cotton (*Gossypium hirsutum* L.). *Euphytica*. 165:391-399.
- Wu, J., Jenkins, J.N., McCarty Jr., J.C., Thaxton, P. 2009. Seed trait associations with *Gossypium barbadense* L. chromosomes/arms in a *G. hirsutum* L. background. *Euphytica*. 167:371-380.
- Wubben, M., Callahan, F.E., Triplett, B.A., Jenkins, J.N. 2009. Phenotypic and Molecular Evaluation of Cotton Hairy Roots as a Model System for Studying Nematode Interactions. *Plant Cell Reports*. 28:1399-1409.
- Abdurakhmonov, I.Y., Saha, S., Jenkins, J.N., Buriev, Z.T., Shermatov, S.E., Scheffler, B.E., Pepper, A.E., Yu, J., Kohel, R.J., Abdurakarimov, A. 2009. Linkage disequilibrium based association mapping of fiber quality traits in *G. hirsutum* L. variety germplasm. *Genetica*. 136:401-417.
- Guo, Y., McCarty Jr., J.C., Jenkins, J.N., An, C., Saha, S. 2009. Genetic detection of node of first fruiting branch in crosses of a cultivar with two exotic accessions of upland cotton. *Euphytica*. 166:317-329.
- Gutierrez, O.A., Wubben, M., Howard, M., Roberts, B., Hanlon, E., Wilkinson, J.R. 2009. Role of phytohormones ethylene and auxin in plant-nematode interactions: mini-review. *Russian Journal of Plant Physiology*. 56:1-5.
- Wu, J., Gutierrez, O. A., Jenkins, J.N., McCarty Jr., J.C., Zhu, J. 2009. Quantitative analysis and QTL mapping for agronomic and fiber traits in an RI population of Upland cotton. *Euphytica*. 165:231-245.
- Wu, J., McCarty Jr., J.C., Saha, S., Jenkins, J.N., Hayes, R.W. 2009. Genetic Changes in Plant Growth and their Associations with Chromosomes from *Gossypium hirsutum* L. in *G. hirsutum* L. *Genetica*. 137:56-66.

- An, C., Saha, S., Jenkins, J.N., Ma, D., Scheffler, B.E., Kohel, R.J., Yu, J., Stelly, D.M. 2008. Cotton (*Gossypium* spp.) R2R3-MYB transcription factors SNP identification, phylogenomic characterization, chromosome localization and linkage mapping. *Theoretical and Applied Genetics*. 161:1015-1026.
- Guo, Y., McCarty Jr., J.C., Jenkins, J.N., Saha, S. 2008. QTLs for node of first fruiting branch in a cross of an upland cotton, *Gossypium hirsutum* L., cultivar with primitive accession Texas 701. *Euphytica*. 163:113-122.
- Hsu, C., An, C., Saha, S., Ma, D., Jenkins, J.N., Scheffler, B.E., Stelly, D. 2008. Molecular and SNP characterization of two genome specific transcription factor genes GhMyb8 and GhMyb10 in cotton species. *Euphytica*. 159:259-273.
- Jenkins, J.N., McCarty Jr., J.C., Gutierrez, O.A., Hayes, R.W., Bowman, D.T., Watson, C.E., Jones, D.C. 2008. Registration of RMUP-C5, a random mated population of Upland cotton germplasm. *Journal of Plant Registrations*. 2:239-242.
- McCarty Jr., J.C., Wu, J., Jenkins, J.N. 2008. Genetic association of cotton yield with its component traits in derived primitive accessions crossed by elite Upland cultivars using the conditional ADAA genetic model. *Euphytica*. 161:337-352.
- Saha, S., Jenkins, J.N., Wu, J., McCarty Jr., J.C., Stelly, D.M. 2008. Genetic analysis of agronomic and fiber traits using four interspecific chromosome substitution lines in cotton. *Plant Breeding*. 127:612-618.
- Wu, J., Jenkins, J.N., McCarty Jr., J.C., Saha, S., Percy, R.G. 2008. Genetic association of lint yield with its components in cotton chromosome substitution lines. *Euphytica*. 164:199-207.
- Wu, Z., Soliman, K.M., Bolton, J.J., Saha, S., Jenkins, J.N. 2008. Identification of differentially expressed genes associated with cotton fiber development in a chromosomal substitution line (CS-B22sh). *Genomics*. 8:166-174.
- Wubben, M., Callahan, F.E., Hayes, R.W., Jenkins, J.N. 2008. Molecular characterization and temporal expression analyses indicate that the MIC (Meloidogyne Induced Cotton) gene family represents a novel group of root-specific defense-related genes in upland cotton (*Gossypium hirsutum* L.). *Planta*. 228:111-123.
- Wubben, M., Jing, J., Baum, T.J. 2008. Cyst nematode parasitism of *Arabidopsis thaliana* is inhibited by salicylic acid (SA) and elicits uncoupled SA-independent pathogenesis-related gene expression in roots. *Molecular Plant-Microbe Interactions*. 21:424-432.
- An, C., Saha, S., Jenkins, J.N., Scheffler, B.E., Wilkins, T.A., Stelly, D.M. 2007. Transcriptome profiling, sequence characterization, and SNP-based chromosomal assignment of the EXPANSIN genes in cotton. *Molecular Genetics and Genomics*. 278:539-553.
- Creech, R.G., Jenkins, J.N., McCarty Jr., J.C., Hayes, R.W., Creech, J.B., Haire, D., Cantrell, R. 2007. Registration of MS-01RKN, MS-24RKN, MS-30RKN, MS-33RKN, MS-35RKN and MS-37RKN cotton germplasm lines with resistance to root-knot nematode. *Journal of Plant Registrations*. 1:147-148.
- Jenkins, J.N., McCarty Jr., J.C., Wu, J., Saha, S., Gutierrez, O.A., Hayes, R.W., Stelly, D.M. 2007. Genetic effects of thirteen *Gossypium barbadense* L. chromosome substitution lines in topcrosses with upland cotton cultivars: II. Fiber quality traits. *Crop Science*. 47:561-572.
- McCarty Jr., J.C., Wu, J., Jenkins, J.N. 2007. Use of primitive derived cotton accessions for agronomic and fiber traits improvement: Variance components and genetic effects. *Crop Science*. 47:100-110.
- Ynturi, P., Jenkins, J.N., McCarty Jr., J.C., Gutierrez, O.A., Saha, S. 2006. Association of root-knot nematode resistance genes with simple sequence repeat markers on two chromosomes in cotton. *Crop Science*. 46:2670-2674.

6408-21410-008-00D

DISCOVERY AND DEVELOPMENT OF NATURAL PRODUCT FOR PHARMACEUTICAL AND AGRICHEMICAL APPLICATIONS – Stephen Duke (P); University, Mississippi

- Ding, Y., Li, X., Ferreira, D. 2010. 4-Arylflavan-3-ols as Proanthocyanidin Models: Absolute Configuration via Density Functional Calculation of Electronic Circular Dichroism. *Journal of Natural Products*. 73:435-440.
- Heratch, W., Khan, I.A. 2010. Microbial Metabolism. Part 11. Metabolites of Flutamide. *Chemical and Pharmaceutical Bulletin*. 58(4):562-564.
- Wang, W., Ali, Z., Li, X., Khan, I.A. 2010. A New ent-Labdane Diterpene Glycoside from the Leaves of *Casearia sylvestris*. *Natural Product Communications*. 5(5):771-774.
- Wang, W., Ali, Z., Shen, Y., Li, X., Khan, I.A. 2010. Ursane Triterpenoids from the Bark of *Terminalia arjuna*. *Fitoterapia*. 81:480-484.
- Bialonska, D., Kasimsetty, S.G., Khan, S.I., Ferreira, D. 2009. Urolithins, Intestinal Microbial Metabolites of Pomegranate Ellagitannins, Exhibit Potent Antioxidant Activity in Cell-Based Assay. *Journal of Agricultural and Food Chemistry*. 57(21):10181-10186.
- Xu, W., Jacob, M.R., Agarwal, A.K., Clark, A.M., Liang, Z., Li, X. 2009. ent-Kaurane Glycosides from *Tricalysia okelensis*. *Chemical and Pharmaceutical Bulletin*. 58:261-264.
- Wang, W., Ali, Z., Li, X., Khan, I.A. 2009. Neolignans from the Leaves of *Casearia sylvestris* Swartz. *Helvetica Chimica Acta*. 93(1):139-146.

6410-21000-014-00D

GENETIC IMPROVEMENT OF SUGARCANE BY CONVENTIONAL AND MOLECULAR APPROACHES – Thomas Tew (P), Anna Hale, Edward Richard Jr, Yong Bao Pan, and William White; Houma, Louisiana

- Hodnett, G.L., Hale, A.L., Packer, D.J., Stelly, D., Da Silva, J., Rooney, W.L. 2010. Elimination of a reproductive barrier facilitates intergeneric hybridization of *Sorghum bicolor* and *Saccharum*. *Crop Science*. 50:1188-1195.
- Liang, J., Pan, Y., Li, Y.R., Fang, F.X., You, J.H. 2010. Assessment of Genetic Diversity in *Saccharum* Using SSR Markers and Capillary Electrophoresis. *Guihaia*. 30(1):106-111.
- Pan, Y.-B. 2010. Databasing molecular identities of sugarcane (*Saccharum* spp.) clones constructed with microsatellite (SSR) DNA markers. *American Journal of Plant Sciences*. 1(2):87-94.
- Tew, T.L., Pan, Y. 2010. Microsatellite (SSR) Marker-Based Paternity Analysis of a Seven-Parent Polycross in Sugarcane. *Crop Science*. 50:1401-1408.
- Tew, T.L., Dufrene Jr, E.O., Garrison, D.D., White, W.H., Grisham, M.P., Pan, Y., Richard Jr, E.P., Legendre, B.L., Miller, J.D. 2009. Registration of HoCP 00-950" Sugarcane. *Journal of Plant Registrations*. 39(1):42-50.
- Chen, P.H., Pan, Y., Chen, R.R., Xu, L.P., Chen, Y.Q. 2009. SSR marker-based analysis of genetic relatedness among sugarcane cultivars (*Saccharum* spp. hybrids) from breeding programs in China and other countries. *Sugar Tech*. 11(4):347-354.
- Alvi, A.K., Iqbal, J., Shah, A.H., Pan, Y. 2008. DNA Based Genetic Variation for Red Rot Resistance in Sugarcane. *Pakistan Journal of Botany*. 40(4):1419-1425.
- Gao, S.J., Pan, Y., Chen, R.K., Zhang, H., Xu, L. 2008. Quick detection of *Leifsonia xyli* subsp. *xyli* by PCR and nucleotide sequence analysis of PCR amplicons from Chinese *Leifsonia xyli* subsp. *xyli* isolates. *Sugar Tech*. 10(4):334-340.
- Tew, T.L., Cobill, R.M., Richard Jr, E.P. 2008. Evaluation of Sweet Sorghum and Sorghum x Sudangrass Hybrids as Feedstocks for Ethanol Production. *BioEnergy Research*. 1:147-152.
- Chen, P., Pan, Y., Chen, R. 2008. High-Throughput Procedure for Single Pollen Grain Collection and Polymerase Chain Reaction in Plants. *Journal of Integrative Plant Biology*. 50:375-383.
- Pan, Y.-B., Scheffler, B.S., Richard, Jr., E.P. 2007. High-throughput genotyping of commercial sugarcane clones with microsatellite (SSR) DNA markers. *Sugar Tech*. 9:176-181.
- Pan, Y.-B. 2006. Highly polymorphic microsatellite DNA markers for sugarcane germplasm evaluation and variety identity testing. *Sugar Tech*. 8(4):246-256.

6435-21000-016-00D

MOLECULAR ANALYSIS OF DEVELOPMENT TO IMPROVE COTTON FIBER – David Fang (P) and Hee Kim; New Orleans, Louisiana

- Naoumkina, M.A., Dixon, R. 2011. Characterization of the mannan synthase promoter from guar (*Cyamopsis tetragonoloba*). *Plant Cell and Environment*. DOI 10.1007/s00299-011-1003-4.
- Hinchliffe, D.J., Meredith Jr, W.R., Delhom, C.D., Thibodeaux, D.P., Fang, D.D. 2011. Elevated growing degree days influence transition stage timing during cotton (*Gossypium hirsutum* L.) fiber development and result in increased fiber strength. *Crop Science*. 51:1683-1692, DOI 10.2135/cropsci2010.10.0569.
- Kim, H.J., Murai, N., Fang, D.D., Triplett, B.A. 2010. Functional analysis of *Gossypium hirsutum* cellulose synthase catalytic subunit 4 promoter in transgenic *Arabidopsis* and cotton tissues. *Plant Science*. 180(2011):323-332.
- Hinchliffe, D.J., Meredith Jr, W.R., Yeater, K.M., Kim, H. 2010. Near-Isogenic Cotton Germplasm Lines that Differ in Fiber-Bundle Strength have Temporal Differences in Fiber Gene Expression Pattern as Revealed by Comparative High-Throughput Profiling. *Theoretical and Applied Genetics*. 120:1347-1366.
- Xiao, J., Fang, D.D., Bhatti, M., Hendrix, B., Cantrell, R.G. 2009. A SNP Haplotype Associated with a gene resistant to *Xanthomonas axonopodis* pv. *malvacearum* in Upland Cotton (*Gossypium hirsutum* L.). *Molecular Breeding*. 25(4):593-602.
- Fang, D.D., Xiao, J., Canci, P.C., Cantrell, R.G. 2009. A New SNP Haplotype associated with blue disease resistance gene in cotton (*Gossypium hirsutum* L.). *Journal of Theoretical and Applied Genetics*. 120:943-953.

6602-21000-021-00D

GENETIC AND GENOMIC APPROACHES TO IMPROVE PEANUT AND CORN RESISTANCE TO DISEASE AND AFLATOXIN CONTAMINATION – Baozhu Guo (P) and Brian Scully; Tifton, Georgia

- Guo, B., Krakowsky, M.D., Ni, X., Scully, B.T., Lee, R., Coy, A., Widstrom, N.W. 2011. Registration of Maize Inbred Line GT603. *Journal of Plant Registrations*. 5(2):211-214.

- Baozhu, G., Butron, A., Scully, B.T. 2010. Maize silk antibiotic polyphenol compounds and molecular genetic improvement of resistance to corn earworm (*Helicoverpa zea* Boddie) in sh2 sweet corn. *International Journal of Plant Biology*. 1:13-18.
- Dang, P.M., Scully, B.T., Lamb, M.C., Guo, B.Z. 2010. Analysis and RT-PCR identification of viral sequences in peanut (*Arachis hypogaea* L.) expressed sequence tags from different peanut tissues. *Plant Pathology Journal*. 9:14-22.
- Fountain, J.C., Chen, Z., Scully, B.T., Kemerait, R., Lee, R., Guo, B. 2010. Pathogenesis-related gene expressions in different maize genotypes under drought stressed conditions.. *African Journal of Plant Science*. Vol. 4(11), pp. 433-440.
- Hong, Y., Chen, X., Liang, X., Liu, H., Zhou, G., Li, S., Wen, S., Holbrook, C., Guo, B. 2010. A SSR-based composite genetic linkage map for the cultivated peanut (*Arachis hypogaea* L.) genome. *Biomed Central (BMC) Plant Biology*. 10:17.
- Jiang, T., Wu, L., Gao, F., Guo, B. 2010. Genetic Linkage Maps of White Birchs (*Betula platyphylla* Suk. and *B. pendula* Roth) based on RAPD and AFLP markers. *Molecular Breeding*. (2011)27:347-356.
- Luo, M., Liu, J., Lee, D., Scully, B.T., Guo, B. 2010. Monitoring the Expression of Maize Genes in Developing Kernels under Drought Stress using Oligo-microarray. *Journal of Integrative Plant Biology*. 52(12):1059-1074.
- Guo, B., Chen, X., Hong, Y., Liang, X., Dang, P., Brennenman, T., Holbrook Jr., C., Culbreath, A. 2009. Analysis of Gene Expression Profiles in Leaf Tissues of Cultivated Peanuts and Development of EST-SSR Markers and Gene Discovery. *International Journal of Plant Genomics*. Article ID 715605:14pp.
- Liang, X., Chen, X., Hong, Y., Liu, H., Zhou, G., Li, S., Guo, B. 2009. Utility of EST-derived SSR in cultivated peanut (*Arachis hypogaea* L.) and *Arachis* wild species. *BMC Plant Biology*. 9:35.
- Guo, B., Yu, J., Holbrook Jr., C., Cleveland, T., Niernan, W.C., Scully, B. 2009. Strategies in Prevention of Preharvest Aflatoxin Contamination in Peanuts: Aflatoxin Biosynthesis, Genetics and Genomics. *Peanut Science*. 36:11-20.
- Jiang, H.F., Ren, X.P., Liao, B.S., Huang, J.Q., Lei, Y., Chen, B.Y., Guo, B., Holbrook, C.C., Upadhyaya, H.D. 2008. Peanut core collection established in China and compared with ICRISAT mini core collection. *Acta Agronomica Sinica*. 34(1):25-30.
- Guo, B., Chen, X., Dang, P., Scully, B.T., Liang, X., Holbrook, C.C., Yu, J., Culbreath, A.K. 2008. Peanut gene expression profiling in developing seeds at different reproduction stages during *Aspergillus parasiticus* infection. *BMC Developmental Biology*. 8:12
- Guo, B., Chen, Z., Lee R.D., Scully B. 2008. Drought Stress and Preharvest Aflatoxin Contamination in Agricultural Commodity: Genetics, Genomics and Proteomics. *Journal of Integrative Plant Biology*. 50(10):1281-1291.
- Guo, B., Liang, X., Chung, S., Holbrook, C.C., Maleki, S.J. 2008. Proteomic analysis of peanut seed storage proteins and genetic variation in a potential peanut allergen. *Protein & Peptide Letters*. 15:567-577.
- Guo, B.Z., Liang, X., Chung, S.Y., Maleki, S.J. 2008. Proteomic screening points to the potential importance of Ara h 3 basic subunit in allergenicity of peanut. *Inflammation & Allergy - Drug Targets*. 7(3):163-166
- Luo, M., Liu, J., Lee, R.D., Guo, B. 2008. Characterization of gene expression profiles in developing kernels of maize (*Zea mays* L.) inbred Tex6. *Plant Breeding*. 127:569-578.
- Guo, B.Z., Widstrom, N.W., Lee, R.D., Coy, A.E., Lynch, R.E. 2007. Registration of maize germplasm GT601 (AM-1) and GT602 (AM-2). *Journal of Plant Registrations*. 1(2):153-154.
- Liang, X.Q., Luo, M., Holbrook, C.C., Guo, B.Z. 2006. Storage protein profiles in Spanish and runner market type peanuts and potential markers. *BMC Plant Biology*. 6:24.

6602-21000-022-00D

DEVELOPMENT OF IMPROVED PEANUT GERMLASM AND RESISTANCE TO DISEASE AND NEMATODE PESTS – C. Holbrook Jr (P); Tifton, Georgia

- Girdthai, T., Jogloy, S., Kesmala, T., Vorasoot, N., Akkasaeng, C., Wongkaew, S., Holbrook Jr, C.C., Patanothai, A. 2010. Relationship between root characteristics of peanut in hydroponics and pot studies. *Crop Science*. 50:159-167.
- Nagy, E.D., Chu, Y., Guo, Y., Khanal, S., Tang, S., Li, Y., Dong, W.B., Timper, P., Taylor, C., Ozias-Akins, P., Holbrook Jr, C.C., Beilinson, V., Nielsen, N.C., Stalker, H., Knapp, S.J. 2010. Recombination is suppressed in an alien introgression on chromosome 5A of peanut harboring Rma, a dominant root-knot nematode resistance gene. *Molecular Breeding*. 26:357-370.
- Chu, Y., Holbrook Jr, C.C., Ozias-Akins, P. 2009. Two alleles of ahFAD2B control the high oleic acid trait in cultivated peanut. *Crop Science*. 49:2029-2036.
- Dong, W.B., Brennenman, T.B., Holbrook Jr, C.C., Timper, P., Culbreath, A.K. 2009. The interaction between *Meloidogyne arenaria* and *Cylindrocladium parasiticum* in runner peanut. *Plant Pathology* 58:71-79.
- Holbrook Jr, C.C., Guo, B., Wilson, D.M., Timper, P. 2009. The U.S. breeding program to develop peanut with drought tolerance and reduced aflatoxin contamination. *Peanut Science*. 36:50-53.

- Holbrook Jr, C.C., Ozias-Akins, P., Timper, P., Wilson, D.M., Cantonwine, E., Guo, B., Sullivan, D.G., Dong, W. 2009. Research from the Coastal Plain Experiment Station, Tifton, Georgia, to minimize contamination in peanut. *Toxin Reviews*. 27:391-410.
- Pimratch, S., Jogloy, S., Vorasoot, N., Toomsan, B., Kesmala, T., Patanothai, A., Holbrook Jr, C.C. 2009. Heritability of N₂ fixation traits, and phenotypic and genotypic correlations between N₂ fixation traits with drought resistance traits and yield in peanut under different water regimes. *Crop Science* 49:791-800.
- Songsri, P., Jogloy, S., Holbrook Jr, C.C., Kesmala, T., Vorasoot, N., Akkasaeng, C., Patanothai, A. 2009. Association of root, specific leaf area and SPAD chlorophyll meter reading to water use efficiency of peanut under different available soil water. *Agricultural Water Management* 96:790-798.
- Songsri, P., Jogloy, S., Kesmala, T., Vorasoot, N., Akkasaeng, C., Patanothai, A., Holbrook Jr, C.C. 2009. Heritability of drought resistance traits and correlation of drought resistance and agronomic traits in peanut. *Crop Science*. 48:2245-2253.
- Songsri, P., Jogloy, S., Kesmala, T., Vorasoot, N., Akkasaeng, C., Patanothai, A., Holbrook Jr, C.C. 2009. Response of reproductive characters of drought resistant peanut genotypes to drought. *Asian Journal of Plant Sciences*. 7(5):427-439.
- Songsri, P., Jogloy, S., Vorasoot, N., Akkasaeng, C., Patanothai, A., Holbrook Jr, C.C. 2009. Root distribution of drought-resistant peanut genotypes in response to drought. *Journal of Agronomy and Crop Science*. 194:92-103.
- Stalker, H.T., Weissinger, A.K., Milla-Lewis, S., Holbrook Jr, C.C. 2009. Genomics: An evolving science in peanut. *Peanut Science*. 36:2-10.
- Dong, W.B., Brenneman, T.B., Holbrook Jr, C.C., Culbreath, A.K. 2008. Evaluation of resistance to *Cylindrocladium parasiticum* in peanut in the greenhouse and in inoculated or naturally infested fields. *Peanut Science*. 35:139-148.
- Pimratch, S., Jogloy, S., Vorasoot, N., Toomsan, B., Patanothai, A., Holbrook Jr, C.C. 2008. Relationship between biomass production and nitrogen fixation under drought stress conditions in peanut genotypes with different levels of drought resistance. *Journal of Agronomy and Crop Science* 194:15-24.
- Cantonwine, E.G., Culbreath, A.K., Holbrook Jr, C.C., Gorbet, D.W. 2008. Disease Progress of Early Leaf Spot and Components of Resistance to *Cercospora Arachidicola* and *Cercosporidium Personatum* in Runner-type Peanut Cultivars. *Peanut Science* 35:1-10.
- Dong, W.B., Holbrook Jr, C.C., Timper, P., Brenneman, T.B., Chu, Y., Ozias-Akins, P. 2008. Resistance in peanut cultivars and breeding lines to three root-knot nematode species. *Plant Disease* 92:631-638.
- Holbrook Jr, C.C., Culbreath, A.K. 2008. Registration of 'Georganic' Peanut. *Journal of Plant Registrations*. 2:17.
- Holbrook Jr, C.C., Timper, P., Culbreath, A.K. 2008. Registration of Peanut Germplasm Line TifGP-1 with Resistance to the Root-knot Nematode and Tomato Spotted Wilt Virus. *Journal of Plant Registrations* 2:57.
- Holbrook Jr, C.C., Timper, P., Culbreath, A.K., Kvien, C.K. 2008. Registration of 'Tifguard' Peanut. *Journal of Plant Registrations*. 2:92-94.
- Holbrook Jr, C.C., Timper, P., Dong, W., Kvien, C.K., Culbreath, A.K. 2008. Development of near isogenic peanut lines with and without resistance to the peanut root-knot nematode. *Crop Science* 48:194-198.
- Chu, Y., Holbrook Jr, C.C., Timper, P., Ozias-Akins, P. 2007. Development of a PCR-based Molecular Marker to Select for Nematode Resistance in Peanut. *Crop Sci.* 47:841-845.
- Chu, Y., Ramos, L., Holbrook Jr, C.C., Ozias-Akins, P. 2007. Frequency of a loss-of-function mutation in Oleoyl-PC Desaturase (ahFAD2A) in the minicore of the U.S. peanut germplasm collection. *Crop Science* 47:2372-2378.
- Dong, W., Holbrook Jr, C.C., Timper, P., Brenneman, T.B., Mullinix, B.G. 2007. Effects of Inoculum Type, Inoculum Level, Inoculation Date, and Assessment Date on Evaluating Resistance to *Meloidogyne arenaria* in Peanut. *J. of Nematology* 39(2):169-175.

6602-21220-014-00D

GENETIC IMPROVEMENT OF MAIZE AND PEARL MILLET FOR RESISTANCE TO INSECTS AND AFLATOXIN – Jeffrey Wilson (P), C Holbrooke Jr., and Xinzhi Ni; Tifton, Georgia

- Park, Y., Gobel, C., Ni, X., Feussner, I., Kolomiets, M.V. 2010. Comparative molecular and biochemical characterization of segmentally duplicated 9-lipoxygenase genes ZmLOX4 and ZmLOX5 of maize. *Planta*. 231:1425-1437.
- Zhang, C., Luo, X., Ni, X., Zhang, Y., Li, X. 2010. Functional characterization of cis-acting elements mediating flavone-inducible expression of CYP321A1. *Insect Biochemistry and Molecular Biology*.
- Chen, Y., Ni, X., Buntin, G. 2009. Physiological, nutritional and biochemical bases of corn resistance to foliage-feeding fall armyworm (Lepidoptera: noctuidae). *Journal of Chemical Ecology* 35:297-306.
- Chen, Y., Ni, X., Cottrell, T.E., Wood, B.W., Buntin, G. 2009. Changes of oxidase and hydrolase activities in pecan leaves elicited by black pecan aphid (Hemiptera: Aphididae) feeding. *Journal of Economic Entomology* 102:1262-1269.

- Maas, A., Ni, X. 2009. Inheritance of chinch bug resistance in grain pearl millet. *Journal of Semi-Arid Tropical (SAT) Agricultural Research* 7. http://ejournal.icrisat.org/Volume7/Sorghum_Millet/PM703.pdf.
- Macedo, T.B., Peterson, R.D., Weaver, D.K., Ni, X. 2009. Impact of *Duraphis noxia* (Mordvilko) and *Rhopalosiphum padi* (L.) (Hemiptera: Aphididae) on primary physiology of four near-isogenic wheat lines. *Journal of Economic Entomology* 102:412-421.
- Ni, X., Wilson, J.P., Buntin, G. 2009. Differential responses of forage pearl millet genotypes to chinch bug (Heteroptera:Blissidae) feeding. *Journal of Economic Entomology*. 102:1960-1969.
- Ni, X., Da, K., Buntin, G., Brown, S.L. 2008. Physiological basis of fall armyworm (Lepidoptera: Noctuidae) resistance in the seedlings of maize inbred lines with varying levels of silk maysin. *Florida Entomologist* 91:537-545.
- Ni, X., Gunawan, G., Brown, S.L., Sumner, P.E., Buntin, G., Holbrook Jr, C.C., Lee, G., Streett, D.A., Throne, J.E., Campbell, J.F. 2008. Insect-attracting and antimicrobial properties of antifreeze for monitoring storage insect pests and natural enemies in stored corn. *Journal of Economic Entomology*. 101:631-636.
- Ni, X., Krakowsky, M.D., Buntin, G., Rector, B.G., Guo, B., Snook, M.E. 2008. Identification of multiple ear-colonizing insect and disease resistance in CIMMYT maize inbred lines with varying levels of silk maysin. *Journal of Economic Entomology*. 101:1455-1465.
- Ni, X., Wilson, J.P., Rajewski, J.A., Buntin, G.D., Dweikat, I.M. 2007. Field evaluation of pearl millet for chinch bug (heteroptera:blissidae) resistance. *Journal of Entomological Science*. 42:467-480.
- Ni, X., Xu, W., Krakowsky, M.D., Buntin, G.D., Brown, S.L., Lee, R.D., Coy, A.E. 2007. Field Screening of Experimental Corn Hybrids and Inbred Lines for Multiple Ear-Feeding Insect Resistance. *Journal of Economic Entomology*. 100(5):1704-1713.

6604-21000-003-00D

PHYSIOLOGICAL MECHANISMS THAT DETERMINE CROP RESPONSE TO IRRIGATION, DISEASE, AND PRODUCTION PRACTICES – Charles Chen (P), Marshall Lamb, and Phat Dang; Dawson, Georgia

- Rowland, D., Beasley, J., Faircloth, W.H. 2009. Genotypic patterns of peanut (*Arachis hypogaea* L.) phenology and the inability of current irrigation scheduling methods to modify them. *Peanut Science*. v. 36 p 133-137.
- Dang, P.M., Rowland, D., Faircloth, W.H. 2009. Comparison of ELISA and RT-PCR testing for Tomato spotted wilt virus (TSWV) infection in peanut. *Peanut Science*. v 36, p 133-137.

6606-21000-002-00D

STONE FRUIT BREEDING AND DEVELOPMENT – Thomas Beckman (P); Byron, Georgia

- Okie, W.R. and B. Blackburn. 2011. Increasing chilling reduces heat requirement for floral budbreak in peach. *HortScience* 46: 245-252.
- Fan, S., Bielenberg, D., Zhebentyayeva, T., Reighard, G.L., Okie, W.R., Holland, D., Abbott, A. 2010. Mapping quantitative trait loci associated with chilling requirement, heat requirement and bloom date in peach [*Prunus persica* (L.) Batsch]. *New Phytologist*. 185(4):917-930.
- Noratto, G., Cisneros-Zevallos, L., Byrne, D.H., Okie, W.R., Porter, W. 2009. Molecular mechanisms involved in the inhibition of MDA-MB-435 breast cancer cells by phenolic acids from the red-flesh peach BY00P6653. *Acta Horticulturae*. 841:67-72.
- Anderson, P.C., Mizell, III, R.F., Brodbeck, B.V., Beckman, T.G., Krewer, G. 2008. Abundance and consumption rate of glassy-winged sharpshooter (Hemiptera: Cicadellidae) on peaches and plums. *Journal of Entomological Science*. 43(4):394-407.
- Lea, M.A., Ibeh, C., Desbordes, C., Vizzotto, M., Cisneros-Zevallos, L., Byrne, D.H., Okie, W.R., Moyer, M.P. 2008. Inhibition of growth and induction of differentiation of colon cancer cells by peach and plum phenolic compounds. *Anticancer Research*. 28(4B):2067-2076.
- Wells, L., Wood, B.W. 2008. Foliar boron and nickel applications reduce water-stage fruit-split of pecan. *HortScience*. 43(5):1437-1440.
- Beckman, T.G. 2008. Nursery performance of peach seedling rootstocks. *Journal of American Pomological Society*. 62(2):46-51.
- Beckman, T.G., Chaparro, J.X., Sherman, W.B. 2008. 'Sharpe', a clonal plum rootstock for peach. *HortScience*. 43(7):2236-2237.
- Beckman, T.G., Krewer, G.W., Chaparro, J.X., Sherman, W.B. 2008. Potential of non-melting flesh peaches for the early season fresh market. *Journal of American Pomological Society*. 62(2):52-57.
- Krewer, G.W., Beckman, T.G., Chaparro, J.X., Sherman, W.B. 2008. Gulfcrimson Peach. *HortScience*. 43:1596-1597.
- Okie, W.R., Layne, D.R. 2008. 'Early Augustprince' and 'Augustprince' Peaches. *HortScience*. 43(5):1600-1602.

- Okie, W.R., Layne, D.R. 2008. 'Scarletprince' and 'Julyprince' Peaches. *HortScience*. 43(5):1603-1605.
- Okie, W.R., Reighard, G.L., Nyczepir, A.P. 2009. Importance of scion cultivar in peach tree short life. *Journal of American Pomological Society*. 63:58-63.
- Okie, W.R. 2006. Introgression of *Prunus* species in plum. *New York State Fruit Quarterly*. 14(1):29-37.

6607-21000-010-00D

CONSERVATION, CHARACTERIZATION, AND EVALUATION OF CROP GENETIC RESOURCES AND ASSOCIATED INFORMATION – Gary Pederson (P), Ming Wang, John Morris, Roy Pittman, Melanie Harrison Dunn, and Robert Jarret; Griffin, Georgia

- Barkley, N.L., Wang, M.L., Pittman, R.N. 2011. A real-time PCR genotyping assay to detect FAD2A SNPs in peanuts (*Arachis hypogaea* L.). *Electronic Journal of Biotechnology*. 14(1).
- Gremillion, S.K., Culbreath, A.K., Gorbet, D.W., Mullinix Jr, B.G., Pittman, R.N., Stevenson, K.L., Todd, J.W., Escobar, R.E., Condori, M.M. 2011. Field evaluations of leaf spot resistance and yield in peanut genotypes in the United States and Bolivia. *Plant Disease*. 95(3):263-268.
- Morris, J.B. 2011. Morphological, Phenological, and Reproductive Trait Analysis for the Pasture Species, Siratro [*Macroptilium atropurpureum* (DC.) Urb.]. *Tropical Grasslands*. 44/266-273.
- Chen, X., Wang, M.L., Holbrook Jr, C.C., Culbreath, A., Liang, X., Brennenman, T., Guo, B. 2010. Identification and characterization of a multigene family encoding germin-like proteins in cultivated peanut (*Arachis hypogaea* L.). *Plant Molecular Biology Reporter*. DOI 10.1007/s11105-010-0237-6
- Chen, Z., Wang, M.L., Barkley, N.L., Pittman, R.N. 2010. A Simple Allele-Specific PCR Assay for Detecting FAD2 Alleles in Both A and B Genomes of the Cultivated Peanut for High Oleate Trait Selection. *Plant Molecular Biology Reporter*. 28:542-548.
- Antonious, G.F., Snyder, J.C., Burke, T., Jarret, R.L. 2010. Screening *Capsicum chinense* fruits for heavy metals bioaccumulation. *Journal of Environmental Health*. Part B 45:562-571.
- Morris, J.B. 2010. Morphological and reproductive characterization of guar genetic resources regenerated in Georgia, USA. *Genetic Resources and Crop Evolution*. 10.1007/s10722-010-9538-8.
- Wang, M.L., Barkley, N.L., Chinnan, M., Stalker, T., Pittman, R.N. 2010. Oil content and fatty acid composition variability in wild peanut species. *Plant Genetic Resources*. 8:232-234.
- Wang, M.L., Morris, J.B., Pinnow, D.L., Davis, J., Raymer, P., Pederson, G.A. 2010. A survey of the castor oil content, seed weight and seed-coat colour on the United States Department of Agriculture germplasm collection.. *Plant Genetic Resources: Characterization and Utilization*. 8:229-231.
- Antonious, G., Berke, T., Jarret, R.L. Pungency in *Capsicum Chinense*: Variation Among Countries of Origin. 2009 *Journal of Environmental Science and Health*. 44(B)(2): 179-184.
- Antonious, G.F., Berke, T., Jarret, R.L. 2009. Antioxidants in *Capsicum chinense*: Variation among countries of origin. *Journal of Environmental Science and Health*. DOI: 10.1080/03601230903000727 44(B):621-626
- Barkley, N.L., Chamberlin, K.D., Wang, M.L., Pittman, R.N. 2009. Development of a real-time PCR genotyping assay to identify high oleic acid peanuts (*Arachis hypogaea* L.). *Molecular Breeding*. DOI 10.1007/s11032-009-9338-z 25(3):541-548.
- Barkley, N.L., Pinnow, D.L., Wang, M.L., Pederson, G.A. 2009. First Report of Tomato Spotted Wilt Virus Infecting African Clover (*Trifolium temense*) in Georgia. *Plant Disease*. 93(2):202.
- Morris, J.B. 2009. Characterization of Medicinal Senna genetic resources. *Plant Genetic Resources: Characterization and Utilization*. DOI: 10.1017/s1479262109344111 7(3):257-259.
- Morris, J.B. 2009. Characterization of sesame (*Sesamum indicum* L.) germplasm regenerated in Georgia, U.S.A.. *Genetic Resources and Crop Evolution*. DOI 10.1007/S10722-009-9411-9.
- Morris, J.B. 2009. Morphological and reproductive characterization in Hyacinth bean, *Lablab purpureus* (L) Sweet germplasm with clinically proven nutraceutical and pharmaceutical traits for use as a medicinal food. *Journal of Dietary Supplement*. DOI: 10.1080/19390210909070830 6(3):263-279
- Naeem, M., Khan, M., Morris, J.B. 2009. Agrobotanical attributes, nitrogen-fixation, enzyme activities and nutraceuticals and tyrosinase enzyme of hyacinth bean (*Lablab purpureus* L.) - a bio-functional medicinal legume.. *American Journal of Plant Physiology*. 4:58-69.
- Singh, S., Jarret, R.L., Russo, V.M., Shimkus, J.M., Majetich, G., Bushway, R., Perkins, L.B. 2009. Determination of capsinoids by HPLC-DAD in *Capsicum* ssp. Part I. *Journal of Agricultural and Food Chemistry*. DOI.10.1021/jf8040287
- Wang, M., Barkley, N.A., Jenkins, T.M. 2009. Microsatellite Markers in Plants and Insects Part I: Applications of Biotechnology. *Genes, Genomes, and Genomics*. 3(1):54-67.
- Wang, M.L., Chen, C.Y., Davis, J., Guo, B., Stalker, T., Pittman, R.N. 2009. Assessment of Oil content and fatty acid composition variability in different peanut subspecies and botanical varieties. *Plant Genetic Resources*. doi: 10.1017/S1479262109990177. 8:71-73

- Wang, M.L., Pinnow, D.L., Barkley, N.L., Pittman, R.N. 2009. Plant Resistance to TSWV and Seed Accumulation of Resveratrol within Peanut Germplasm and Its Wild Relatives in the US Collection. *Plant Pathology Journal* ISSN 1812-5387 . 8(2):53-61.
- Wang, M.L., Zhu, C., Barkley, N.L., Chen, Z., Erpelding, J.E., Murray, S., Tesso, T., Pederson, G.A., Yu, J. 2009. Genetic Diversity and Population Structure Analysis of Accessions in the US Historic Sweet Sorghum Collection. *Theoretical and Applied Genetics*. Online DOI 10.1007/s00122-009-1155-6. 120:13-23.
- Zhenbang, C., Wang, M.L., Waltz, C., Raymer, P. 2009. Genetic Diversity of Warm-Season Turfgrass: Seashore Paspalum, Bermudagrass, and Zoysiagrass Revealed by AFLPs. *FLORICULTURE, ORNAMENTAL AND PLANT BIOTECHNOLOGY*. Online ISSN 1749-0308 3:1 p.20-24.
- Barkley, N.L., Krueger, R., Federici, C.T., Roose, M.L. 2009. What phylogeny and gene genealogy analyses reveal about homoplasy in citrus microsatellite alleles. *Plant Systematics and Evolution*. 282:71-86
- Jarret, R.L., Berke, T., Baldwin, E.A., Antonious, G. 2009. Variability for Free Sugars and Organic Acids in Capsicum Chinense Jacq.. *Chemistry and Biodiversity*. 6(2):138-145.
- Wang, M.L., Jorge, M., Morris, J.B., Zhenbang, C., Barkley, N.L., Pederson, G.A. Evaluation of Lespedeza Germplasm Genetic Diversity and Its Phylogenetic Relationship with the Genus Kummerowia. *Conservation Genetics*: 10:79-85. 2009 DOI 10.1007/s10592-008-9524-2
- Dida, M., Wanyera, N., Harrison Dunn, M.L., Bennetzen, J., Devos, K. 2008. Population Structure and Diversity in Finger Millet (*Eleusine coracana*) Germplasm.. *Tropical Plant Biology* 1: 131-141.
- Jarret, R.L. 2008. DNA Barcoding in a Crop Genebank: Resolving the Capsicum annum Species Complex.. *The Open Biology Journal*. 1:35-42.
- Wang, M.L., Barkley, N.L., Gillaspie, A.G., Pederson, G.A. 2008. Phylogenetic relationships and genetic diversity of the USDA Vigna germplasm collection revealed by gene-derived markers and sequencing.. *Genetical Research*. 90(6):467-480.
- Wang, M.L., Pittman, R.N. 2008. Resveratrol Content in Seeds of Peanut Germplasm Quantified by HPLC.. *Plant Genetic Resources: Characterization and Utilization*. 7(1) 80-83 (2008).
- Barkley, N.L., Wang, M.L. 2008. Application of TILLING and EcoTILLING as Reverse Genetic Approaches to Elucidate the Function of Genes in Plants and Animals. *Current Genomics*. 9(4):212-226.
- Barkley, N.L., Wang, M.L., Gillaspie Jr, A.G., Dean, R.E., Pederson, G.A., Jenkins, T.M. 2008. Discovering and Verifying DNA Polymorphism in a Mung Bean [*V. radiata* (L). R. Wilczek] Collection by Ecotilling and Sequencing. *Biomed Central (BMC) Notes* 1:28.
- Jarret, R.L., Berke, T. 2008. Variation for Fruit Morphological Characteristics within a Germplasm Collection of Capsicum Chinense Jacq.. *HortScience*. 43:1694-1697.
- Jarret, R.L., Gillaspie Jr, A.G., Barkley, N.L., Pinnow, D.L. 2008. The Occurrence and Control of Pepper Mild Mottle Virus in the USDA/ARS Capsicum Germplasm Collection. *Seed Technology Journal*. 30(1):26-36.
- Morris, J.B. 2008. Characterization of regenerated butterfly pea (*Clitoria ternatea* L.) accessions for morphological, phenology, reproductive and potential nutraceutical, pharmaceutical trait utilization.. *Genetic Resources and Crop Evolution*. 56:421-427
- Morris, J.B. 2008. *Macrotyloma Axillare* and *M. uniflorum*: Descriptor Analysis, Anthocyanin Indexes, and Potential Uses. *Genetic Resources and Crop Evolution* 55: 5-8.
- Morris, J.B. 2008. *Rhynchosia minima* (L.) dc. - regeneration, characterization and potential uses for natural products and flavonoids. *Plant Genetic Resources Newsletter*. 153:15-19
- Williamson, K.S., Morris, J.B., Pye, Q.N., Kamat, C.D., Hensley, K. 2008. A survey of sesamin and composition of tocopherol variability from seeds of 11 diverse sesame (*sesamum indicum* L.) genotypes using hplc-pda-eed. *Phytochemical Analysis* 19: 311-322.
- Fu, Y.B., Chong, J., Fetch, T., Wang, M.L. 2007. Microsatellite Variation in *Avena Sterilis* Oat Germplasm. *Theoretical and Applied Genetics*. *Theoretical Applied Genetics* (2007) 114:1029-1038
- Gillaspie Jr, A.G. 2007. Attempts to Improve the Method of Screening Cowpea Germplasm for Resistance to Cucumber Mosaic Virus and Blackeye Cowpea Mosaic Virus. *Plant Pathology Journal*. 6(2):202-205.
- Jarret, R.L. 2007. Morphological Variation for Fruit Characteristics in the USDA/ARS Capsicum Baccatum L. Germplasm Collection. *HortScience* 42: 1303-1305.
- Wang, M.L., Morris, J.B., Barkley, N.L., Dean, R., Jenkins, T., Pederson, G.A. 2007. Evaluation of Genetic Diversity of the USDA Lablab Purpureus Germplasm Collection Using Simple Sequence Repeat Markers. *Journal of Horticultural Science and Biotechnology*. 82:571-578.
- Wang, M.L., Pinnow, D.L., Pittman, R.N. 2007. Preliminary Screening of Peanut Germplasm in the U.S. Collection for TSWV Resistance and High Flavonoid Content. *Plant Pathology Journal*. 6(3): 219-226.
- Gillaspie Jr, A.G., Barkley, N.L., Morris, J.B. 2006. An unusual strain of alfalfa mosaic virus detected in *crotalaria* spp. germplasm in georgia. *Plant Pathology Journal* 5(3): 397-400.
- Jarret, R.L., Antonious, G., Kochhar, T.S., Snyder, J.C. 2006. Antioxidants in Hot Pepper: Variation Among Accessions. *Journal of Environmental Science and Health, Part B*. 41:1237-1243.

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FUNCTIONAL GENOMICS OF AGRONOMIC TRAITS IN DEVELOPING SEED AND POLLEN IN MAIZE AND SORGHUM – Prem Chourey (P), Alisa Huffaker, and Peter Teal; Gainesville, Florida

- Huffaker, A., Dafoe, N.J., Schmelz, E.A. 2011. ZmPep1, an ortholog of Arabidopsis elicitor Peptide 1, regulates maize innate immunity and enhances disease resistance. *Plant Physiology*. 155(30):1325-1338.
- Xiong, Y., Li, Q., Kang, B., Chourey, P.S. 2011. Discovery of Genes Expressed In Basal Endosperm Transfer Cells in Maize Using 454 Transcriptome Sequencing. *Planta*. DOI 10.1007/s11105-011-0291-8.
- Chourey, P.S., Li, Q., Dibyendu, K. 2010. Sugar - hormone crosstalk in seed development: Two redundant pathways of IAA biosynthesis are regulated differentially in the invertase-deficient miniature1 (mn1) seed mutant in maize.. *Molecular Plant*. 3:1026-1036.
- Rijavec, T., Jain, M., Dermastia, M., Chourey, P.S. 2010. Spatial and temporal profiles of cytokinin biosynthesis and accumulation in developing caryopsis of maize. *Annals Of Botany*. 107: 1235–1245.
- Jain, M., Chourey, P.S., Boote, K.J., Allen Jr, L.H. 2010. Short-term high temperature growth conditions during vegetative-to-reproductive phase transition irreversibly compromise cell wall invertase-mediated sucrose catalysis and microspore meiosis in grain sorghum. *Journal of Plant Physiology*. 167:578-582.
- Yamaguchi, Y., Huffaker, A., Bryan, A.C., Tax, F.E., Ryan, C.A. 2010. PEPR2 is a second receptor for the Pep1 and Pep2 peptides and contributes to defense responses in arabidopsis. *The Plant Cell*. 22:508-522.
- Leclere, S.L., Schmelz, E.A., Chourey, P.S. 2010. Sugar levels regulate tryptophan-dependent auxin biosynthesis in developing maize kernels. *Plant Physiology*. 153: 306-318.

6618-21000-013-00D

GENETIC IMPROVEMENT OF CITRUS – Kim Bowman (P), Randall Niedz, Eddie Stover, and Thomas McCollum; Fort Pierce, Florida

- Maul, D.P., Mccollum, T.G., Guy, C.L., Porat, R. 2011. Temperature conditioning alters transcript abundance of genes related to chilling stress in 'Marsh' grapefruit flavedo. *Postharvest Biology and Technology*. 60:177-185.
- Aradhya, M.K., Stover, E.W., Velasco, D., Koehmstedt, A. 2010. Genetic structure and differentiation in cultivated fig (*Ficus carica* L.). *Genetica*. 138:681-694
- Niedz, R.P., Evens, T.J. 2010. Effect of Agrobacterium culture and inoculation density on transformation efficiency of a citrange (*Citrus reticulata* x *Poncirus trifoliata*). *Journal of Forestry and Horticulture*. 2(3):30-37.
- Niedz, R.P., Evens, T.J. 2010. The effects of benzyladenine and meta-topolin on in vitro shoot regeneration of a citrus citrandarin rootstock. *Research Journal of Agricultural & Biological Sciences*. 6:45-53.
- Stover, E.W., Dangl, G., Yang, J., Bautista, J.C., Aradhya, M.K. 2010. Investigations into the origin of 'Norton' grape using SSR markers. *Proceedings of Florida State Horticultural Society*. 122:19-24.
- Muganu, M., Dangl, G., Aradhya, M.K., Frediani, M., Scossa, A., Stover, E.W. 2009. Ampelographic and DNA Characterization of Local Grapevine Accessions of the Tuscia Area (Latium, Centre of Italy). *American Journal of Enology and Viticulture*. 60:110-115.
- Niedz, R.P., Evens, T.J. 2008. The effects of nitrogen and potassium nutrition on the growth of nonembryogenic and embryogenic tissue of sweet orange (*Citrus sinensis* (L.) Osbeck). *Biomed Central (BMC) Plant Biology*. 8:126. doi:10.1186/1471-2229-8-126
- Bautista, J., Dangl, G., Yang, J., Reisch, B., Stover, E.W. 2008. Use of Genetic Markers to Assess Pedigrees of Grape Cultivars and Breeding Program Selections. *American Journal of Enology and Viticulture*. 59:248-254.
- Ryugo, K., Stover, E.W. 2008. Release of hairless kiwifruit 'Eldorado' and 'Nugget', and 'Early Bird' pollinizer for further evaluation.. *Journal of American Pomological Society*. 62:137-138.
- Stover, E.W., Riaz, S., Walker, M. 2008. Pcr screening for xylella fastidiosa in grape genebank accessions collected in the Southeast US. *American Journal of Enology and Viticulture*. 59:437-439.
- Albrecht, U., Bowman, K.D. 2008. Gene expression in *Citrus sinensis* (L.) Osbeck after infection with the bacterial pathogen *Candidatus Liberibacter asiaticus* causing Huanglongbing in Florida. *Plant Science*. 175(3):291-306
- Maul, D.P., Mccollum, T.G., Popp, M., Guy, C.L., Porat, R. 2008. Transcriptome profiling of grapefruit following exposure to low temperature and conditioning treatments uncovers principal molecular components involved in chilling tolerance and susceptibility.. *Plant Cell and Environment*. 31(6):752-768.
- Albrecht, U., Bowman, K.D. 2007. Inducible Proteins in Citrus Rootstocks with Different Tolerance Towards the Root Rot Pathogen *Phytophthora palmivora*. *Journal of Phytopathology*. 155:606-615
- Chen, C., Bowman, K.D., Choi, Y.A., Dang, P.M., Rao, M., Huang, S., Soneji, J.R., McCollum, T.G., Gmitter, Jr., F.G. 2007. EST-SSR genetic maps for *Citrus sinensis* and *Poncirus trifoliata*. *Tree Genetics & Genomes* [serial online]. Available: <http://www.springerlink.com/content/a2t5271785626233/fulltext.html>
- Colova(Tsolova), V.M., Bordallo, P.N., Phils, B.R., Bausher, M.G. 2007. Synchronized somatic embryo development in embryogenic suspensions of grapevine (*muscadinia rotundifolia* small and *vitis vinifera* L.). *Vitis*. 46(1):15-18.

- Mozoruk, J.J., Hunnicutt, L.E., Hunter, W.B., Bausher, M.G. 2007. CsV03-3, a gene that encodes a protein with DNA binding activity is a member of a novel gene family from *Citrus sinensis* whose expression is responsive to defense signals and abiotic stress.. *Journal of Plant Physiology*. Available: <http://www.ScienceDirect.com>
- Niedz, R.P., Evens, T.J. 2007. Regulating plant tissue growth by mineral nutrition. *In Vitro Cellular and Developmental Biology - Plants*. 43(4):370-381.
- Samson, N., Bausher, M.G., Lee, S., Jansen, R.K., Daniell, H. 2007. The complete nucleotide sequence of the coffee (*Coffea arabica* L.) chloroplast genome: organization and implications for biotechnology and phylogenetic relationships among angiosperms.. *Plant Biotechnology Journal*. 5(2):339-353

6625-21000-003-00D

ENHANCEMENT OF SUGARCANE GERMPLASM FOR DEVELOPMENT OF CULTIVARS AND SUSTAINABLE PRODUCTION
– Barry Glaz (P), Neil Glynn, Duli Zhao, and Serge Edme; Canal Point, Florida

- Davidson, R., Milligan, S.B., Glaz, B.S., Comstock, J.C., Hu, C., Glynn, N.C., Edme, S.J., Holder, D.G., Gilbert, R.A., Sood, S.G., Del Blanco, I.A., Zhao, D. 2011. Registration of 'CPCL 99-4455' Sugarcane. *Journal of Plant Registrations*. 5:54-61.
- Glaz, B.S., Shine, J.M., Irey, M.S., Perdomo, R., Powell, G., Comstock, J.C. 2011. Fiber Content of Three Sugarcane Cultivars in Three Crop Cycles on Sand and Muck Soils. *Agronomy Journal*. 103:211-220.
- Glynn, N.C., Dixon, L.J., Castlebury, L.A., Szabo, L.J., Comstock, J.C. 2010. PCR assays for the sugarcane rust pathogens *Puccinia kuehnii* and *P. melanocephala* and detection of a SNP associated with geographic distribution in *P. kuehnii*. *Plant Pathology*. 59:703-711.
- Glynn, N.C., Gilbert, R.A., Comstock, J.C., Davis, M.J. 2010. Transmission of a Sugarcane Yellow Leaf Virus Resistance Transgene to Sexual Progeny and Screening by Kanamycin Inoculation. *Field Crops Research* 119:308-313.
- Del Blanco, I.A., Glaz, B.S., Edme, S.J. 2010. Improving efficiency of sugarcane genotype selection in Florida. *Crop Sci.*, doi: 10.2135/cropsci2009.09.0539 50:1744-1753.
- Zhao, D., Glaz, B.S., Comstock, J.C. 2010. Sugarcane Response to Water-Deficit Stress during Early Growth on Organic and Sand Soils. *American Journal of Agricultural and Biological Science*. 5(3):403-414.
- Zhao, D., Glaz, B.S., Edme, S.J., Del Blanco, I.A. 2010. Precision of Sugarcane Biomass Estimates in Pot Studies Using Fresh and Dry Weights. *Journal of American Society of Sugar Cane Technologists*. 30:37-49.
- Zhao, D., Reddy, R., Kakani, G.V., Read, J.J. 2010. Remote sensing algorithms for estimating nitrogen uptake and nitrogen use efficiency in cotton. *Acta Agriculturae Scandinavica, Section B-Plant Soil Science* 60:500-509.
- Zhao, D., Wright, D., Marois, D., Mackowiak, C., Brennan, M. 2010. Improved growth and nutrient status of an oat cover crop in sod-based versus conventional peanut-cotton rotations. *Agronomy for Sustainable Development*. 30:497-504.
- Glynn, N.C., Gilbert, R.A., Glaz, B.S., Comstock, J.C., Kang, M.S., Daren, C.W., Tai, P.Y., Miller, J.D. 2009. Repeatability between two intermediate sugarcane genotype selection stages in Florida. *Journal of Crop Improvement* volume 23:252-265
- Lam, E., Shine, J., Da Silva, J., Lawton, M., Bonos, S., Calvino, M., Carrer, H., Castro Silva F, M., Glynn, N.C., Helsel, Z., Ma, J., Richard Jr, E.P., Souza, G., Ming, R. 2009. Improving Sugarcane for Biofuel: Engineering for an even better feedstock. *Global Change Biology*. J. 1: 251-255. 2009.
- Milligan, S.B., Davidson, R.W., Edme, S.J., Comstock, J.C., Hu, C.J., Holder, D.G., Glaz, B.S., Glynn, N.C., Gilbert, R.A. 2009. Registration of 'CPCL 97-2730' Sugarcane. *Journal of Plant Registrations*. 3:158-164.
- Edme, S.J., Davidson, R.W., Gilbert, R.A., Comstock, J.C., Glynn, N.C., Glaz, B.S., Del Blanco, I.A., Miller, J.D., Tai, P. 2009. Registration of 'CP 01-1372' Sugarcane. *Journal of Plant Registrations*. 3:150-157.
- Glaz, B.S., Edme, S.J., Davidson, W.R., Gilbert, R.A., Comstock, J.C., Glynn, N.C., Miller, J.D., Tai, P.Y. 2009. Registration of 'CP 00-2180' Sugarcane. *Journal of Plant Registrations*. 3:35-41.
- Glaz, B.S., Morris, D.R. 2009. Sugarcane Responses to Water-Table Depth and Periodic Flood. *Agronomy Journal*. 102:372-380.
- Glynn, N.C., Mccorkle, K., Comstock, J.C. 2009. Diversity among mainland USA sugarcane cultivars examined by SSR genotyping. *American Society of Sugar Cane Technologists*. volume 29:36-52.
- Comstock, J.C., Glaz, B.S., Edme, S.J., Davidson, W.R., Gilbert, R.A., Glynn, N.C., Miller, J.D., Tai, P.Y. 2009. Registration of 'CP 00-1446' Sugarcane. *Journal of Plant Registrations*. 3:28-34.
- Glaz, B.S., Kang, M.S. 2008. Location Contributions Determined via GGE Biplot Analysis of Multievironment Sugarcane Genotype-Performance Trials. *Crop Science*. 48:941-950 (2008).
- Glaz, B.S., Reed, S.T., Albano, J.P. Sugarcane Response to Nitrogen Fertilization on a Histosol with shallow Water Table and Periodic Flooding. *Journal of Agronomy and Crop Science*. 194:369-379.
- Juarez, J., Miller, J.D., Orozco, H., Solares, E., Tai, P.Y., Comstock, J.C., Glaz, B.S., Queme De Leon, J., Ovalle, W., Edme, S.J., Glynn, N.C., Deren, C. 2008. Registration of 'CP 88-1165' Sugarcane. *Journal of Plant Registrations*. 2:102-109 (2008).

- Suman, A., Kimbeng, C., Edme, S.J., Veremis, J.C. 2008 Sequence-Related Amplified Polymorphism (SRAP) markers for assessing interrelationships and genetic diversity among members of the *Saccharum* complex. *Diversity Plant Genetic Resources Journal*. 6:1-10.
- Gilbert, R.A., Comstock, J.C., Glaz, B.S., Edme, S.J., Davidson, W.R., Glynn, N.C., Miller, J.D., Tai, P.Y. 2008. Registration of 'CP 00-1101' Sugarcane. *Journal of Plant Registrations*. 2:95-101.
- Gilbert, R.A., Glynn, N.C., Comstock, J.C., Davis, M.J. Agronomic performance and genetic characterization of sugarcane transformed for resistance to sugarcane yellow leaf virus. *Field Crops Research*. 111:39-46.
- Gilbert, R.A., Miller, J.D., Comstock, J.C., Glaz, B., Edme, S.J. 2007. Performance of exogenous sugarcane germplasm on mineral soils of Florida, USA. *International Society of Sugar Cane Technologists Proceedings*. 26:388-393.
- Glaz, B.S. 2007. Sugarcane response to month and duration of preharvest flood. *Journal of Crop Improvement*. 20 1/2(39/40):137-152.

6631-21000-017-00D

DEVELOPMENT OF AN INTERNATIONAL MARKER ASSISTED SELECTION PROGRAM FOR CACAO – Raymond Schnell (P) and David Kuhn; Miami, Florida

- Aikpokpodion, P.O., Kolesnikova-Allen, M., Adetimirin, V.O., Guiltinan, M.J., Eskes, A.B., Motamayor, J.C., Schnell li, R.J. 2010. Population structure and molecular characterization of Nigerian field genebank collections of cacao, *Theobroma cacao* L. *Silvae Genetica*. 59(6):273-285.
- Livingstone, D., Motamayor, J.C., Schnell li, R.J., Cariaga, K.A., Freeman, B.L., Meerow, A.W., Brown, J.S., Kuhn, D.N. 2010. Development of Single Nucleotide Polymorphism markers in *Theobroma cacao* and comparison to Simple Sequence Repeat markers for genotyping of Cameroon clones. *Molecular Breeding*. DOI 10.1007/s11032-010-9416-2.
- Kuhn, D.N., Figueira, A., Lopes, U., Motamayor, J., Meerow, A.W., Cariaga, K.A., Freeman, B.L., Livingstone, D., Schnell li, R.J. 2010. Evaluating *Theobroma grandiflorum* for comparative genomic studies with *Theobroma cacao*. *Tree Genetics and Genomes*. DOI 10.1007/s11295-010-0291-0.
- Borrone, J., Brown, J.S., Tondo, C.L., Mauro-Herrera, M., Kuhn, D.N., Violi, H., Sautter, R.T., Schnell li, R.J. 2009. An EST-SSR based linkage map for *Persea americana* Mill. (avocado). *Tree Genetics and Genomes*. 10.1007/s11295-009-0208-y.
- Efombagn, M., Sounigo, O., Eskes, A.B., Motamayor, J.C., Manzanares-Dauleux, J.J., Schnell li, R.J., Nyasse, S. 2009. Parentage analysis and outcrossing patterns in cacao (*Theobroma cacao* L.) farms in Cameroon. *Heredity*. p.1-8.
- Livingstone, D., Freeman, B.L., Tondo, C.L., Cariaga, K.A., Kuhn, D.N. 2009. Improvement of High-throughput Genotype Analysis After Implementation of a Dual-curve Sybr Green I-based Quantification and Normalization Procedure. *HortScience*. 44: 1228-1232.
- Pokou, N.D., N'Goran, J.A., Lachenaud, P., Eskes, A.B., Motamayor, J.C., Schnell li, R.J., Kolesnikova-Allen, M., Clement, D., Sangare, A. 2009. Recurrent selection of cocoa populations in Cote d'Ivoire: comparative genetic diversity between the first and second cycles. *Plant Breeding*. 128: 514-520.
- Violi, H.A., Brown, J.S., Tondo, C.L., Borrone, J.W., Ploetz, R., Schnell li, R.J. 2009. Microsatellite markers reveal low breeding system efficacy and pollen contamination can limit production of full-sib avocado progeny. *Scientia Horticultureae*. doi:10.1016/j.scienta.2008.11.011.
- Araujo, I.S., De Souza Filho, G.A., Faleiro, F.G., Queiroz, V.T., Moreira, M.A., De Barros, E.G., Guimaraes, C.T., Machado, R.C., Pires, J.L., Lopes, U.V., Schnell li, R.J. 2008. Mapping of quantitative trait loci for butter content and hardness in cocoa beans (*Theobroma cacao* L.). *Plant Molecular Biology Reporter*. DOI 10.1007/s11105-008-0069-9.
- Argout, X., Fouet, O., Wincker, P., Gramacho, K., Legavre, T., Sabau, X., Risterucci, A.M., Da Silva, C., Cascardo, J., Allegre, M., Kuhn, D.N., Verica, J., Courtois, B., Loo, G., Babin, R., Sounigo, O., Ducamp, M., Guiltinan, M.J., Ruiz, M., Alemanno, L., Machado, R., Phillips, W., Schnell li, R.J., Gilmour, M., Rosenquist, E., Butler, D., Maximova, S., Lanaud, C. 2008. Towards the understanding of the cocoa transcriptome: Production and analysis of an exhaustive dataset of ESTs of *Theobroma cacao* L. generated from various tissues and under various conditions. *Biomed Central (BMC) Genomics*. 2008.9:512.
- Ayala Silva, T., Gordon, G.G., Heath, R.R. 2008. Use of Satellite Data to Study the Impact of Land-Cover/Land-Use Change in Madison County Alabama. *American Journal of Applied Sciences* 6(4):688-692.
- Borrone, J.W., Tondo, C.T., Kuhn, D.N., Brown, J.S., Schnell li, R.J., Violi, H.A. 2008. Out-crossing in Florida avocados as measured using microsatellite markers. *J. Amer. Soc. Hort Sci.* 133(2):255-261.
- Borrone, J.W., Schnell li, R.J., Violi, H.A., Ploetz, R. 2007. Seventy microsatellite markers from *Persea americana* Miller (avocado) expressed sequence tags. *Molecular Ecology*. 7:439-444.
- Ayala Silva, T., Beyl, C., Dortch, G. 2006. *Agrobacterium* rhizogenes mediated-transformation of *Asimina triloba* L. seedling cuttings. *International Journal of Biological Sciences*.

Ayala Silva, T., Beyl, C., Heath, R.R. 2006. Classification of hyperspectral data and neural networks to differentiate between typical leaves of wheat and those deficient in nitrogen, phosphorus, potassium and calcium. *International Journal of Biological Sciences*.

6631-21000-018-00D

CONSERVATION, CHARACTERIZATION, AND GENETIC IMPROVEMENT OF SUBTROPICAL AND TROPICAL ORNAMENTAL GERMLASM – Alan Meerow (P) and Tomas Ayala Silva; Miami, Florida

- Bay-Smidt, M.B., Jager, A.K., Krydsfeldt, K., Meerow, A.W., Stafford, G.I., Van Staden, Ronsted, N. 2010. Phylogenetic selection of target species in Amaryllidaceae tribe Haemantheae for acetylcholinesterase inhibition and affinity to the serotonin reuptake transport protein. *South African Journal of Botany*. doi:10.1016/j.sajb.2010.07.016.
- Calonje, M., Meerow, A.W., Stevenson, D.W. 2010. The typification of *Zamia erosa* and the priority of that name over *Z. amblyphyllidia*. *Taxon*. 59(2) April 2010:637-642.
- Flagg, R.O., Smith, G.L., Meerow, A.W. 2010. New combinations in *Habranthus* (Amaryllidaceae) in Mexico and Southwestern U.S.A. *Novon*. 20:33-34
- Pathikonda, S., Meerow, A.W., He, J., Mopper, S. 2010. Salinity tolerance and genetic variability in freshwater and brackish *Iris hexagona* colonies. *American Journal of Botany*. 97(9):1438-1443.
- Snijman, D.A., Meerow, A.W. 2010. Floral and macroecological evolution within *Cyrtanthus* (Amaryllidaceae) inferences from combined analyses of plastid *ndhF* and nrDNA ITS sequences. *South African Journal of Botany*. 76:217-238.
- Meerow, A.W. 2010. Convergence or reticulation? Mosaic evolution in the canalized American Amaryllidaceae, in Seberg, O., Petersen, G., Barfod, A.S. and Davis, J.I. (eds.) *Diversity, phylogeny and evolution in the monocotyledons*, pp. 145-168. Aarhus University Press, Aarhus.
- Meerow, A.W. 2010. Tilting at windmills: 20 years of *Hippeastrum* breeding. *Israel Journal of Plants Sciences*. 57:303-31.
- Meerow, A.W., Ayala Silva, T., Irish, B.M. 2010. *Turnera diffusa* 'Luisa', a drought tolerant small shrub for warm climates. *HortScience*. 45(12):1895-1896.
- Namoff, S., Veloz, A., Jimenez, F., Rodriguez-Pena, R.A., Peguero, B., Lewis, C., Moynihan, J., Abdo, M., Maunder, M., Meerow, A.W., Von Wettberg, E., Francisco-Ortega, J. 2010. Sweet drinks are made of this: Conservation genetics of an endemic palm species from the Dominican Republic. *Journal of Heredity*. 102:1-10.
- Mauro-Herrera, M., Meerow, A.W., Perera, L., Russell, J., Schnell li, R.J. 2009. Ambiguous genetic relationships among coconut (*Cocos nucifera* L.) cultivars: the effects of outcrossing, sample source and size, and method of analysis. *Genetic Resources and Crop Evolution*. 57:203-217.
- Meerow, A.W., Borrone, J.W., Couvreur, T.L., Mauro-Herrera, M., Hahn, W.J., Kuhn, D.N., Nakamura, K., Schnell li, R.J. 2009. Phylogenetic analysis of seven *WRKY* genes across the palm subtribe *Attaleinae* (Arecaceae) identifies *Syagrus* as sister to the coconut. *PLoS One*. DOI <http://dx.plos.org/10.1371/journal.pone.0007353>
- Oleas, N.H., Meerow, A.W., Francisco-Ortega, J. 2009. Eight microsatellite loci in *Phaedranassa schizantha* Baker (Amaryllidaceae) and cross-amplification in other *Phaedranassa* species. *Conservation Genetics*. DOI 10.1007/s10592-009-9846-8.
- Maunder, M., Levia, A., Santiago-Valentín, E., Stevenson, D.W., Acevedo-Rodríguez, P., Meerow, A.W., Mejía, M., Clubbe, C., Francisco-Ortega, J. 2008. Plant Conservation in the Caribbean Island Biodiversity Hotspot. *Botanical Review*. 74:197-207.
- Meerow, A.W., Ayala Silva, T. 2008. Three cultivars of *Tecoma guarume*. *HortScience*. 43:286-583. Available: <http://hortsci.ashspublications.org/cgi/content/full/43/2/546?ijkey=RERwggwjhNLghug&keytype=ref>
- Borrone, J.W., Meerow, A.W., Kuhn, D.N., Whitlock, B.A., Schnell li, R.J. 2007. The potential of the *WRKY* gene family for phylogenetic reconstruction: an example from the *Malvaceae*. *Molecular Phylogenetics and Evolution*. 44:1141-1154.
- Francisco-Ortega, J., Santiago-Valentin, E., Acevedo-Rodriguez, P., Lewis, C., Pipoly Iii, J., Meerow, A.W., Maunder, M. 2007. Seed plant genera endemic to the Caribbean Island biodiversity hotspot: A review and a molecular phylogentic perspective. *Botanical Review*. 73:183-234.
- Mauro Herrera, M., Meerow, A.W., Borrone, J.W., Kuhn, D.N., Schnell li, R.J. 2007. Usefulness of *WRKY* gene-derived markers for assessing genetic diversity of Florida coconut cultivars. *Scientia Horticultureae*. 115:19-26.
- Meerow, A.W., Gideon, M., Kuhn, D.N., Motamayor, J., Nakamura, K. 2007. Genetic structure and gene flow among South Florida populations of *Iris hexagona* Walt. (Iridaceae) assessed with 19 microsatellite DNA loci. *International Journal of Plant Science*. 168:1291-1309.
- Meerow, A.W., Nakamura, K. 2007. Ten microsatellite loci from *Zamia integrifolia* (Zamiaceae). *Molecular Ecology Notes*. 7(5):824-826.

- Meerow, A.W., Reveal, J.L., Snijman, D.A., Dutilh, J.H. Superconservation proposal to conserve Amaryllidaceae(1805)over Alliaceae(1797). *Taxon* 56: 1299-1300. 2007.
- Moynihan, J., Meerow, A.W., Francisco-Ortega, J. 2006. Isolation, characterization and cross-species amplification of microsatellite loci in the cycad genus *dioon* (zamiaceae). potential utilization in population genetics studies of *dioon* edule. *Molecular Ecology Notes*. 7:72-74.

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CONSERVATION AND UTILIZATION OF SUBTROPICAL/TROPICAL FRUIT CROPS, SUGARCANE, AND TRIPSACUM GENETIC RESOURCES – Raymond Schnell (P), David Kuhn, and Tomas Ayala Silva; Miami, Florida

- Ayala Silva, T., Schnell II, R.J. 2010. The tropical and subtropical germplasm collections at the national germplasm repository in Miami, Fl. *Journal of American Pomological Society*. 64(1):28-37.
- Ayala Silva, T., Schnell II, R.J., Meerow, A.W., Brown, J.S., Gordon, G.G. 2009. A Study on the Morphological and PhysicoChemical Characteristics of Five Cooking Bananas. *Journal of Agronomy*. P.1-6.
- Brown, J.S., Schnell II, R.J., Ayala Silva, T., Moore, J.M., Tondo, C.L., Winterstein, M.C. 2009. Broad-sense heritability estimates for fruit color and morphological traits from open-pollinated half-sib mango families. *HortScience*. 44(6):1-5.
- Schnell II, R.J., Tondo, C.L., Brown, J.S., Ayala Silva, T., Borrone, J.W., Davenport, T.L., Kuhn, D.N. 2009. Out-crossing between 'Bacon' pollinizers and adjacent 'Hass' avocado trees and the description of two new lethal mutants. *HortScience*. 44(6):1522-1526.
- Ying, Z., Davenport, T.L., Faber, B., Zhang, T., Schnell II, R.J., Tondo, C.L. 2009. Re-evaluation of honeybees and wind on pollination of avocado. *Journal of Horticultural Science and Biotechnology*. 84(3)255-260.
- Ying, Z., Davenport, T.L., Zhang, T., Schnell II, R.J., Tondo, C.L. 2009. Selection of Highly Informative Microsatellite Markers to Identify Pollen Donors in 'Hass' Avocado Orchards. *Plant Mol Biol Rep* 27:374-380.
- Aikpokpodion, P.O., Kolesnikova-Allen, M., Adetimirin, V.O., Motamayor, J., Adu-Ampomah, Y., Eskes, A.B., Schnell II, R.J. 2009. Genetic diversity assessment of sub-samples of cacao, *Theobroma cacao* L. collections in West Africa using simple sequence repeats marker.. *Tree Genetics & Genomics*. DOI 10.1007/s11295-009-0221-1.
- Brown, J.S., Sautter, R.T., Tondo, C.T., Borrone, J.W., Kuhn, D.N., Motamayor, J.C., Schnell II, R.J. 2008. A composite linkage map from the combination of three crosses made from commercial clones of cacao, *T. cacao* L.. *Tropical Plant Biology*. DOI 10.1007/s12042-008-9011-4.
- Efombagan, I.B., Motamayor, J.C., Sounigo, O., Eskes, A.B., Nyasse, S., Cilas, C., Schnell II, R.J., Manzanares-Dauleux, M., Kolesnikova-Allen, M. 2008. Genetic diversity and structure of farm and genebank accessions of cacao (*Theobroma cacao* L.) in Cameroon revealed by microsatellite markers. *Tree Genetics & Genomics*. DOI 10.1007/s11295-008-0155-z.
- Kuhn, D.N., Motamayor, J.C., Meerow, A.W., Borrone, J.W., Schnell II, R.J. 2008. SSCP markers provide a useful alternative to microsatellites in genotyping and estimating genetic diversity in populations and germplasm collections of plant specialty crops. *Electrophoresis*. 29(18):1-14.
- Motamayor, J.C., Lachenaud, P., Wallace, J., Loo, G., Kuhn, D.N., Brown, J.S., Schnell II, R.J. 2008. Geographic and genetic population differentiation of the Amazonian chocolate tree. *PLoS One*. Available: DOI: 10.1007/s12042-008-9011-4
- Brown, J.S., Phillips-Mora, W., Power, E.J., Krol, C.A., Cervantesmartinez, C., Motamayor, J., Schnell II, R.J. 2007. Mapping QTL for resistance to frosty pod and black pod diseases, and for horticultural traits in *Theobroma cacao* L.. *Crop Science*. 47:1851-1858.
- Opoku, S.Y., Bhattacharjee, R., Kolesnikova-Allen, M., Motamayor, J.C., Schnell II, R.J., Ingelbrecht, I., Enu-Kwesi, L., Adu-Ampomah, Y. 2007. Genetic diversity in cocoa (*Theobroma cacao*, L.) germplasm collection from Ghana. *Journal of Crop Improvement*. Vol. 20(1/2) (#39/40)p.73-87.
- Schnell II, R.J., Brown, J.S., Kuhn, D.N., Cervantesmartinez, C., Borrone, J.W., Olano, C.T., Motamayor, J.C., Phillips, W., Johnson, E., Monteverde-Penso, E.J., Amores, F., Lopes, U. 2007. Current challenges of tropical tree crop improvement: integrating genomics into an applied cacao breeding program. *Acta Horticulturae: Proc IS on biotechnology of temperate fruit crops and tropical species*. 738:p129-144.
- Schnell II, R.J., Kuhn, D.N., Brown, J.S., Tondo, C.T., Motamayor, J.C. 2007. Development of marker assisted selection program for cacao. *Journal of Phytopathology*. 97(12):1664-1669.
- Faleiro, F.G., Queiroz, V.T., Lopes, U.V., Guimaraes, C.T., Pires, J.L., Yamada, M.M., Araujo, I.S., Pereira, M.G., Schnell II, R.J., De Souza Filho, G.A. 2006. Mapping QTLs for Witches'Broom (*Crinipellis perniciosa*) Resistance in cacao (*Theobroma cacao* L.). *Euphytica*. 149:227-235

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GENETIC RESOURCE MANAGEMENT OF TROPICAL/SUBTROPICAL GENETIC RESOURCES AND ASSOCIATED INFORMATION – Brain Irish (P) and Ricardo Goenaga; Mayagüez, Puerto Rico

- Irish, B.M., Goenaga Portela, R.J., Zhang, D., Schnell li, R.J., Brown, J.S., Motamayor, J. 2010. Microsatellite fingerprinting of the USDA-ARS Tropical Agriculture Research Station cacao (*Theobroma cacao* L.) germplasm collection. *Crop Science*. 50:656-667.
- Irish, B.M., Crespo, A., Goenaga, R.J., Ayala Silva, T., Niedz, R.P. 2009. Ploidy level and genomic composition of the USDA-ARS Tropical Agriculture Research Station Musa sp. Germplasm Collection. *Journal of Agriculture of the University of Puerto Rico*, 93(1-2):1-22.
- Irish, B.M., Correll, J.C., Koike, S.T., Morelock, T.E. 2007. Three new races of the spinach downy mildew pathogen identified by a modified set of spinach differentials. *Plant Disease*. 91(11):1392-1396.
- Irish, B.M., Goenaga, R., Park, S., Kang, S. 2007. First Report of *Phytophthora palmivora*, causal agent of black pod, on cacao (*Theobroma cacao* L.) in Puerto Rico. *Plant Disease*. 91:1051.

6635-21000-048-00D

CHARACTERIZATION, CONVERSION, AND IMPROVEMENT OF COMMON BEAN GERmplasm – Timothy Porch Clay (P); Mayagüez, Puerto Rico

- Beaver, J.S., Porch Clay, T.G., Zapata, M. Registration of 'Badillo' Light Red Kidney Bean. *Journal of Plant Registrations*. 4:1-4.
- Porch Clay, T.G., Smith, J.R., Beaver, J.S., Griffiths, P.D., Canaday, C. 2010. Registration of TARS-HT1 and TARS-HT2 heat tolerant dry bean germplasm lines. *HortScience*. 1278-1280.
- Porch, T.G., Ramirez, V.H., Santana, D., Harmsen, E.W. 2009. Evaluation of drought tolerance in common bean germplasm in Juana Díaz, Puerto Rico. *Journal of Agronomy and Crop Science* 195:1-7.
- Porch, T.G., Blair, M., Lariguet, P., Galeano, C., Pankhurst, C., Broughton, W.J. 2009. Generation of a mutant population for TILLING common bean genotype BAT 93. *Journal of the American Society for Horticultural Science*. 134:348-355.
- Beaver, J.S., Porch, T.G., Zapata, M. 2008. Registration of 'verano' white bean. *Journal of Plant Registrations*. 2:187-189.
- Ramirez, V.H., Porch, T.G., Harmsen, E.W. 2008. Development of linear models for estimation of leaflet area in common bean (*Phaseolus vulgaris* L.). *J.Agric. Univ. P.R.* 92(3-4):171-182.

6635-21000-049-00D

CONSERVATION, MANAGEMENT, AND GENETIC ENHANCEMENT OF SORGHUM GENETIC RESOURCES AND ASSOCIATED INFORMATION – Hugo Cuevas (P); Mayagüez, Puerto Rico

- Erpelding, J.E. 2010. Anthracnose disease response in the Burundi sorghum germplasm collection. *Agriculture and Biology Journal of North America*. 1(6):1119-1125.
- Erpelding, J.E. 2010. Field assessment of anthracnose disease response for the sorghum germplasm collection from the Mopti region of Mali. *American Journal of Agricultural and Biological Science* 5(3):363-369.
- Erpelding, J.E., Prom, L.K. 2009. Response to anthracnose infection for a subset of Ethiopian sorghum germplasm. *Journal of Agriculture of the University of Puerto Rico*. 93(3-4):195-206.
- Erpelding, J.E. 2009. Anthracnose disease response for photoperiod-insensitive Ethiopian germplasm from the U.S. sorghum collection. *World Journal of Agricultural Sciences*. 5(6):707-713.
- Erpelding, J.E. 2008. Field evaluation of anthracnose resistance for sorghum germplasm from the Sikasso region in Mali. *The Open Agriculture Journal*. (2):113-120.
- Erpelding, J.E. 2008. Field evaluation of anthracnose disease response for the sorghum germplasm collection from the Kayes region of Mali. *Tropical and Subtropical Agroecosystems*. (8):291-296.
- Erpelding, J.E. 2008. Field evaluation of foliar anthracnose disease response for sorghum germplasm from the Matabeleland North Province of Zimbabwe. *African Journal of Agricultural Research*. Vol 3(11), pp. 792-796.
- Erpelding, J.E. 2008. Sorghum germplasm resistance to anthracnose. *The Americas Journal of Plant Science and Biotechnology*. 2(1):42-46.
- Erpelding, J.E. 2007. Inheritance of Anthracnose Resistance for the Sorghum Cultivar Redlan. *Plant Pathology Journal*. 6 (2): 187-190.
- Erpelding, J.E., Wang, M. L. 2007. Response to Anthracnose Infection for a Random Selection of Sorghum Germplasm. *Plant Pathology Journal* 6(2): 127-133.

6645-21000-028-00D

PHYSIOLOGICAL/BIOCHEMICAL MECHANISMS ASSOCIATED WITH GENETIC ALTERATION OF SOYBEAN QUALITY AND PRODUCTIVITY – Robert Upchurch (P), Earl Taliencio, Thomas Carter Jr., and Joseph Burton; Raleigh, North Carolina

- Upchurch, R.G. and Ramirez, M.E. 2010. Defense-related gene expression in soybean leaves and seeds inoculated with *Cercospora kikuchii* and *Diaporthe phaseolorum* var. *meridinales*. *Physiological and Molecular Plant Pathology*.
- Taliencio, E.W., Scheffler, J.A., Scheffler, B.E. 2010. Characterization of Two Cotton (*Gossypium hirsutum*) Invertase Genes. (Available online in *Molecular Biology Reports*).
- Taliencio, E.W., Scheffler, J.A., and Kwanyuen, P. 2010. Nitrogen Metabolism in Cotton Stems and Roots during Reproductive. *Journal of Cotton Science*, 14:107–112.
- Gilsinger, J.J., Burton, J.W., Carter Jr, T.E. 2010. Maternal Effects on Fatty Acid Composition of Soybean Seed Oil. *Crop Sci.* 50:1874-1881.

6645-21220-012-00D

INCREASING THE COMPETITIVE POSITION OF U.S. SOYBEANS IN GLOBAL MARKETS THROUGH GENETIC DIVERSITY AND PLANT BREEDING – Thomas Carter (P), Joseph Burton, and Robert Upchurch; Raleigh, North Carolina

- Place, G.T., Reberg-Horton, S.C., Carter Jr, T.E., Smith, A.N. 2011. Effects of Soybean Seed Size on Weed Competition. *Agronomy Journal*. 103:175-181.
- Carter Jr, T.E., Rzewnicki, P.E., Burton, J.W., Villagarcia, M.R., Bowman, D.T., Taliencio, E.W., Kwanyuen, P. 2010. Registration of N6202 soybean germplasm with high protein, good yield potential, large seed and diverse pedigree. *J. of Plant Registrations*. 4:77-79.
- Upchurch, R.G. and Ramirez, M.E. 2010. Gene expression profiles of soybeans with mid-oleic acid seed phenotype. *J. of the Am. Oil Chem. Soc.* 87:857-864.
- Carter Jr, T.E., Burton, J.W., Rzewnicki, P.E., Villagarcia, M.R., Fountain, M.O., Taliencio, E.W., Bowman, D. 2009. Registration of 'N8101' small-seeded soybean. *Journal of Plant Registrations*, 3:22-27.
- Lee, J., Smothers, S.L., Dunn, D., Villagarcia, M.R., Shumway, C.R., Carter Jr, T.E., Shannon, G. Evaluation of a Simple Method to Screen Soybean Genotypes for Salt Tolerance. *Crop Sci.* 2008 48: 2194-2200.
- Carter Jr, T.E., Burton, J.W., Fountain, M.O., Rzewnicki, P.E., Villagarcia, M.R., Bowman, D.T. 2008. Registration of soybean cultivar 'N8001'. *Journal of Plant Registrations*, 2: 22-23.
- Naegle, E., Kwanyuen, P., Burton, J.W., Carter Jr, T.E., Rufty, T. 2008. Seed Nitrogen Mobilization in Soybean: Effects of Seed Nitrogen Content and External Nitrogen Fertility. *Journal of Plant Nutrition*, 31:367-379.
- Upchurch, R.G. 2008. Fatty acid unsaturation, mobilization, and signaling in the response of plants to stress. *Biotechnol. Lett.* 30:967-977.
- Wrather, A., Shannon, G., Carter Jr, T.E., Bond, J., Rupe, J., Almeida, A. 2008. Reaction of Drought Tolerant Soybean Genotypes to *Macrophomina phaseolina*. *Plant Health Progress*. Online. doi:10.1094/PHP-2008-0618-01-RS.
- Xue, H., Upchurch, R.G., Kwanyuen, P. 2008. Relationships between oleic and linoleic acid content and seed colonization by *Cercospora kikuchii* and *Diaporthe phaseolorum*. *Plant Disease* 92:1038-1042.
- Zhang, P., Burton, J.W., Upchurch, R.G., Shanklin, J., Whittle, E., Dewey, R. 2008. Mutations in a novel 9-stearoyl-ACP-desaturase gene are associated with enhanced stearic acid levels in soybean seeds. *Crop Sci.* 48:2305-2313.
- Byfield, G., Upchurch, R.G. 2007. Effect of temperature on microsomal omega-3 linoleate desaturase gene expression and linolenic acid content in developing soybean seeds. *Crop Science*, 47:2445-2452.
- Carter Jr, T.E., Burton, J.W., Fountain, M.O., Rzewnicki, P.E., Villagarcia, M.R., Bowman, D.T. 2007. Registration of soybean cultivar 'N7002'. *Journal of Plant Registrations*, 1:93-94.
- Hufstetler, V.E., Boerma, R., Carter Jr, T.E., Earl, H.J. 2007. Genotypic variation for three physiological traits affecting drought tolerance in soybean. *Crop Sci.* 47:25-35.

6645-21220-013-00D

USING GENETIC DIVERSITY OF IMPROVE QUANTITATIVE DISEASE RESISTANCE AND AGRONOMIC TRAITS OF CORN – James Holland (P) and Peter Balint Kurti; Raleigh, North Carolina

- Kump, K., Bradbury, P., Buckler Iv, E.S., Belcher, A., Oropeza-Rosas, M., Wissner, R., Zwonitzer, J., Kresovich, S., McMullen, M.D., Ware, D., Balint Kurti, P.J., Holland, J.B. 2011. Genome-Wide Association Study of Quantitative Resistance to Southern Leaf Blight in the Maize Nested Association Mapping Population. *Nature Genetics*. 43; 163-168.

- Balint Kurti, P.J., Yang, J., Van Esbroek, G., Jung, J., Smith, M. 2010. Use of an advanced intercross line population for mapping of quantitative trait loci for northern leaf blight resistance in maize and for the investigation of multiple disease resistance. *Crop Science*. 50: 458-466.
- Balint Kurti, P.J., Simmons, S., Blum, J., Ballare, C., Stapleton, A. 2010. Maize Leaf Epiphytic Bacteria Diversity Patterns Are Genetically Correlated with Resistance to Fungal Pathogen Infection. *Biology of Plant Microbe Interactions*. 23: 473-484.
- Chintamanani, S., Hulbert, S., Johal, G., Balint Kurti, P.J. 2010. Identification of a Maize Locus that Modulates the Hypersensitive Defense Response, Using Mutant-Assisted Gene Identification and Characterization (MAGIC). *Genetics*. 184:813-825.
- Chung, C., Longfellow, J.M., Walsh, E.K., Kerdieh, Z., Van Esbroek, G., Balint Kurti, P.J., Nelson, R.J. 2010. Resistance loci affecting distinct stages of fungal pathogenesis in maize: use of introgression lines for QTL mapping and characterization. *Biomed Central (BMC) Plant Biology*. 10:103.
- Coles, N.D., McMullen, M.D., Balint Kurti, P.J., Pratt, R.C., Holland, J.B. 2010. Genetic Control of Photoperiod Sensitivity in Maize. *Genetics*. 184:799-812.
- Eller, M.S., Payne, G.A., Holland, J.B. 2010. Selection for Reduced Fusarium Ear Rot and Fumonisin Content in Advanced Backcross Maize Lines and Their Topcross Hybrids. *Crop Science*. 50:2249-2260.
- Glover, J.D., Reganold, J.P., Bell, L.W., Borevitz, J., Brummer, E.C., Buckler IV, E.S., Cox, C.M., Cox, T., Crews, T.E., Culman, S.W., Dehann, L.R., Eriksson, D., Gill, B., Holland, J.B., Hu, F.Y., Hulke, B.S., Ibrahim, A., Jackson, W., Jones, S., Murray, S., Paterson, A.H., Ploschuk, E., Sacks, E.J., Snapp, S., Tao, D.Y., Van Tassel, D., Wade, L., Wyse, D., Xu, Y. 2010. Increasing Food and Ecosystem Security through Perennial Grain Breeding. *Science*. 328:1638-1639.
- Gonzalo, M., Holland, J.B., Vyn, T., McIntyre, L. 2010. Direct Mapping Of Density Response in Recombinant Inbred Lines of Maize (*Zea mays* L.). *Heredity*. 104:583-599.
- Zwonitzer, J., Coles, N., Krakowsky, M.D., Holland, J.B., McMullen, M.D., Pratt, R., Balint Kurti, P.J. 2010. Mapping Disease Resistance QTL for Three Foliar Diseases of Maize in a RIL Population. *Journal of Phytopathology*. 100:72-79.
- Yan, W., Holland, J.B. 2010. A HERITABILITY-ADJUSTED GGE BILOT FOR TEST ENVIRONMENT EVALUATION. *Euphytica*. 171(3):355-369.
- Eller, M.S., Robertson-Hoyt, L., Payne, G., Holland, J.B. 2009. Grain Yield and Fusarium Ear Rot of Maize Hybrids Developed From Lines With Varying Levels of Resistance. *Maydica*. 53:231-237.
- Poland, J., Balint Kurti, P.J., Wisser, R., Pratt, R., Nelson, R. 2009. Shades of gray: The world of quantitative disease resistance. *Trends in Plant Science*. 4:21-29.
- Zwonitzer, J.C., Bubeck, D.M., Bhattaramakki, D., Goodman, M.M., Arellano, C., Balint Kurti, P.J. 2009. Use of backcross recurrent selection and QTL mapping to identify loci contributing to southern leaf blight resistance in a highly resistant maize line. *Theoretical and Applied Genetics*. 118:911-925.
- Balint Kurti, P.J., Zwonitzer, J., Wisser, R. 2008. Use of an Advanced Intercross Line Population for Precise Mapping of Quantitative Trait Loci for Gray Leaf Spot Resistance in Maize. *Crop Science*. 48:1696-1703.
- Balint Kurti, P.J., Zwonitzer, J., Wisser, R., Pe, E., Pea, G., Lee, M., Cardinal, A. 2008. Identification of Quantitative Trait Loci for Resistance to Southern Leaf Blight and Days to Anthesis in Two Maize Recombinant Inbred Line Populations. *Phytopathology*. 98:315-320.
- Eller, M.S., Payne, G., Holland, J.B. 2008. Breeding for Improved Resistance to Fumonisin Contamination in Maize. *Toxin Review*. 27:371-389.
- Johal, G., Balint Kurti, P.J., Weil, C. 2008. Mining and harnessing natural variation - a little MAGIC. *Crop Science*. 48:2066-2073.
- Nelson, P., Coles, N., Holland, J.B., Bubeck, D., Smith, S., Goodman, M. 2008. Molecular Characterization of Maize Inbreds with Expired U.S. Plant Variety Protection. *Crop Science*. 48:1673-1685.
- Van Esbroeck, G., Ruiz, J., Sanchez, J., Holland, J.B. 2008. A comparison of leaf appearance rates among teosinte, maize landraces and modern maize. *Maydica*. 53:117-153.
- Ruiz Corral, J., Duran Puga, N., Sanchez, J., Ron Parra, J., Gonzalez, D., Holland, J.B., Medina, G. 2008. Climatic Adaptation and Ecological Descriptors of 42 Mexican Maize (*Zea Mays* L.) Races. *Crop Science*. 48:1502-1512.
- Balint Kurti, P.J., Zwonitzer, J., Wisser, R., Oropeza-Rosas, M., Holland, J.B., Szalma, S., Carson, M.L. 2007. Precise mapping of Quantitative Trait Loci for Resistance to Southern Leaf Blight, caused by *Cochliobolus heterostrophus* race O, using an Advanced Intercross Maize Population. *Genetics*. 176:645-657.
- Holland, J.B., Robertson-Hoyt, L.A., Kleinschmidt, C.E., White, D.G., Payne, G.A., Maragos, C.M. 2007. Relationships of resistance to Fusarium ear rot and fumonisin contamination with agronomic performance of maize. *Crop Science*. 47(5):1770-1778.
- Gonzalo, M., Vyn, T.J., Holland, J.B., McIntyre, L.M. 2007. Mapping reciprocal effects and interactions with plant density stress in *zea mays* L. *Heredity*. 99:14-30.
- Holland, J.B., Robertson-Hoyt, L., Betran, J., Payne, G., White, D., Isakeit, T., Maragos, C.M., Molnar, T. 2007. Relationships among resistances to Fusarium and Aspergillus ear rots and contamination by fumonisin and aflatoxin in maize. *Phytopathology*. 97(3):311-317.

Jines, M.P., Balint Kurti, P.J., Robertson-Hoyt, L.A., Molnar, T., Holland, J.B., Goodman, M.M. 2006. Mapping resistance to southern corn rust in a semi-tropical recombinant inbred topcross population.. *Theoretical and Applied Genetics*. 114:659-667.

6645-22000-016-00D

EVALUATION AND IMPROVEMENT OF CEREAL GERMLASM FOR DISEASE RESISTANCE AND WINTER-HARDINESS – Gina Brown Guedira (P), Christina Cowger, David Marshall, and David Livingstone; Raleigh, North Carolina

- Kang, J., Clark, A., Vansanford, D., Griffey, C., Brown Guedira, G.L., Dong, Y., Murphy, J.P., Costa, J. 2011. Exotic Scab Resistance Quantitative Trait Loci (QTL) Effects on Soft Red Winter Wheat. *Crop Science*. 51:924-933.
- Griffey, C., Brooks, W., Vaughn, M., Thomason, W., Paling, J., Pitman, R., Dunaway, D., Corbin, R., Kenner, J., Hokanson, E., Behl, H., Beahm, B., Liu, S., Gundrum, P., Brann, D., Whitt, D., Custis, J., Starner, D., Gulick, S., Ashburn, S., Jones, N., Marshall, D.S., Fountain, M.O., Tuong, T.D., Premakumar, R., Livingston, D.P., Hicks, K.B., Kurantz, M.J., Taylor, F., Moreau, R.A. 2011. Registration of 'Dan' Winter Hulless Barley. *Crop Science*. doi: 10.3198/jpr2010.03.0131crc.
- Costa, J.M., Bockelman, H., Brown Guedira, G.L., Cambron, S., Chen, X., Cooper, A., Cowger, C., Dong, Y., Grybauskas, A., Jin, Y., Kolmer, J., Murphy, J.P., Sneller, C., Souza, E. 2010. Registration of the Soft Red Winter Wheat Germplasm MD01W233-06-1 Resistant to Fusarium Head Blight. *Journal of Plant Registrations*. 4:1-6.
- Mundt, C.C., Sackett, K.E., Wallace, L.D., Cowger, C., Dudley, J.P. 2010. Aerial dispersal and multiple-scale spread of epidemics. *EcoHealth*. DOI: 10.1007/s10393-009-0251-z.
- Olson, E.L., Brown Guedira, G.L., Marshall, D.S., Jin, Y., Mergoum, M., Lowe, I., Dubcovsky, J. 2010. Genotyping of U.S. Wheat Germplasm for Presence of Stem Rust Resistance Genes Sr24, Sr36 and Sr1RSAmigo. *Crop Science*. 50:668-675.
- Cowger, C., Arellano, C. 2010. Plump Kernels with High Deoxynivalenol Linked to Late Gibberella zeae Infection and Marginal Disease Conditions in Winter Wheat. *Phytopathology*. 100(7):719-728.
- Cowger, C., Weisz, R., Anderson, J.M., Horton, J. 2010. Maize Debris Increases Barley Yellow Dwarf Virus Severity in North Carolina Winter Wheat. *Agronomy Journal*. 102:688-696.
- Griffey, C.A., Thomason, W.E., Pitman, R.M., Beahm, B.R., Paling, J.J., Chen, J., Fanelli, J.K., Dunaway, D.W., Brooks, W.S., Vaughn, M.E., Hokanson, E.G., Behl, H.D., Corbin, R.A., Hall, M.D., Liu, S., Custis, J.T., Waldenmaier, C.M., Starner, D.E., Gulick, S.A., Ashburn, S.R., Whitt, D.L., Bockelman, H.E., Souza, E.J., Brown Guedira, G.L., Kolmer, J.A., Long, D.L., Jin, Y., Chen, X., Cambron, S.E. 2010. Registration of 'Jamestown' Wheat. *Journal of Plant Registrations*. 4:28-33.
- Griffey, C.A., Thomason, W.E., Pitman, R.M., Beahm, B.R., Paling, J.J., Chen, J., Gundrum, P.G., Fanelli, J.K., Kenner, J.C., Dunaway, D.W., Brooks, W.S., Vaughn, M.E., Hokanson, E.G., Behl, H.D., Corbin, R.A., Hall, M.D., Liu, S., Custis, J.T., Waldenmaier, C.M., Starner, D.E., Gulick, S.A., Ashburn, S.R., Jones, E.H., Whitt, D.L., Bockelman, H.E., Souza, E.J., Brown Guedira, G.L., Kolmer, J.A., Long, D.L., Jin, Y., Chen, X., Cambron, S.E. 2010. Registration of '3434' Wheat. *Journal of Plant Registrations*. 4:44-49.
- Griffey, C.A., Thomason, W.E., Pitman, R.M., Beahm, B.R., Paling, J.J., Chen, J., Gundrum, P.G., Fanelli, J.K., Kenner, J.C., Dunaway, D.W., Brooks, W.S., Vaughn, M.E., Hokanson, E.G., Behl, H.D., Corbin, R.A., Hall, M.D., Liu, S., Custis, J.T., Waldenmaier, C.M., Starner, D.E., Gulick, S.A., Ashburn, S.R., Jones, E.H., Whitt, D.L., Bockelman, H.E., Souza, E.J., Brown Guedira, G.L., Kolmer, J.A., Long, D.L., Jin, Y., Chen, X., Cambron, S.E. 2010. Registration of 'Shirley' Wheat. *Journal of Plant Registrations*. 4:38-44.
- Hall, M.D., Tucker, D., Griffey, C.A., Liu, S., Sneller, C., Guttieri, M., Van Sanford, D., Costa, J., Marshall, D.S., Brown Guedira, G.L. 2010. Registration of USG 3209/Jaypee Wheat Recombinant Inbred Line Mapping Population. *Journal of Plant Registrations*. 4:159-162.
- Li, R., Qu, R., Bruneau, A., Livingston, D.P. 2010. Selection for freezing tolerance in St. Augustine grass under controlled conditions. *Crop Science*. 129:417-421.
- Cowger, C., Parks, W.R., Marshall, D.S. 2009. Appearance of Powdery Mildew of Wheat (caused by *Blumeria graminis* f. sp. *Tritici*) on Pm17-bearing Cultivars in North Carolina. *Plant Disease*. 93:1219.
- Cowger, C., Patton Ozkurt, J.L., Brown Guedira, G.L., Perugini, L. 2009. Post-flowering moisture increased Fusarium head blight and deoxynivalenol levels in North Carolina field experiment with winter wheat. *Phytopathology*. 99:320-327.
- Lagudah, E.S., Krattinger, S.G., Herrera-Foessel, S., Singh, R.P., Huerta-Espino, J., Spielmeier, W., Brown Guedira, G.L., Selter, L.L., Keller, B. 2009. Gene-specific markers for the wheat gene Lr34/Yr18/Pm38 which confers resistance to multiple fungal pathogens. *Theoretical and Applied Genetics*. 119:889-898.
- Livingston, D.P., Tuong, T.D., Haigler, C., Avci, U., Tallury, S. 2009. Microwave Processing of Crowns from Winter Cereals for Light Microscopy.. *Journal of Microscopy*. 49:1837-1842.

- Livingston, D.P., Tuong, T.D., Owen, S. 2009. Carbohydrate changes in winter oat crowns during recovery from freezing. *Dynamic Biochemistry, Process Biotechnology and Molecular Biology*. 3:16-22.
- Maxwell, J.J., Lyerly, J.H., Cowger, C., Marshall, D.S., Brown Guedira, G.L., Murphy, J.P. 2009. MIAG12: A *Triticum timopheevii*-Derived Mildew Resistance Gene in Common 1 Wheat on Chromosome 7AL. *Theoretical and Applied Genetics*. 119:1489–1495.
- Parks, R., Carbone, I., Murphy, J., Cowger, C. 2009. Population Genetic Analysis of an Eastern U.S. Wheat Powdery Mildew Population Reveals Geographic and Recent Common Ancestry with U.K. and Israeli Populations. *Phytopathology*. 99:840-849.
- Wooten, D.R., Livingston, D.P., Lyerly, H.J., Holland, J.B., Jellen, E.N., Marshall, D.S., Murphy, J. 2009. Quantitative Trait Loci and Epistasis for Oat Winter Hardiness Component Traits. *Crop Science*. 48:149-157.
- Hall, M.D., Fritz, A.K., Brown Guedira, G.L. 2009. Genetic Analysis of Resistance to Soil-borne Wheat Mosaic Virus Derived from *Aegilops tauschii*. *Euphytica*. 169:169-176.
- Mundt, C.C., Sackett, K.E., Wallace, L.D., Cowger, C., Dudley, J.P. 2009. Long Distance Dispersal and Accelerating Waves of Disease: Empirical Relationships. *The American Naturalist*. 173:456-466.
- Brown Guedira, G.L., Griffey, C., Kolb, F., Mckendry, A., Murphy, P., Vansanford, D. 2008. Breeding FHB-resistant soft winter wheat: progress and prospects. *Cereal Research Communications*. 36:31S-35S.
- Coram, T., Brown Guedira, G.L., Chen, X. 2008. Using Transcriptomics to Understand the Wheat Genome. *CAB Reviews: Perspectives in Agriculture, Veterinary Science, Nutrition and Natural Resources*. 3:083.
- Cowger, C., Brunner, P.C., Mundt, C.C. 2008. Frequency of Sexual Recombination by *Mycosphaerella graminicola* in Mild and Severe Epidemics. *Phytopathology*. 98:752-759.
- Cowger, C., Weisz, R. 2008. Blends of Soft Red Winter Wheat Varieties Increased Yield in North Carolina. *Agronomy Journal*. 100:169-177.
- Parks, W., Carbone, I., Murphy, J., Marshall, D.S., Cowger, C. 2008. Virulence structure of the eastern U.S. wheat powdery mildew population. *Plant Disease*. 92:1074-1082.
- Wooten, D., Livingston, D.P., Lyerly, J., Holland, J.B., Jellen, E., Marshall, D.S., Murphy, J. 2008. Quantitative trait loci and epistasis for crown freezing tolerance in the Kanota x Ogle hexaploid oat mapping population.. *Crop Science*. 48:149-157.
- Wooten, D.R., Livingston, D.P., Lyerly, J., Holland, J.B., Fellon, E.N., Marshall, D.S., Murphy, P.J. 2008. Quantitative Trait Loci and Epistasis for Oat Winter Hardiness Component Traits. *Crop Science*. 48:149-157.
- Cowger, C., Murphy, P. 2007. Artificial Inoculation of Wheat for Selecting Resistance to *Stagonospora Nodorum* Blotch. *Plant Disease*. 91:539-545.
- Costs, J.M., Griffey, C.A., Bockelman, H.E., Cambron, S.E., Cooper, A., Graybosch, R.A., Grybauskas, A., Kratochvil, R.J., Jin, Y., Marshall, D.S. 2007. Registration of 'Chesapeake' Wheat. *Journal of Plant Registrations*. doi: 10.3198/jpr2006.10.0630crc.
- Kirigwi, F.M., Van Ginkel, M., Brown Guedira, G.L., Gill, B.S., Paulsen, G., Fritz, A.K. 2007. Markers associated with a qtl for grain yield in wheat under drought. *Molecular Breeding*. 20(4):401-403.
- Miranda, L., Murphy, J., Marshall, D.S., Cowger, C., Leath, S. 2007. Chromosomal location of Pm35, a novel *Aegilops tauschii* derived powdery mildew resistance gene introgressed into common wheat (*Triticum aestivum* L.). *Theoretical and Applied Genetics*. 114:1451-1456.
- Miranda, L.M., Perugini, L., Srnic, G., Brown Guedira, G.L., Marshall, D.S., Leath, S., Murphy, J.P. 2007. Genetic Mapping of a *Triticum monococcum*-derived Powdery Mildew Resistance Gene in Common Wheat. *Crop Science*. 47:2323-2329.
- Murphy, J.P., Navarro, R.A., Marshall, D.S., Cowger, C., Cox, T., Kolmer, J.A., Leath, S., Gaines, C.S. 2007. Registration of NC06BGTAG12, and NC06BGTAG13 Powdery Mildew Resistant Wheat Germplasm.. *Journal of Plant Registrations*. 1:75-77.
- Perugini, L., Murphy, J., Marshall, D.S., Brown Guedira, G.L. 2007. Pm37, a new broadly effective powdery mildew resistance gene from *Triticum timopheevii* . *Journal of Theoretical and Applied Genetics*. 116:417-425.
- Tucker, D.M., Griffey, C.A., Liu, S., Brown Guedira, G.L., Marshall, D.S., Saghai-Maroo, M. 2007. Confirmation of Three Quantitative Trait Loci Conferring Adult Plant Resistance to Powdery Mildew in Two Winter Wheat Populations. *Euphytica*. 155:1-13.
- Verges, V.L., Van Sanford, D., Brown Guedira, G.L., Tekonry, D. 2007. Heritability estimates and response to selection for fusarium head blight resistance in segregating populations of soft red winter wheat . *Crop Science*.
- Wooten, D.R., Livingston, D.P., Jellen, E.N., Boren, K.J., Marshall, D.S., Murphy, J. 2007. An intergenomic reciprocal translocation associated with oat winterhardiness. *Crop Science*. 47:1832-1840.
- Engle, J.S., Lipps, P.E. 2006. Distribution and pathogenic characterization of *pyrenophora tritici-repentis* and *stagonospora nodorum* in ohio. *Biological and Cultural Tests for Control of Plant Diseases*.
- Engle, J.S., Lipps, P.E., Minyo, R.J. 2006. Reaction of commercial soft red winter wheat cultivars to *stagonospora nodorum* in the greenhouse and field. *Plant Disease*.
- Engle, J.S., Madden, L.V., Lipps, P.E. 2006. Distribution and pathogenic characterization of *pyrenophora tritici-repentis* and *stagonospora nodorum* in ohio. *Phytopathology* 1355-1362.

- Miranda, L.M., Murphy, J.P., Marshall, D.S., Leath, S. 2006. Pm34, a new powdery mildew resistance gene transferred from *aegilops tauschii* coss. into common wheat (*triticum aestivum* l.). *Theoretical and Applied Genetics*. 113:1497-1504.
- Olien, C.R., Livingston, D.P. 2006. Understanding freeze stress in biological tissues: thermodynamics of interfacial water. *Thermochimica Acta*. 451:9-13.

6657-21000-006-00D

GENETIC AND CULTURAL PRACTICE IMPROVEMENT FOR SUSTAINABLE COTTON PRODUCTION – Philip Bauer (P) and Benjamin Campbell; Florence, South Carolina

- Campbell, B.T., Saha, S., Jenkins, J.N., Park, W., Percy, R.G., Frelichowski, J.E., Mayee, C., Gotmare, V., Dessauw, D., Giband, M., Du, X., Jia, Y., Constable, G., Dillon, S., Abdurakhmonov, I.Y., Abdurakarimov, A., Rizaeva, S.M., Adullaev, A.A., Barroso, P.V., Padua, J.G., Hoffman, L.V., Podolnaya, L. 2010. Status of the global cotton germplasm resources. *Crop Science*. 50:1161-1179.
- Mi, X., Eskridge, K., Wang, D., Baenziger, P.S., Campbell, B.T., Gill, K.S., Dweikat, I., Bovaird, J. 2010. Regression-based multi-trait QTL mapping using a structural equation model. *Genetics Research*. 9:Article 38, 21 pp.
- Mi, X., Eskridge, K.M., Wang, D., Baenziger, P.S., Campbell, B.T., Gill, K.S., Dweikat, I. 2010. Bayesian mixture structural equation modelling in multiple-trait QTL mapping. *Genetics Research*. 92:239-250.
- Bauer, P.J., Park, D.M., Campbell, B.T. 2009. Cotton production in rotation with summer legumes. *Journal of Cotton Science*. 13:183-188.

6659-21000-016-00D

GENETIC IMPROVEMENT OF SOUTHERNPEAS AND PEPPERS – Richard Fery (P); Charleston, South Carolina

- Fery, R.L., Thies, J.A. 2011. 'Truhart-NR', A Root-knot Nematode Resistant, Pimento-type Pepper. *HortScience*. 46(5):815-816.
- Fery, R.L., Thies, J.A. 2011. PA-566, A Root-knot Nematode Resistant, Pimento-type Pepper. *HortScience*. 46(4):668-669.
- Fery, R.L., Thies, J.A. 2010. PA-559, a Root-knot Nematode Resistant, Red-fruited, Habanero-type Pepper. *HortScience*. 45(5):822-823.
- Fery, R.L. 2009. 'WhiteAcre-DG', a Small-seeded, Cream-type Southernpea with an Enhanced Persistent Green Seed Phenotype. *HortScience*. 44(5):1472-1473.
- Fery, R.L. 2009. 'ZipperCream-GC', a Large-seeded, Cream-type Southernpea with a Green Cotyledon Phenotype. *HortScience*. 44(5):1474-1475.
- Fery, R.L. 2007. 'GreenPack-DG', a Pinkeye-type Southernpea with an Enhanced Persistent Green Seed Phenotype. *HortScience*. 42:692-693.
- Fery, R.L., Buckley, B., Marsh, D.B. 2007. 'WhipperSnapper', a Dual-purpose Southernpea for the Production of Both Snaps and Fresh-shell Peas. *HortScience*. 42:1297-1298.
- Fery, R.L., Thies, J.A. 2007. 'TigerPaw-NR', a Root-Knot Nematode Resistant, Habanero-type Pepper. *HortScience*. 42(7):1721-1722.

6659-21000-017-00D

GENETICS AND GENETIC IMPROVEMENT OF DISEASE RESISTANCE AND QUALITY TRAITS IN WATERMELON, BROCCOLI, AND LEAFY GREEN BRASSICAS – Mark Farnham (P) and Amnon Levi; Chaleston, South Carolina

- Farnham, M.W. 2010. Glossy and Nonglossy Near-isogenic Lines USVL115-GL, USVL115-NG, USVL188-GL and USVL188-NG of Broccoli Derived from Doubled Haploids. *HortScience*. 45:660-662.
- Levi, A., Harris-Shultz, K.R., Wechter, W.P., Kousik, C.S., Thies, J.A. 2010. DNA markers and pollen morphology reveal that *Praecitrullus fistulosus* is more closely related to *Benincasa hispida* than to *Citrullus* spp.. *Genetic Resources and Crop Evolution*. 135:369-378.
- Nimmakayala, P., Jeong, J., Tomason, Y., Levi, A., Perumal, R., Reddy, U.K. 2010. Molecular Phylogeny of *Citrullus* Species as Inferred from AFLPs and SSRs. *Plant Genetic Resources*. 8:16-25.
- Farnham, M.W., Kopsell, D.A. 2009. Importance of Genotype on Carotenoid and Chlorophyll Levels in Broccoli Heads. *HortScience*. 44:1248-1253.
- Harris, K.R., Ling, K., Levi, A., Wechter, W.P. 2009. Identification and Utility of Markers Linked to the Zucchini Yellow Mosaic Virus Resistance Gene in Watermelon. *Journal of the American Society for Horticultural Science*. 134:529-534.

- Harris, K.R., Wechter, W.P., Levi, A. 2009. Isolation, Sequence Analysis, and Linkage Mapping of NBS-LRR Disease Resistance Gene Homologs in Watermelon. *Journal of the American Society for Horticultural Science*. 134:649-657.
- Iniguez-Luy, F.L., Lukens, L., Farnham, M.W., Amasino, R.M., Osborn, T.C. 2009. Development of Public Immortal Mapping Populations, Molecular Markers and Linkage Maps for Rapid Cycling Brassica rapa and B. oleracea. *Theoretical and Applied Genetics*. 120:31-43.
- Levi, A., Thies, J.A., Ling, K., Simmons, A.M., Kousik, C.S., Hassell, R. 2009. Genetic Diversity of Lagenaria Siceraria and Identification of Accessions Useful for the Development of Disease and Pest Resistant Rootstocks. *Diversity Plant Genetic Resources Journal*, <http://journals.cambridge.org/action/displayIssue?jid=PGR>. doi: 10.1017/S1479262109225354.
- Levi, A., Wechter, W.P., Harris-Shultz, K.R., Davis, A.R., Fie, Z. High-frequency Oligonucleotides in Watermelon Expressed Sequenced Tag-unigenes Are Useful in Producing Polymorphic Polymerase Chain Reaction Markers among Watermelon Genotypes. *Journal of the American Society for Horticultural Science*. 135(4):369-378.
- Nimmakayala, P., Tomason, Y.R., Jeong, J., Vajja, G., Levi, A., Gibson, P., Reddy, U.K. 2009. Molecular Diversity in Ukrainian Melon Collection as Revealed by AFLP and Microsatellite Markers. *Plant Genetic Resources*, online publication <http://journals.cambridge.org/action/displayIssue?jid=PGR>. doi:10.1017/S1479262108098481.
- Levi, A., Wechter, W.P., Davis, A.R. 2009. EST-PCR Markers Representing Watermelon Fruit Genes are Polymorphic among Watermelon Heirloom Cultivars Sharing a Narrow Genetic Base. *Plant Genetic Resources*. 7:16-32.
- Farnham, M.W., Davis, E.H., Morgan, J.T., Smith, J.P. 2008. Neglected Landraces of Collard (Brassica oleracea L.) from the Carolinas (USA). *Genetic Resources and Crop Evolution*. 55:797-801.
- Salman, A., Levi, A., Wolf, S., Trebitsh, T. 2008. Acc Synthase Genes are Polymorphic in Watermelon (citrullus spp.) and Differentially Expressed in Flowers and in Response to Auxin and Gibberellin. *Plant And Cell Physiology*. 49:740-750.
- Gao, M., Li, G., Yang, B., Qiu, D., Farnham, M.W., Quiros, C. 2007. High Density Brassica Oleracea Linkage Map: Identification of Useful New Linkages. *Theoretical and Applied Genetics*. 115:277-287.
- Hale, A.L., Farnham, M.W., Nzaramba, M.N., Kimbeng, C.A. 2007. Heterosis for Horticultural Traits in Broccoli. *Theoretical and Applied Genetics*. 115:351-360.
- Joobeur, T., Gusmini, G., Zhang, X., Levi, A., Yong, X., Wehner, T., Oliver, M., Dean, R. 2006. Construction of a watermelon BAC library and identification of SSRs anchored to melon or Arabidopsis genomes. *Theoretical and Applied Genetics*. 112:1553-62.

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APPENDIX 3

National Program 301 Plant Genetic Resources, Genomics, and Genetics Improvement

ACCOMPLISHMENT REPORT 2006 – 2011

ARS Variety and Germplasm Public Releases*

545 Total Releases

<i>Cucurbits (11 Releases)</i>	<i>Oil Crops (65 Releases)</i>
<i>Fiber Crops (41 Releases)</i>	<i>Small Fruit Crops (29 Releases)</i>
<i>Floriculture and Nursery Crops (25 Releases)</i>	<i>Sugar Crops (52 Releases)</i>
<i>Forage Crops (46 Releases)</i>	<i>Tree Fruits (28 Releases)</i>
<i>Grain Crops (128 Releases)</i>	<i>Tree Nuts (4 Releases)</i>
<i>Legumes (47 Releases)</i>	<i>Vegetables (69 Releases)</i>

CUCURBITS (11 Releases)

- Cucumber:** EOM 402-10 High B-Carotene Cucumber, released by ARS scientists in Logan, Utah, FY 2011
- Cucumber:** European Long Greenhouse Cucumber Inbred Backcross Line Population, released by ARS scientists in Logan, Utah, FY 2011
- Cucumber:** USDA, ARS Beit Alpha Cucumber Inbred Backcross Line Population, released by ARS scientists in Logan, Utah, FY 2011
- Cucumber:** USDA, ARS Cucumis Hystrix-Derived U.S. Processing Cucumber Inbred Backcross Line Population, released by ARS scientists in Logan, Utah, FY 2011
- Melon:** USDA 846-1 Fractal Melon, released by ARS scientists in Logan, Utah, FY 2010
- Watermelon:** PI 525088-PMR: A Watermelon Line Useful for Introducing Race 1 Powdery Mildew Resistance into Watermelon Lines, released by ARS scientists in Lane, Oklahoma, FY 2006
- Watermelon:** USVL-220, Watermelon, released by ARS scientists in Charleston, South Carolina, FY 2006
- Watermelon:** LSW-177, Watermelon, released by ARS scientists in Lane, Oklahoma, FY 2007
- Watermelon:** LSW-194, Watermelon, released by ARS scientists in Lane, Oklahoma, FY 2007
- Watermelon:** MSW-28, Watermelon, released by ARS scientists in Lane, Oklahoma, FY 2007
- Watermelon:** USVL-230 Watermelon, released by ARS scientists in Charleston, South Carolina, FY 2009

* A number of these Releases are collaboratively released with our University and Industry partners

FIBER CROPS (41 Releases)

- Cotton:** SJ-U86, one upland cotton germplasm line, released by ARS scientists in Shafter, California, FY 2006
- Cotton:** NMRIL recombinant inbred line population of upland cotton, released by ARS scientists in Maricopa, Arizona, FY 2006
- Cotton:** MS-01RKN, MS-24RKN, MS-30RKN, MS-33RKN, MS-35RKN, and MS-37RKN, root-knot nematode resistant upland cotton germplasm lines, released by ARS scientists in Mississippi State, Mississippi, FY 2006
- Cotton:** Lonren-1, Cotton, released by ARS scientists in College Station, Texas, FY 2007
- Cotton:** Lonren-2, Cotton, released by ARS scientists in College Station, Texas, FY 2007
- Cotton:** RMUP-C5, Random Mated Population of Upland Cotton Germplasm, released by ARS scientists in Mississippi State, Mississippi, FY 2007
- Cotton:** PD 98066 and PD 99035, Cotton, released by ARS scientists in Florence, South Carolina, FY 2007
- Cotton:** SJ-07P-FR01, SJ-07P-FR02, SJ-07P-FR03, and SJ-07P-FR04, released by ARS scientists in Shafter, California, FY 2007
- Cotton:** SP156, SP177, SP179, SP205, SP225, released by ARS scientists in Stoneville, Mississippi, FY 2008
- Cotton:** PSI 425 and PSI 113 Pima Cotton, released by ARS scientists in College Station, Texas, FY 2008
- Cotton:** JC Cotton, released by ARS scientists in Stoneville, Mississippi, FY 2009
- Cotton:** SCM3-4-3, SCM3-7-3, RM3-8-1, EM4-3-1, Cotton, released by ARS scientists in Stoneville, Mississippi, FY 2009
- Cotton:** CRB 252 Upland Cotton, released by ARS scientists in College Station, Texas, FY 2009
- Cotton:** GVS1, GVS2, GVS3, GVS4, GVS5, Upland Cotton, released by ARS scientists in Stoneville, Mississippi, FY 2009
- Cotton:** MT2468 REN1, MT2468 REN2, and MT2468 REN3, released by ARS scientists in Mississippi State, Mississippi, FY 2009
- Cotton:** MD9NE, MD25, Cotton, released by ARS scientists in Stoneville, Mississippi, FY 2010
- Cotton:** GA 120R1B3 Germplasm Line of Cotton, released by ARS scientists in Tifton, Georgia, FY 2010

FLORICULTURE AND NURSERY CROPS (25 Releases)

- Beautyberry:** Duet, Beautyberry, released by ARS scientists in Washington, DC, FY 2006
- Black Chokeberry:** McKenzie, released by ARS scientists in Bismarck, North Dakota, FY 2008
- Camellia:** Anacostia, Camellia, released by ARS scientists in Washington, DC, FY 2010
- Chinese Hibiscus:** USS Arizona, Chinese Hibiscus, released by ARS scientists in Poplarville, Mississippi, FY 2010
- Chinese Hibiscus:** USS California, Chinese Hibiscus, released by ARS scientists in Poplarville, Mississippi, FY 2010
- Damiana:** Luisa, Damiana, released by ARS scientists in Miami, Florida, FY 2011
- Fringe Flower:** Snow Panda, Fringe Flower, released by ARS scientists in Washington, DC, FY 2011
- Hackberry:** Prairie Harvest, released by ARS scientists in Bismarck, North Dakota, FY 2009

- Halberd-Leaved Rose Mallow:** Lufkin White, Halberd-Leaved Rose Mallow, Ornamental Hibiscus, released by ARS scientists in Poplarville, Mississippi, FY 2011
- Halberd-Leaved Rose Mallow:** Lufkin Red Hibiscus, Halberd-Leaved Rose Mallow, Ornamental Hibiscus, released by ARS scientists in Poplarville, Mississippi, FY 2011
- Hibiscus:** Sahara Sunset Red Leaf Hibiscus, released by ARS scientists in Poplarville, Mississippi, FY 2008
- Japanese Snowbell:** Japanese Snowbell (NA 71587) 'Spring Showers,' released by ARS scientists in McMinnville, Tennessee, FY 2010
- Lilac:** Old Glory, Lilac, released by ARS scientists in Washington, DC, FY 2006
- Lilac:** Declaration, Lilac, released by ARS scientists in Washington, DC, FY 2006
- Oakleaf Hydrangea:** Ruby Slippers, Oakleaf Hydrangea, released by ARS scientists in McMinnville, Tennessee, FY 2010
- Oakleaf Hydrangea:** Muchkin, Oakleaf Hydrangea, released by ARS scientists in McMinnville, Tennessee, FY 2010
- 'Ohelo Berry:** Nene (N06-6) 'Ohelo Berry, released by ARS scientists in, Hilo, Hawaii, FY 2008
- 'Ohelo Berry:** Kilauea (N06-7) 'Ohelo Berry, released by ARS scientists in Hilo, Hawaii, FY 2008
- 'Ohelo Berry:** Red Button (N06-9) 'Ohelo Berry, released by ARS scientists in Hilo, Hawaii, FY 2008
- Ornamental Ginger:** Ramata, Ornamental Ginger, released by ARS scientists in Poplarville, Mississippi, FY 2010
- Petunia:** USNA-76202, Petunia, released by ARS scientists in Washington, DC, FY 2007
- Tecoma:** Tangelo, Tecoma, released by ARS scientists in Miami, Florida, FY 2008
- Tecoma:** Miami Sunrise, Tecoma, released by ARS scientists in Miami, Florida, FY 2008
- Tecoma:** Miami Sunset, Tecoma, released by ARS scientists in Miami, Florida, FY 2008
- Viburnum:** Nantucket, released by ARS scientists in Washington, DC, FY 2008

FORAGE CROPS (43 Releases)

- Alfalfa:** IAFAL-C3, Yellow Flowered Alfalfa, released by ARS scientists in Madison, Wisconsin, FY 2007
- Alfalfa:** Falcata, Alfalfa, released by ARS scientists in Logan, Utah, FY 2008
- Bahiagrass:** Tifquik, Bahiagrass germplasm, released by ARS scientists in Tifton, Georgia, FY 2006
- Bahiagrass:** T18 and T23, Bahiagrass, released by ARS scientists in Tifton, Georgia, FY 2007
- Basalt Milkvetch:** NBR-1 Germplasm, released by ARS scientists in Logan, Utah, FY 2008
- Basin Wildrye:** Continental, Basin Wildrye, released by ARS scientists in Logan, Utah, FY 2008
- Birdsfoot Trefoil:** Witt, Birdsfoot Trefoil, released by ARS scientists in Madison, Wisconsin, FY 2007
- Bottlebrush Squirreltail:** Rattlesnake Germplasm, released by ARS scientists in Logan, Utah, FY 2006
- Bottlebrush Squirreltail:** Antelope Creek, Bottlebrush Squirreltail, released by ARS scientists in Logan, Utah, FY 2009
- Bottlebrush Squirreltail:** Pleasant Valley, Bottlebrush Squirreltail, released by ARS scientists in Logan, Utah, FY 2009
- Crested Wheatgrass:** Hycrest II, Crested Wheatgrass, released by ARS scientists in Logan, Utah, FY 2008

Crested Wheatgrass: Vavilov II, Crested Wheatgrass, released by ARS scientists in Logan, Utah, FY 2008

Dallisgrass: Sabine Dallisgrass, released by ARS scientists in College Station, Texas, FY 2008

Eastern Gamagrass: Meadowcrest, released by ARS scientists in Woodward, Oklahoma, FY 2006

Forage Soybean: Greencastle Forage Soybean, released by ARS scientists in Beltsville, Maryland, FY 2008

Green Needlegrass: Fowler Germplasm, Green Needlegrass, released by ARS scientists in Logan, Utah, FY 2006

Hairy Vetch: Purple Bounty, Hairy Vetch, released by ARS scientists in Beltsville, Maryland, FY 2006

Hairy Vetch: Purple Prosperity, Hairy Vetch, released by ARS scientists in Beltsville, Maryland, FY 2008

Indian Ricegrass: White River Germplasm, released by ARS scientists in Logan, Utah, FY 2006

Indiangrass: Warrior Indiangrass, released by ARS scientists in Lincoln, Nebraska, FY 2008

Indiangrass: Scout Indiangrass, released by ARS scientists in Lincoln, Nebraska, FY 2008

Indiangrass: Chief Indiangrass, released by ARS scientists in Lincoln, Nebraska, FY 2008

Kura Clover: Kura1, Kura Clover, released by ARS scientists in Madison, Wisconsin, FY 2006

Ryegrass: Floregon Italian Ryegrass (*Lolium multiflorum*) with resistance to Crown Rust, released by ARS scientists in Corvallis, Oregon, FY 2006

Shattercane: N614, A3N615, N616, N617, Shattercane, released by ARS scientists in Lincoln, Nebraska, FY 2011

Snake River Wheatgrass: Discovery, Snake River Wheatgrass, released by ARS scientists in Logan, Utah, FY 2008

Switchgrass: Bomaster, Switchgrass, released by ARS scientists in Raleigh, North Carolina, FY 2006

Switchgrass: Performer, Switchgrass, released by ARS scientists in Raleigh, North Carolina, FY 2006

Switchgrass: Tem-Lodorm, Switchgrass, released by ARS scientists in College Station, Texas, FY 2007

Switchgrass: Colony, released by ARS scientists in Raleigh, North Carolina, FY 2009

Switchgrass: ALB280 and ALB5B, Switchgrass, released by ARS scientists in Albany, California, FY 2010

Western Prairie Clover: Majestic Germplasm, released by ARS scientists in Logan, Utah, FY 2010

Western Prairie Clover: Spectrum Germplasm, released by ARS scientists in Logan, Utah, FY 2010

Wheatgrass: Slender Wheatgrass (Firststrike), released by ARS scientists in Logan, Utah, FY 2006

Wheatgrass: DGE-1, Durum / Wheatgrass Disomic Addition Line, released by ARS scientists in Fargo, North Dakota, FY 2006

Wheatgrass: Manifest Intermediate Wheatgrass, released by ARS scientists in Fort Collins, Colorado, FY 2007

Wheatgrass: Recovery Western Wheatgrass, released by ARS scientists in Logan, Utah, FY 2009

Wildrye: Homestead Canada Wildrye, released by ARS scientists in Lincoln, Nebraska, FY 2006

GRAIN CROPS (131 Releases)

Barley: Sidney and Stoneham, Russian Wheat Aphid-Resistant Two-Rowed Drought Hardy Spring Feed Barley Varieties, released by ARS scientists in Stillwater, Oklahoma, FY 2006

Barley: Herald Barley, released by ARS scientists in Aberdeen, Idaho, FY 2006

- Barley:** STARS 0601B, STARS 0602B, STARS 0603B, STARS 0604B, STARS 0605B, STARS 0606B, STARS 0607B, STARS 0608B, STARS 0609B, STARS 0610B, STARS 0611B, STARS 0612B, STARS 0613B, STARS 0614B, STARS 0615B, STARS 0616B, STARS 0617B, STARS 0618B, and STARS 0619B, nineteen spring, 6-rowed, malting barley germplasm, released by ARS scientists in Stillwater, Oklahoma, FY 2006
- Barley:** STARS 0620B, STARS 0621B, STARS 0622B, STARS 0623B, STARS 0624B, STARS 0625B, STARS 0626B, STARS 0627B, STARS 0628B, STARS 0629B, STARS 0630B, STARS 0631B, STARS 0632B, STARS 0633B, STARS 0634B, STARS 0635B, and STARS 0636B seventeen spring, 2-rowed, malting barley germplasm, released by ARS scientists in Stillwater, Oklahoma, FY 2006
- Barley:** STARS 0637B, STARS 0638B, STARS 0639B, STARS 0640B, STARS 0641B, STARS 0642B, and STARS 0643B, seven spring, 2-rowed, feed barley germplasm, released by ARS scientists in Stillwater, Oklahoma, FY 2006
- Barley:** Tetonia, Barley, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Barley:** A Set of Four Low-Phytate and Wild Type Sib Pairs, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Barley:** 95SR316A Barley, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Barley:** RWA 1758, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Barley:** Clearwater, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Barley:** Lenetah, released by ARS scientists in Aberdeen, Idaho, FY 2008
- Barley:** Endeavor Barley, released by ARS scientists in Aberdeen, Idaho, FY 2009
- Barley:** LP1-2581; LP640-1304; LP3-1159; LP1-2163H, Barley, released by ARS scientists in Aberdeen, Idaho, FY 2010
- Barley:** 03AH3054-51, Barley, released by ARS scientists in Aberdeen, Idaho, FY 2010
- Barley:** 03AH6561-94, Barley, released by ARS scientists in Aberdeen, Idaho, FY 2010
- Barley:** STARS 1006B, STARS 1007B, STARS 1008B, STARS 1009B, STARS 1010B, STARS 1011B, STARS 1012B, and STARS 1013B, eight winter, six-rowed, feed barley (*Hordeum vulgare* L.) germplasm lines, released by ARS scientists in Stillwater, Oklahoma, FY 2010
- Corn:** CRW3(S1)C6, released by ARS scientists in Columbia, Missouri, FY 2007
- Corn:** TZAR101, 102, 103, 104, 105, 105, Corn, released by ARS scientists in New Orleans, Louisiana, FY 2008
- Corn:** MP 718 and MP 719, released by ARS scientists in Mississippi State, Mississippi, FY 2011
- Hop:** Teamaker, Hop, released by ARS scientists in Corvallis, Oregon, FY 2006
- Hop:** Kazak 2000, Hop, released by ARS scientists in Corvallis, Oregon, FY 2008
- Hop:** Mt. Rainier, Hop, released by ARS scientists in Corvallis, Oregon, FY 2009
- Rice:** Indica-14 and -15 Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2006
- Rice:** Indica-16,-17,-18 Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2007
- Rice:** M-202/IR50 (MI) Recombinant Inbred Mapping Populations, released by ARS scientists in Davis, CA, FY 2006
- Rice:** TIL:455, Rice, released by ARS scientists in Beaumont, Texas, FY 2006
- Rice:** TIL:514, Rice, released by ARS scientists in Beaumont, Texas, FY 2006
- Rice:** TIL:642, Rice, released by ARS scientists in Beaumont, Texas, FY 2006

- Rice:** RU9101001/KATY, released by ARS scientists in Stuttgart, Arkansas, FY 2007
- Rice:** Deletion Mutant Population of Katy Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2007
- Rice:** Presidio Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2008
- Rice:** Deltabelle Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2009
- Rice:** Rondo Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2009
- Rice:** JES Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2009
- Rice:** RIL103, Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2010
- Rice:** RIL158, Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2010
- Rice:** RIL186, Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2010
- Rice:** RIL220, Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2010
- Rice:** RIL221, Rice, released by ARS scientists in Stuttgart, Arkansas, FY 2010
- Rice:** M-105, Rice Cultivar, released by ARS scientists in Davis, California, FY 2011
- Sorghum:** Sorghum Genetic Stocks: BN611, A/BN612, RN613, released by ARS scientists in Lincoln, Nebraska, FY 2007
- Spring Oat:** Maverick, Spring Oat, released by ARS scientists in Aberdeen, Idaho, FY 2006
- Spring Oat:** Monico, Spring Oat, released by ARS scientists in Aberdeen, Idaho, FY 2006
- Waxy Proso Millet:** Plateau, released by ARS scientists in Lincoln, Nebraska, FY 2009
- Wheat:** ARS15144 and ARS14142, Hard Red Winter Wheat, released by ARS scientists in Pullman, Washington, FY 2008
- Wheat:** Appalachian White, Wheat, released by ARS scientists in Raleigh, North Carolina, FY 2009
- Wheat:** Camelot, Winter Wheat, released by ARS scientists in Lincoln, Nebraska, FY 2008
- Wheat:** Cara, Wheat, released by ARS scientists in Pullman, Washington, FY 2008
- Wheat:** Centerfield, Wheat, released by ARS scientists in Stillwater, Oklahoma, FY 2007
- Wheat:** Duster, Wheat, released by ARS scientists in Stillwater, Oklahoma, FY 2007
- Wheat:** Hard Kernel Puroindoline Allele Near-Isogenic Line (NIL), released by ARS scientists in Pullman, Washington, FY 2007
- Wheat:** Mace Hard Red Winter Wheat Cultivar, released by ARS scientists in Lincoln, Nebraska, FY 2008
- Wheat:** NE01481, Hard Red Winter Wheat, released by ARS scientists in Lincoln, Nebraska, FY 2010
- Wheat:** NE01643, Hard Red Winter Wheat, released by ARS scientists in Lincoln, Nebraska, FY 2007
- Wheat:** NH03614 CL, Winter Wheat, released by ARS scientists in Lincoln, Nebraska, FY 2008
- Wheat:** NI04421, Hard Red Winter Wheat, released by ARS scientists in Lincoln, Nebraska, FY 2010
- Wheat:** Anton Hard White Winter Wheat Cultivar, released by ARS scientists in Lincoln, Nebraska, FY 2008
- Wheat:** Oregon Feed Wheat #5, Winter Wheat, released by ARS scientists in Pullman, Washington, FY 2008
- Wheat:** Stars 0601W Wheat, released by ARS scientists in Stillwater, Oklahoma, FY 2007
- Wheat:** WQL7CENTL; WQL7CENTH; WQL8AWSL; WQL8DAWSH, Wheat Genetic Stocks, released by ARS scientists in Pullman, Washington, FY 2010
- Wheat:** Nueast, Wheat, released by ARS scientists in Raleigh, North Carolina, FY 2009

Wheat: Waxy-Pen, Wheat, released by ARS scientists in Pullman, Washington, FY 2006

LEGUMES (47 Releases)

Black Bean: TARS-MST1 and SB-DT1, Multiple-Stress Tolerant Black Bean Germplasm, released by ARS scientists in Mayaguez, Puerto Rico, FY 2011

Chickpea: Dylan, released by ARS scientists in Pullman, Washington, FY 2006

Chickpea: Sawyer Chickpea, released by ARS scientists in Pullman, Washington, FY 2008

Chickpea: CA0469C025C, Chickpea, released by ARS scientists in Pullman, Washington, FY 2010

Common Bean: ABC Weihing, Common Bacterial Blight, Rust and Mosaic Resistant, Semi-Upright, High Seed Quality Great Northern Bean Germplasm Line, released by ARS scientists in Lincoln, Nebraska, FY 2007

Common Bean: Verano Common Bean, released by ARS scientists in Mayaguez, Puerto Rico, FY 2008

Common Bean: NE1-06-12 Great Northern Common Bean, released by ARS scientists in Lincoln, Nebraska, FY 2008

Common Bean: Badillo, Common Bean, released by ARS scientists in Mayaguez, Puerto Rico, FY 2009

Common Bean: TARS-HT1 and TARS-HT2 Common Bean Germplasm Lines, released by ARS scientists in Mayaguez, Puerto Rico, FY 2010

Common Bean: PR0401-259 and PR0650-31, released by ARS scientists in Mayaguez, Puerto Rico, FY 2011

Common Bean: TARS MST1, TARS-MST2, TARS-MST3 and SB-DT1, released by ARS scientists in Mayaguez, Puerto Rico, FY 2010

Common Bean/Dry Bean: USPT-CBB-5 and USPT-CBB-6, released by ARS scientists in Prosser, Washington, FY 2006

Cowpea: Zippercream-GC, Southernpea (Cowpea), released by ARS scientists in Charleston, South Carolina, FY 2008

Cowpea: Whiteacre-DG, SouthernPea (Cowpea), released by ARS scientists in Charleston, South Carolina, FY 2008

Cowpea: US11-36, US11-37, and US11-38, Cowpea, released by ARS scientists in Charleston, South Carolina, FY 2009

Dry Bean: USWK-CBB-17, released by ARS scientists in Prosser, Washington, FY 2006

Dry Bean: BD1003 Cranberry Dry Bean, released by ARS scientists in Prosser, Washington, FY 2008

Dry Bean: USCR-CBB-20 Dry Bean-Cranberry Market Class, released by ARS scientists in Prosser, Washington, FY 2008

Dry Pea: Spector, Dry Pea, released by ARS scientists in Pullman, Washington, FY 2006

Kidney Bean: Silver Cloud White Kidney Bean, released by ARS scientists in Prosser, Washington, FY 2006

Lentil: Riveland, Lentil, released by ARS scientists in Albany, California, FY 2007

Lentil: Cedar, Lentil, released by ARS scientists in Pullman, Washington, FY 2008

Lentil: Shasta, Lentil, released by ARS scientists in Pullman, Washington, FY 2008

Lentil: Essex, Lentil released by ARS scientists in Pullman, Washington, FY 2009

Lentil: Morena, Spanish Brown Type Lentil, released by ARS scientists in Pullman, Washington, FY 2010

- Pea:** Medora, released by ARS scientists in Pullman, Washington, FY 2006
- Pea:** Windham, released by ARS scientists in Pullman, Washington, FY 2006
- Pea:** RIL 846-34, RIL 846-40, RIL 847-36, released by ARS scientists in Pullman, Washington, FY 2007
- Pea:** 00-5001, 00-5003, 00-5004, 00-5005, 00-5006 and 00-5007 Fusarium Wilt and Root Rot Resistant Green Pea Breeding Lines, released by ARS scientists in Prosser, Washington, FY 2010
- Pinto Bean:** Quincy, Pinto Bean, released by ARS scientists in Prosser, Washington, FY 2006
- Southernpea:** Greenpack-DG, Southernpea, released by ARS scientists in Charleston, South Carolina, FY 2006

OIL CROPS (65 Releases)

- Lesquerella:** WCL-L03, released by ARS scientists in Maricopa, Arizona, FY 2006
- Peanut:** C724-19-15, Peanut, released by ARS scientists in, Tifton, Georgia, FY 2007
- Peanut:** TIFGP-1, Peanut, released by ARS scientists in Tifton, Georgia, FY 2007
- Peanut:** Georganic, Peanut, released by ARS scientists in Tifton, Georgia, FY 2007
- Peanut:** GS-CRSP15, Peanut, released by ARS scientists in Griffin, Georgia, FY 2008
- Peanut:** Red River Runner Peanut, released by ARS scientists in Lubbock, Texas, FY 2009
- Peanut:** C724-19-25, Peanut, released by ARS scientists in Tifton, Georgia, FY 2010
- Safflower:** WSRC01, WSRC02, WSRC03, Safflower, released by ARS scientists in Pullman, Washington, FY 2007
- Soybean:** Minimax Soybean, released by ARS scientists in Beltsville, Maryland, FY 2006
- Soybean:** DS4-SCN05, released by ARS scientists in Stoneville, Mississippi, FY 2006
- Soybean:** Nitrasoy Soybean, released by ARS scientists in Raleigh, North Carolina, FY 2006
- Soybean:** Stout-RPS1K, Soybean, released by ARS scientists in Wooster, Ohio, FY 2006
- Soybean:** Prohio, Soybean, released by ARS scientists in Wooster, Ohio, FY 2006
- Soybean:** S01-9364 and S01-9391 Soybean Germplasm lines Resistant to Soybean Cyst Nematode and Seed Oil Low in Linolenic, released by ARS scientists in Columbia, Missouri, FY 2007
- Soybean:** HC99-2763, released by ARS scientists in Wooster, Ohio, FY 2007
- Soybean:** Wooster, Soybean, released by ARS scientists in Wooster, Ohio, FY 2007
- Soybean:** Glycine, Soybean, released by ARS scientists in Raleigh, North Carolina, FY 2007
- Soybean:** N8001, Soybean, released by ARS scientists in Raleigh, North Carolina, FY 2007
- Soybean:** Barc-19, Soybean, released by ARS scientists in Beltsville, Maryland, FY 2008
- Soybean:** N8101, Soybean, released by ARS scientists in Raleigh, North Carolina, FY 2008
- Soybean:** Triple Null Soybean, released by ARS scientists in Columbia, Missouri, FY 2009
- Soybean:** LG-04-6863, released by ARS scientists in Urbana, Illinois, FY 2009
- Soybean:** N6202, Soybean, released by ARS scientists in Raleigh, North Carolina, FY 2009
- Soybean:** SB-01, Soybean, released by ARS scientists in Columbia, Missouri, FY 2010
- Soybean:** DS-880, Soybean, released by ARS scientists in Stoneville, Mississippi, FY 2010
- Soybean:** LG00-6925, Soybean, released by ARS scientists in Urbana, Illinois, FY 2010

- Soybean:** LG04-6000, Soybean, released by ARS scientists in Urbana, Illinois, FY 2010
- Soybean:** N7003CN, Soybean Cultivar, released by ARS scientists in Raleigh, North Carolina, FY 2010
- Sunflower:** RS3, Sunflower Germplasm, released by ARS scientists in Fargo, North Dakota, FY 2006
- Sunflower:** RHA 461, RHA 462, RHA 463, RHA 464, Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2006
- Sunflower:** HA 465, HA 466, HA 467, RHA 468, Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2006
- Sunflower:** HA 458, HA 459, HA 460, Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2006
- Sunflower:** HA 469, RHA 470, RHA 471, Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2007
- Sunflower:** CONFSCLB1, CONFSCLB2, and CONFSCLR1 to CONFSCLR6, released by ARS scientists in, Fargo, North Dakota, FY 2007
- Sunflower:** TOCOB1, TOCOR1, TOCOR2 Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2008
- Sunflower:** RS5, Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2010
- Sunflower:** RHA 472, RHA 473, RHA 474, RHA 475, Oilseed Sunflower, released by ARS scientists in Fargo, North Dakota, FY 2011
- Vernonia:** WCL-VP1, Vernonia, released by ARS scientists in Maricopa, Arizona, FY 2006
- Vernonia:** WCL-VP2, Vernonia, released by ARS scientists in Maricopa, Arizona, FY 2007
- Vernonia:** WCL-VP3, Vernonia, released by ARS scientists in Maricopa, Arizona, FY 2007

SMALL FRUIT CROPS (29 Releases)

- Blackberry:** Wild Treasure Thornless Trailing Blackberry, released by ARS scientists in Corvallis, Oregon, FY 2006
- Blackberry:** Newberry Trailing Blackberry, released by ARS scientists in Corvallis, Oregon, FY 2009
- Blackberry:** Onyx Blackberry, released by ARS scientists in Corvallis, Oregon, FY 2010
- Blueberry:** Blue Suede Blueberry Cultivar, released by ARS scientists in Griffin, Georgia, FY 2008
- Blueberry:** Earlibirdblue, Southern Highbush Blueberry, released by ARS scientists in Poplarville, Mississippi, FY 2011
- Blueberry:** Prince Rabbiteye Blueberry, released by ARS scientists in Poplarville, Mississippi, FY 2008
- Blueberry:** TH-605, released by ARS scientists in Washington, DC, FY 2009
- Blueberry:** Pearl Blueberry, released by ARS scientists in Poplarville, Mississippi, FY 2010
- Blueberry:** Sweetheart Blueberry, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** Razz, Highbush Blueberry Cultivar, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** TH-681, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** TH-743, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** TH-754, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** TH-759, released by ARS scientists in Beltsville, Maryland, FY 2010
- Blueberry:** TH-761, released by ARS scientists in Beltsville, Maryland, FY 2010

Blueberry: TH-763, released by ARS scientists in Beltsville, Maryland, FY 2010

Blueberry: Summer Sunset, Blueberry, released by ARS scientists in Beltsville, Maryland, FY 2011

Gooseberry: Jeanne, Gooseberry Cultivar, released by ARS scientists in Corvallis, Oregon, FY 2007

Grape: Dwarf Grapevine, released by ARS scientists in Geneva, New York, FY 2006

Grape: Eudora, Muscadine Grape, released by ARS scientists in Poplarville, Mississippi, FY 2007

Grape: Sunglo, Raisin Grape Cultivar, released by ARS scientists in Parlier, California, FY 2010

Grape: Scout, Hybrid Grape Cultivar, released by ARS scientists in Geneva, New York, FY 2010

Grape: Matador, Grape, released by ARS scientists in Geneva, New York, FY 2010

Grape: Kingfisher, Grape, released by ARS scientists in Geneva, New York, FY 2010

Grape: Minotaur, Grape, released by ARS scientists in Geneva, New York, FY 2010

Grape: F10, Grape, released by ARS scientists in Parlier, California, FY 2011

Raspberry: Vintage Red Raspberry, released by ARS scientists in Corvallis, Oregon, FY 2011

Strawberry: Stolo Strawberry, released by ARS scientists in Corvallis, Oregon, FY 2007

Strawberry: Valley Red Strawberry, released by ARS scientists in Corvallis, Oregon, FY 2009

SUGAR CROPS (52 Releases)

Sugarbeet: EL53 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2006

Sugarbeet: FC723, Sugarbeet Line, released by ARS scientists in Fort Collins, Colorado, FY 2006

Sugarbeet: FC723 CMS, Sugarbeet Line, released by ARS scientists in Fort Collins, Colorado, FY 2006

Sugarbeet: EL-X1 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2006

Sugarbeet: EL-X2 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2006

Sugarbeet: EL-X3 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2006

Sugarbeet: EL-X4 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2006

Sugarbeet: FC7221 Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2008

Sugarbeet: FC7220 Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2008

Sugarbeet: C812-41 and C812-41CMS Sugarbeet Beta, released by ARS scientists in Salinas, California, FY 2008

Sugarbeet: EL54 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2008

Sugarbeet: F1017, F1018, F1019, F1020, F1021, F1022, F1023, released by ARS scientists in Fargo, North Dakota, FY 2009

Sugarbeet: EL55 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2009

Sugarbeet: SR98 Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2009

Sugarbeet: CN12-446, CN12-770, and CN72-652, Sugarbeet, released by ARS scientists in Salinas, California, FY 2009

Sugarbeet: FC1018, Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2009

Sugarbeet: FC1019, Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2009

Sugarbeet: FC1020, Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2009

Sugarbeet: FC1022, Sugarbeet, released by ARS scientists in Fort Collins, Colorado, FY 2009

- Sugarbeet:** F1024, Sugarbeet, released by ARS scientists in Fargo, North Dakota, FY 2010
- Sugarbeet:** SR98/2, Sugarbeet, released by ARS scientists in East Lansing, Michigan, FY 2011
- Sugarbeet:** F1025, F1026, and F1027, released by ARS scientists in Fargo, North Dakota, FY 2011
- Sugarcane:** HOCP 00-950 Sugarcane Cultivar, released by ARS scientists in Houma, Louisiana, FY 2007
- Sugarcane:** HOCP 91-552, released by ARS scientists in Houma, Louisiana, FY 2007
- Sugarcane:** HO 00-961, Sugarcane, released by ARS scientists in Houma, Louisiana, FY 2007
- Sugarcane:** L 79-1002, High Fiber Sugarcane Variety, released by ARS scientists in St. Gabriel, Louisiana, FY 2007
- Sugarcane:** CP 00-1101, released by ARS scientists in Canal Point, Florida, FY 2007
- Sugarcane:** CP 00-1446, released by ARS scientists in Canal Point, Florida, FY 2007
- Sugarcane:** CP 00-2180, released by ARS scientists in Canal Point, Florida, FY 2007
- Sugarcane:** CP 88-1165, released by ARS scientists in Canal Point, Florida, FY 2007
- Sugarcane:** CP 01-1372 Sugarcane, released by ARS scientists in Canal Point, Florida, FY 2008
- Sugarcane:** CPCL 97-2730 Sugarcane, released by ARS scientists in Canal Point, Florida, FY 2008
- Sugarcane:** CPCL 99-4455 Sugarcane, released by ARS scientists in Canal Point, Florida, FY 2009
- Sugarcane:** CP 03-1912, Sugarcane Cultivar, released by ARS scientists in Canal Point, Florida, FY 2010
- Sugarcane:** CPCL 00-4111, Sugarcane Cultivar, released by ARS scientists in Canal Point, Florida, FY 2010
- Sugarcane:** CPCL 02-0926, released by ARS scientists in Canal Point, Florida, FY 2011
- Sugarcane:** CPCL 02-1295, released by ARS scientists in Canal Point, Florida, FY 2011
- Sugarcane:** CPCL 95-2287, released by ARS scientists in Canal Point, Florida, FY 2011
- Sugarcane:** CP 04-1935, released by ARS scientists in Canal Point, Florida, FY 2011
- Sugarcane:** CP 04-1844, released by ARS scientists in Canal Point, Florida, FY 2011
- Sugarcane:** CP 04-1566, released by ARS scientists in Canal Point, Florida, FY 2011

TREE FRUITS (28 Releases)

- Apple:** G.778, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.890, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.214, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.222, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.707, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.87, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.969, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.210, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apple:** G.179, Apple, released by ARS scientists in Geneva, New York, FY 2010
- Apricot:** Bolaroja, Apricot, released by ARS scientists in Parlier, California, FY 2007
- Apricot:** Primarosa, Apricot, released by ARS scientists in Parlier, California, FY 2007

Apricot: TWOCOT, Apricot, released by ARS scientists in Parlier, California, FY 2011

Citrus Rootstock: US-897, released by ARS scientists in Fort Pierce, Florida, FY 2006

Citrus Rootstock: US-802, released by ARS scientists in Fort Pierce, Florida, FY 2006

Citrus Rootstock: US-942, released by ARS scientists in Fort Pierce, Florida, FY 2010

Mandarin: U.S. Early Pride, released by ARS scientists in Fort Pierce, Florida, FY 2009

Orange: US Seedless Pineapple Orange, released by ARS scientists in Fort Pierce, Florida, FY 2009

Peach: August Prince, Peach, released by ARS scientists in Byron, Georgia, FY 2006

Peach: Early August, Peach, released by ARS scientists in Byron, Georgia, FY 2006

Peach: Gulfcrimson (AP01-7), Peach, released by ARS scientists in Byron, Georgia, FY 2007

Peach: HB X OK '10', released by ARS scientists in Davis, California, FY 2009

Peach: HB X OK '32', released by ARS scientists in Davis, California, FY 2009

Peach: KV981175 Peach, released by ARS scientists in Kearneysville, West Virginia, FY 2009

Peach: KV00398, released by ARS scientists in Kearneysville, West Virginia, FY 2009

Peach: HBOK 50, Peach Rootstock, released by ARS scientists in Parlier, California, FY 2010

Pear: Sunrise, Pear Cultivar, with Superior Resistance to Fire Blight and Excellent Fruit Quality, released by ARS scientists in Kearneysville, West Virginia, FY 2006

Plum: Sharpe (FLA1-1), Plum, released by ARS scientists in Byron, Georgia, FY 2007

Plum: Orablue Plum, released by ARS scientists in Kearneysville, West Virginia, FY 2010

TREE NUTS (4 Releases)

Pecan: Lakota, Pecan, released by ARS scientists in College Station, Texas, FY 2007

Pecan: Mandan Pecan, released by ARS scientists in College Station, Texas, FY 2009

Pecan: Apalachee Pecan, released by ARS scientists in College Station, Texas, FY 2009

Pecan: Lipan, Pecan, released by ARS scientists in College Station, Texas, FY 2011

VEGETABLES (69 Releases)

Broccoli: USVL089, Broccoli, released by ARS scientists in Charleston, South Carolina, FY 2006

Broccoli: USVL115-GL, USVL115-NG, USVL188-GL, USVL188-NG, Broccoli, released by ARS scientists in Charleston, South Carolina, FY 2008

Carrot: B1111, Carrot, released by ARS scientists in Madison, Wisconsin, FY 2007

Lettuce: Lines 04-0344, 04-0350, 04-0353, 04-0363, 04-0368, 04-0375, and 04-0379, seven breeding lines of crisphead lettuce, released by ARS scientists in Salinas, California, FY 2006

Lettuce: RH05-0336, RH05-0339, and RH05-0340, F9 iceberg type lettuce breeding lines with resistance to Verticillium wilt, released by ARS scientists in Salinas, California, FY 2007

Lettuce: MU06-857, released by ARS scientists in Salinas, California, FY 2008

Lettuce: RH04-0157-3, a BLS resistant F4:5 breeding line, released by ARS scientists in Salinas, California, FY 2008

- Lettuce:** RH07-0370M, RH07-0373M, RH07-0379M, RH07-0380M, RH07-0386M, and RH07-0387M, six F6:8 breeding lines selected from RH04-0157-3 for improved horticultural characteristics, released by ARS scientists in Salinas, California, FY 2008
- Lettuce:** MU06-859 Lettuce, released by ARS scientists in Salinas, California, FY 2008
- Lettuce:** 06-831-1 and 06-833-1 Lettuce, released by ARS scientists in Salinas, California, FY 2009
- Lettuce:** 06-810-1 Lettuce, released by ARS scientists in Salinas, California, FY 2009
- Lettuce:** SM09A and SM09B Lettuce, released by ARS scientists in Salinas, California, FY 2010
- Lettuce:** RH08-0472 and RH08-0475, Lettuce, released by ARS scientists in Salinas, California, FY 2011
- Onion:** OH-1, released by ARS scientists in Madison, Wisconsin, FY 2007
- Onion:** SKI-1 A&B, released by ARS scientists in Madison, Wisconsin, FY 2007
- Onion:** B8667 A&B, released by ARS scientists in Madison, Wisconsin, FY 2007
- Pepper:** 05C113-7, Pepper, released by ARS scientists in Washington, DC, FY 2007
- Pepper:** 05C69-12, Pepper, released by ARS scientists in Washington, DC, FY 2007
- Pepper:** 05C37-3, Pepper, released by ARS scientists in Washington, DC, FY 2007
- Pepper:** 06C84, Pepper, released by ARS scientists in Washington, DC, FY 2007
- Pepper:** 05C104-4, Pepper, released by ARS scientists in Washington, DC, FY 2007
- Pepper:** Numex Las Cruces, Cayenne Pepper, released by ARS scientists in Kimberly, Idaho, FY 2011
- Pepper:** PA-559, Habanero-Type Pepper, released by ARS scientists in Charleston, South Carolina, FY 2008
- Pepper:** PA-560, Habanero-Type Pepper, released by ARS scientists in Charleston, South Carolina, FY 2009
- Pepper:** PA-566, Pimento-Type Pepper, released by ARS scientists in Charleston, South Carolina, FY 2010
- Pepper:** Tigerpaw-NR, Habanero-Type Pepper, released by ARS scientists in Charleston, South Carolina, FY 2006
- Pepper:** Truhart-NR, Pimento-Type Pepper, released by ARS scientists in Charleston, South Carolina, FY 2009
- Potato:** Blazer Russet, Potato, released by ARS scientists in Aberdeen, Idaho, FY 2006
- Potato:** Peter Wilcox, Potato, released by ARS scientists in Beltsville, Maryland, FY 2007
- Potato:** Premier Russet, Potato, A93157-6LS, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Potato:** Highland, A9045-7, Potato released by ARS scientists in Aberdeen, Idaho, FY 2007
- Potato:** Yukon Gem, NDA5507-3Y, Potato, released by ARS scientists in Aberdeen, Idaho, FY 2007
- Potato:** A84180-8, released by ARS scientists in Aberdeen, Idaho, FY 2008
- Potato:** A88338-1, released by ARS scientists in Aberdeen, Idaho, FY 2008
- Potato:** Classic Russet, Potato, released by ARS scientists in Aberdeen, Idaho, FY 2009
- Potato:** Clearwater Russet, Potato, released by ARS scientists in Aberdeen, Idaho, FY 2009
- Potato:** Alpine Russett, released by ARS scientists in Aberdeen, Idaho, FY 2009
- Potato:** M1,M2,M3,M4,and M5 Advanced Selections, Potato, released by ARS scientists in Madison, Wisconsin, FY 2011

Spinach: 03-04-9, Spinach Breeding Line, released by ARS scientists in Salinas, California, FY 2006

Spinach: 03-04-63, Spinach Breeding Line, released by ARS scientists in Salinas, California, FY 2007

Sweet Potato: Charleston Scarlet, released by ARS scientists in Charleston, South Carolina, FY 2006

Sweet Potato: W-311 and W-361, Multiple Pest-Resistant Orange-Fleshed Sweet Potato Breeding Lines, released by ARS scientists in Charleston, South Carolina, FY 2007

Sweet Potato: W-315 and W-388, Sweet Potato, released by ARS scientists in Charleston, South Carolina, FY 2007

Table Beet: TBEL1, released by ARS scientists in East Lansing, Michigan, FY 2006

APPENDIX 4

National Program 301 Plant Genetic Resources, Genomics, and Genetics Improvement

ACCOMPLISHMENT REPORT 2006 – 2011

Selected Supporting Information and Documentation for Accomplishments and Impact of NP 301 Research

RELATIONSHIP OF NATIONAL PROGRAM 301 TO THE ARS STRATEGIC PLAN:

Outputs of NP 301 research support the actionable strategies associated with the performance measure shown below from the *ARS Strategic Plan for 2006- 2011*, Strategic Goal 2, Enhance the Competitiveness and Sustainability of Rural and Farm Economies; Objective 2.2: *Increase The Efficiency Of Domestic Agricultural Production And Marketing Systems*.

Performance Measure 2.2.3: Expand, maintain, and protect our genetic resource base, increase our knowledge of genes, genomes, and biological processes, and provide economically and environmentally sound technologies that will improve the production efficiency, health, and value of the Nation's crops.

Target: New technologies developed and used by ARS customers to increase production efficiency and enhance the economic value and quality of U.S. crop production while decreasing the environmental footprint of production systems.

PLANNING AND COORDINATION – SELECTED NP 301 WORKSHOPS AND STAKEHOLDER GATHERINGS:

- NP 301 Customer and Stakeholder Workshop, Baltimore, Maryland, November 2005.
- Plant Germplasm Operations Committee Annual Meetings, Beltsville, Maryland, 2006-2011.
- National Plant Germplasm Committee Annual Meetings, 2006-2011.
- Floriculture and Nursery Research Initiative Researchers' Workshop; Portland, Oregon, June 2006 and Cleveland, Ohio, October 2009.
- ARS Wine/Grape Research Workshops; Kennewick, Washington, July 2007 and Charlottesville, Virginia, October 2010.
- ARS National Sugarcane Workshop, Houma, Louisiana, August 29-31, 2007.
- Potato Expo Research Brainstorming Session, Orlando, Florida, January 6, 2010.
- ARS – Industry Sugarbeet Meeting, Orlando, Florida, February 24, 2009.
- National HLB/ACP Research Coordination Launching Workshop, Washington, D.C., December 2 – 3, 2009.
- Citrus Strategic Research Planning Meeting, Beltsville, Maryland, March 2-3, 2010.
- Citrus Health Research Forum, Denver, Colorado, June 16-18, 2010.
- National Clean Plant Network Stakeholder Workshops; Riverdale, Maryland, May 8-9, 2007, and Davis, California, May 11-13, 2010.

- Pulse Health Initiative Workshop- Planning Meeting, Beltsville, Maryland, November 18, 2009.
- National Vegetable Crop Initiative (NVCI) strategic planning workshop, Denver, Colorado, May 12-13, 2008.
- USDA National Institute of Food and Agriculture (NIFA) Customer Workshop, Washington, D.C., April 2010.
- Crop Germplasm Committee Chairs Biennial Meetings, various locations, 2006, 2008, and 2010.
- GRIN-Global Project Technical Steering Group Meetings, Beltsville, Maryland and Ames, Iowa, 2008, 2009, 2010.
- World Cocoa Foundation-ARS Research Roundtables, quarterly, various locations, 2006-2011.
- Mars-ARS Cocoa Research Initiative Planning Workshop, 2006.
- Mars-ARS Cocoa Research Review, 2008.
- U. S.-Mexico-Canada NORGEN Genetic Resources Annual Meetings, 2007-2011.
- American Seed Research Summit (ARS co-organized), 2008.
- ARS Pacific Basin Agricultural Research Center Customer/Stakeholder Workshop, Hilo, Hawaii, 2008.

EXTERNAL AWARDS:

Peer-reviewed research grants and stakeholder awards are a useful indicator of the quality and value of the NP 301 program. From 2006 to 2011, NP 301 National Program scientists were awarded 555 grants and awards that complemented and enhanced the objectives of the NP 301 Action Plan. As you can see from Table 1 below, the largest portion of the external funding sources were from industry (47.7 percent) – evidence that the research is valued by stakeholders. University grants were the second largest number, and usually involved cooperative research projects jointly conducted with university partners. ARS scientists were also successful in competing outright for grants from other Federal sources (80 in all).

External Grant Sources of Funding for National Program 301 Projects - 2006-2011

UNIVERSITY	GOVERNMENT							INTERNATIONAL	INDUSTRY	MISC
	APHIS	ENERGY	FAS	FS	NIFA	OTHER FEDERAL	STATE/ LOCAL			
147	7	5	10	5	46	7	15	27	265	21

Table 1. Peer-reviewed research grants and awards given to NP 301 project scientists from 2006-2011.

SCIENTIST TRAINING:

The training and preparing of young scientists for careers in agriculture is critical for sustaining our modern society. Table 2 shows that from 2006-2011 NP 301 National Program scientists trained and mentored more than 2,000 postdoctoral fellows and graduate and undergraduate students. Together, the external research grants and awards, and the training of young scientists are important measures of the leadership and community stewardship that NP 301 research leaders and scientists are contributing to agriculture and related sciences.

	Postdoctoral Fellows	Graduate Students	Undergraduate Students
Total	302	394	1,400

Table 2. Students trained by NP 301 research leaders and scientists from 2006-2011.