



Rift Valley Fever – Vector Reservoirs, Inter-epidemic maintenances of the virus

**RVF Workshop – An integrated Approach to
Controlling Rift Valley Fever (RVF) in Africa and the
Middle East. Jan 27th - 29th Cairo, Egypt.**

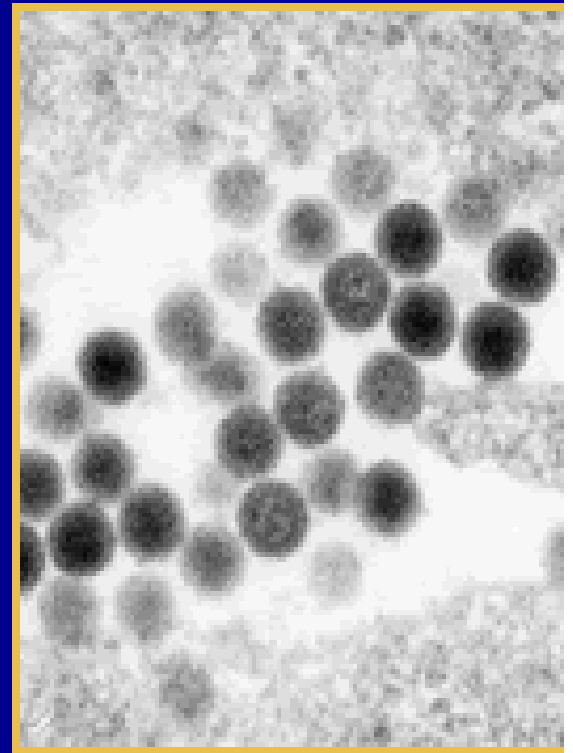
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RVF, Introduction

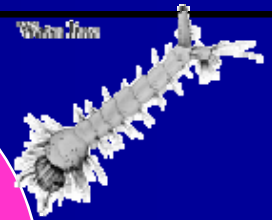
- Mosquito-borne disease caused by a Phelebovirus, Bunyaviridae family
- Affects wide range of mammals - man and domestic livestock
- Epidemics prone disease, impacting human/animal health
- Devastating economic and social consequences.



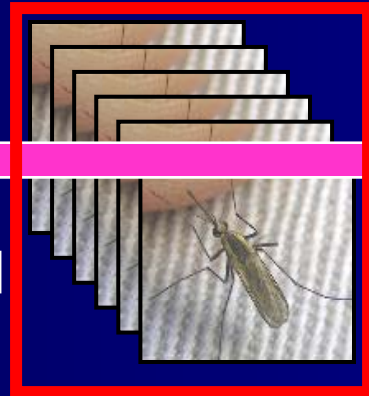
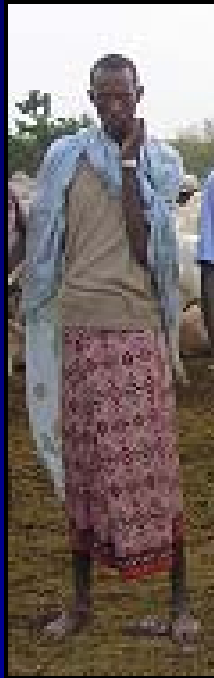
Epidemic Transmission Cycle



“dambo” with TOI eggs



1. Abnormal sustained flooding



Amplifiers



2. Large mosquito populations

OTHER ROUTES: Contact with infected animal blood, tissue & milk

RVF – vector species diversity

Genera:

Aedes

Culex,

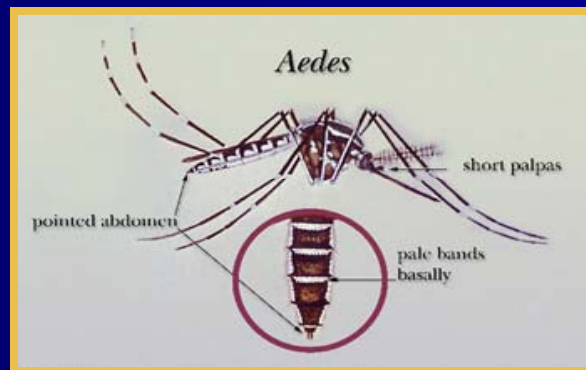
mansonia

Amblyomma

variegatum

Culicoides.

- RVFV isolated from wide range of vector species
- Experimentally, a wide variety of species can transmit RVF
- There is world wide distribution of potential vector of RVFV
- e.g Recent studies in France and Tunisia (2008) found competent vectors

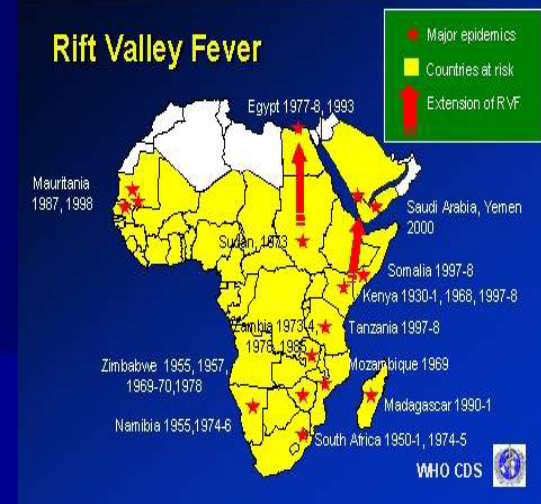


!! Hence likely spread of the virus.



VECTORS IN MAJOR OUTBREAKS

- Kenya 1950/1951 – Over 100,000 livestock dead
 - **Vectors not documented**
- Egypt – 1977 - First Major human involvement
 - **vectors *Culex pipiens*.**
- West Africa 1987 – Senegal & Mauritania
 - Irrigation project + heavy rains
 - **Vectors – *Ae. vexans*, *Ae. Ochraceous*, *Culex poicilipes***
- East Africa/Kenya 1997/98 – Kenya, Tanzania, Somalia --
 - Human and animal catastrophe
 - **Entomologic investigations NOT documented**
- Saudi/Yemen - 2000/2001
 - **Vectors – *Ae. Vexans arabiensis*, *Cu. tritaeniorhynchus***
- East Africa/Kenya 2006/07 – Kenya, Tanzania, Somalia
 - Due to earlier detection (than 97/98)
 - **Peak timing of extensive entomologic surveillance**



INTEREPIDEMIC VECTORS

Previous inter-epidemic surveys
Implicated many species
as RVF vectors in Kenya

<u><i>Aedes</i></u>	<u><i>Culex</i></u>	<u><i>Mansonia</i></u>	<u><i>Anopheles</i></u>
<i>mcintoshi</i>	<i>zombaensis</i> <i>vansomereni</i>	<i>africanus</i>	<i>pharoensis</i>
<i>circumluteolus</i>	<i>theileri</i> <i>antennatus</i>		<i>christyi</i>
<i>dentatus</i>	<i>rubinotus</i>		<i>squamosus</i>



KENYA OUTBREAK 2006/07

- In October/Nov 2006, reports of heavy persistent rains in NEP- Kenya + severe flooding
- On 22nd Dec 2007, outbreak of RVF was declared by GK.
- On 14th Dec GEIS/WRP & KEMRI surveillance team moved to Garissa to asses impact of flooding
- Entomologic survey continued through the outbreak



OBJECTIVES OF RVF OUTBREAK RESPONSE

Laboratory testing of entomologic collections from Garissa, Kilifi, Baringo and Kirinyaga was done:

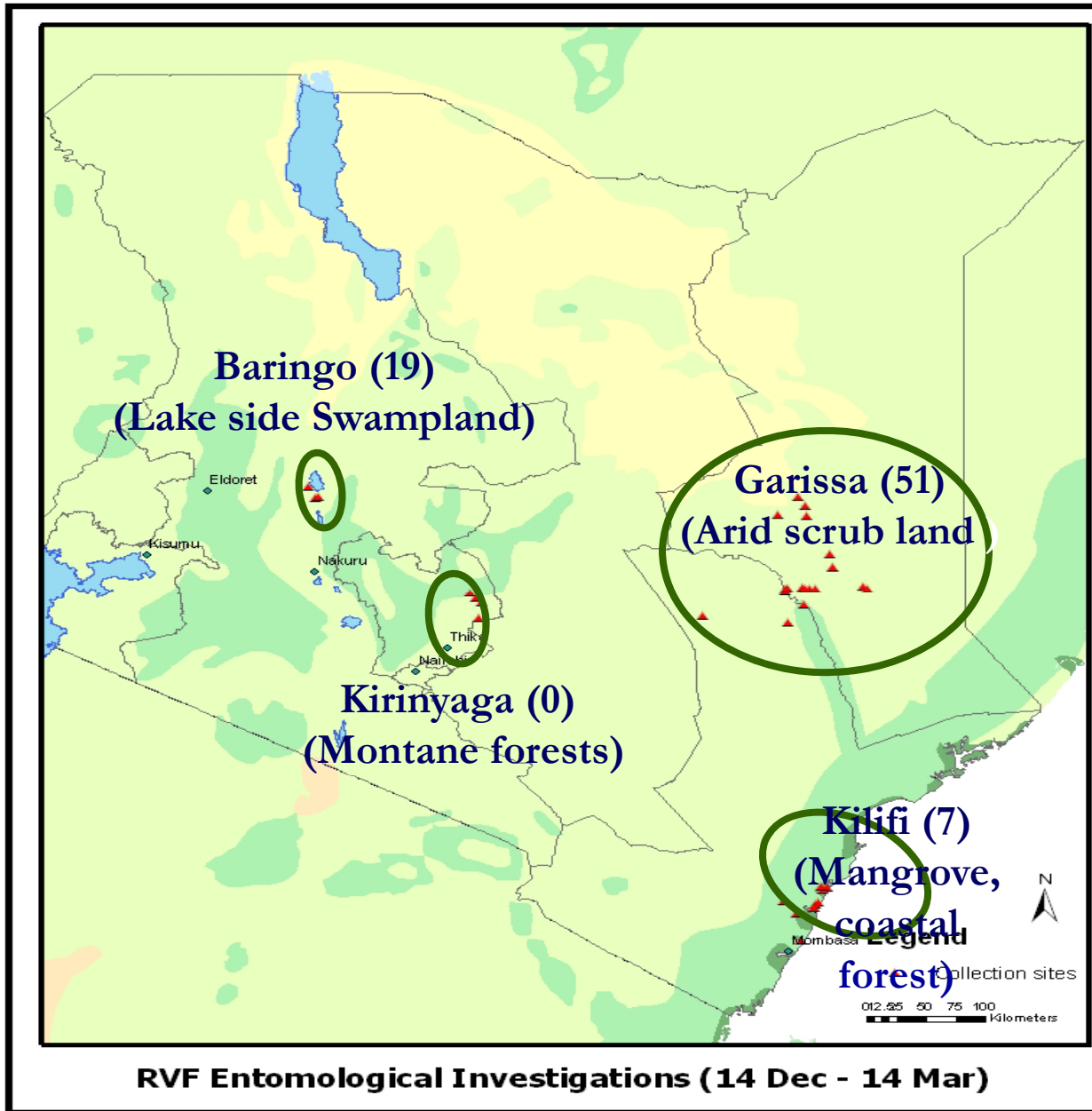
- To determine and document vector species involved in RVF outbreaks in the diverse ecologies.

Later

- To determine the competence of implicated vectors.
- Determine their host preference.



THE SAMPLES CAME FROM DIVERSE ECOLOGIES IN KENYA





Kilifi



Baringo



Garissa



Garissa

MOSQUITO COLLECTION

- Mosquitoes sampled using standard CO² baited light traps
- Set overnight around affected homes, villages, animal pens
- Trapped mosquitoes taken in the morning





MOSQUITO IDENTIFICATION

A team effort





TESTING SAMPLES FOR RVF VIRUS

Each species was grouped in pools (up to 25/pool).
Those with blood in gut were
set aside for subsequent host ID.

Pools homogenised in
medium with serum
and antibiotic supplements

RVFV RNA extracted
from
homogenates

Amplified
RVF confirmed
by sequencing PCR
product

RNA amplified by RT-PCR
(using RVF Specific primers)





TESTING - CAPACITY BUILDING



USAMRIID





PROCEDURES WITH LAB SAFETY BSL-3



Homogenisation of mosquito pools
and virus RNA extraction





PCR LABORATORY



Virus RNA amplification and detection



LAB TEST RESULTS





VIRUS DETECTION IN MOSQUITO SPECIES IN GARISSA

SPECIES	SITES	TESTED POOLS	% PROPORTION OF COLLECTION	RVF +VE POOLS
<i>Ae. mcintoshi</i>	Garissa	500	42.3	26
<i>Ae. ochraceous</i>	Garissa	450	30.0	23
<i>An. squamosus</i>	Garissa	88	5.9	2
<i>TOTAL</i>		1038	78.2%	51



VIRUS DETECTION IN MOSQUITO SPECIES FROM KILIFI

<i>Ae. pembaensis</i>	Kilifi	65	23.2	1*
<i>Cx. poicilipes</i>	Kilifi	100	38.6	3
<i>Cx. bitaeniorhynchus</i>	Kilifi	22	8.0	3*
<i>TOTAL</i>		187	69.8	7

*First time detection of RVFV in species



VIRUS DETECTION IN MOSQUITO SPECIES FROM BARINGO

<i>Cx. quinquefasciatus</i>	Baringo	75	4.0	1
<i>Cx. univittatus</i>	Baringo	8	0.3	1
<i>Ma. uniformis</i>	Baringo	804	66.8	15
<i>Ma. africanus</i>	Baringo	327	20.2	2
TOTAL		1214	91.3	19

INFECTION RATES GARISSA

SITE	RVF+ Sp	IR*	LIMITS	DENSITY
ElHumow	<i>Ae. ochraceous</i>	2.54	1.53 – 3.98	6,192
	<i>Ae. mcintoshi</i>	2.38	1.48 - 3.64	"
	<i>An squamosus</i>	1.11	0.20 – 3.64	"
Kurabul	<i>Ae. mcintoshi</i>	2.0	0.53 – 5.41	178
	<i>Ae. ochraceous</i>	1.97	0.52 – 5.33	"
Dertu/Shan	<i>Ae. ochraceous</i>	10.65	1.97 – 36.11	30
	<i>Ae. mcintoshi</i>	1.25	0.07 – 6.08	"
Desai	<i>Ae. ochraceous</i>	1.11	0.20 – 3.63	516
	<i>Ae. mcintoshi</i>	0.83	0.15 – 2.71	"

Pooled infection rates – Bias corrected maximum likelihood infection rate/1000



INFECTIONS RATES BARINGO AND KILIFI

SITES	RVF + Sp	IR	LIMITS	TOTAL
Logumgum (Baringo)	<i>Cx. Univittatus</i> *	18.01	1.32 – 118.01	5,269
"	<i>Ma. uniformis</i>	0.89	0.52 – 1.44	"
"	<i>Cx. quinquefasciatus</i>	0.71	0.04 – 3.42	"
"	<i>Ma. africana</i>	0.33	0.06 – 1.08	"
Gongoni (Kilifi)	<i>Cx. bitaeniorhyn</i>	6.92	1.84 – 18.94	63
Tezo	<i>Cx. poicilipes</i>	1.28	0.34 – 3.46	"
Uyombo	<i>Ae. pembaensis</i>	0.65	0.04 – 3.17	"



IMPLICATION



Garissa/Baringo

- 77 RVFV isolates, 51 (66%) from Garissa – 19 (24.6%) from Baringo.
- Garissa and Baringo reported highest numbers of suspected human cases
- Both pastoralist zones. Livestock are kept in large herds aids virus amplification.
- Both - flood prone terrain, & high temperatures
- suited for high mosquito densities, rapid virus growth in vectors (short EIP) and hence higher transmission rates.





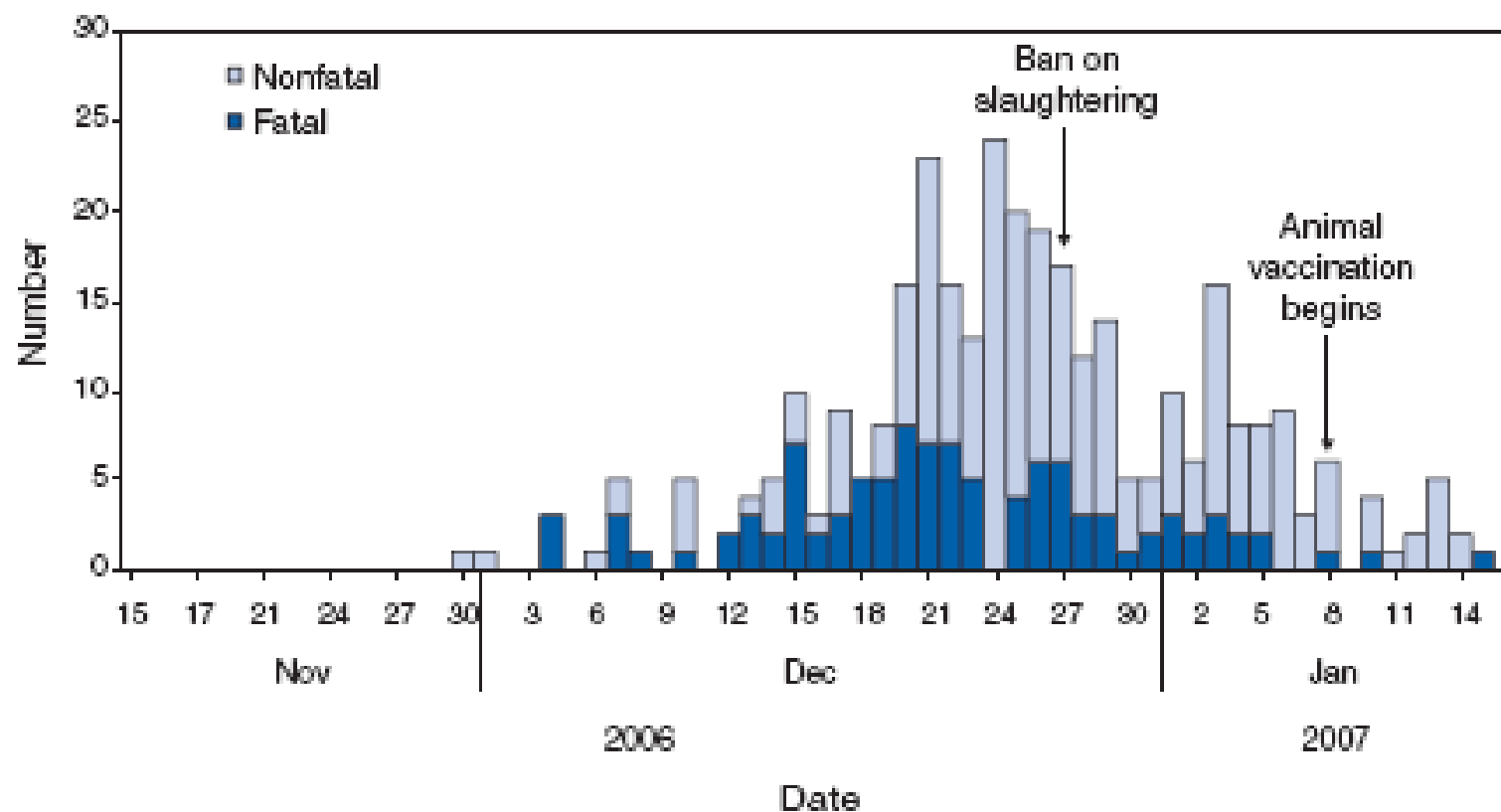


DISCUSSION

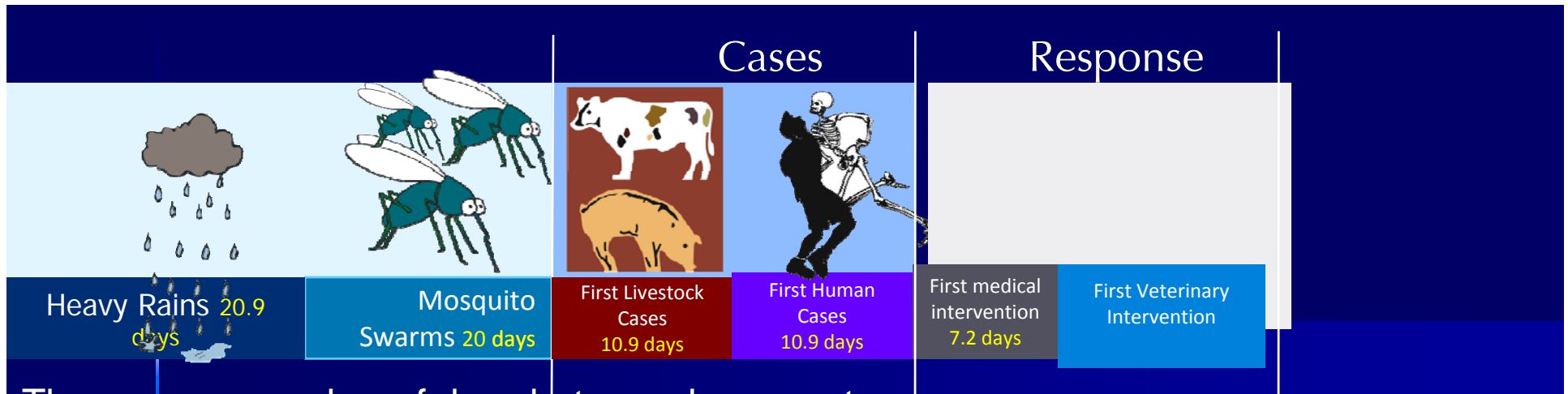
- This is one of the most comprehensive entomologic survey undertaken in a RVF outbreak in Kenya.
- Attributed to the pre-existing surveillance activities by GEIS&KEMRI
- Findings indicate that different mosquito species serve as epizootic and/or epidemic vectors of RVFV in different ecologic settings in Kenya
- Virus infected species included previously known and new ones
- This presents a complex epidemiologic pattern of RVFV in Kenya
- Any effort to come up with control strategies must put this into account



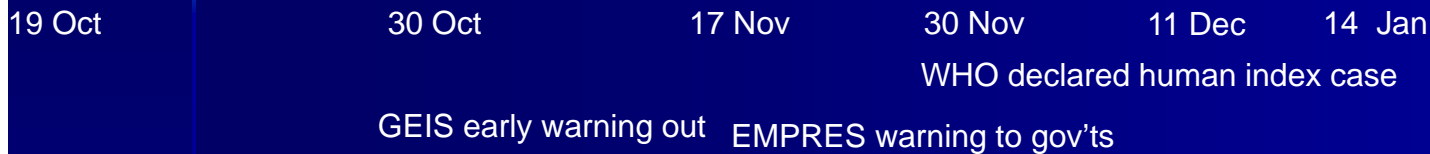
FIGURE 3. Number of reported Rift Valley fever cases (n = 330), by date of illness onset — Kenya November 2006–January 2007*



* As of January 25, 2007, for cases with known date of onset.



The average number of days between key events



How effective disease prevention can be achieved? - Stop the transmission!

1. Advanced Larviciding will:
 - avoid high vector densities
 - reduce vector infection rates
 - reduce virus transmission & amplification
 - reduce animal exposure <<< human exposure

3. Adulticides – During outbreak to reduce transmission – Too late

4. Livestock -insecticide treatment (pour on)?
 - movement of livestock from vector swarms

SUGGESTIONS



- **Vector surveillance necessary in outbreak hotspots/regions during IEP:**

- to facilitate early detection/prevention
- Identify IEP maintenance mechanism
- Localise emergence zones (vectors/reservoirs)

- **Analysis of blood in fed specimens**

- Extent virus transmission to human by mosquitoes,
- Identification of other possible reservoirs, the range of other hosts involved.



Acknowledgements

- KEMRI
- USAMRU Kenya - GEIS
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- NAMRU 3
- USAMRIID
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- USAID
- Kenya's MOPH&S and MOHS + their staff, Garissa, Kilifi, Baringo AND Kirinyaga



AND TO THE PEOPLE



THANK YOU