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# 8.6 BIODIVERSITY OF PARASITIC MITES OF HONEY BEES IN SOUTHEAST ASIA

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Biodiversity considers three levels of biological organization: species, genetic and ecological diversity. Southeast Asia is a biologically rich region (Thailand, Burma, Cambodia, Vietnam, Malaysia, and Indonesia) with many species of bee mites. Many more species are known to occasionally invade a bee colony or attach to a honey bee (Eickwort 1988, Delfinado-Baker and Rath 1992) but they will not be considered here. This paper surveys only those parasitic mites that are known to form significant associations with the six species of Apis in Southeast Asia (Table 1). Most of the parasitic mites of five native Asian honey bees are more specific to their own hosts than they are to the European honey bee, Apis mellifera. The association among species and populations of Asian honey bees and European honey bees of different genetic backgrounds with the 'new' parasites Varroa jacobsoni and Tropilaelaps clareae promises to provide new insights into 'coevolution'. This paper will review the diverse group of mites parasitic on Asian honey bees. Varroa jacobsoni and T. clareae are the most familiar species of mites, while Euvarroa sinhai and E. wongsirii are important on their hosts (Eickwort 1988, Lekprayoon and Tangkanasing 1991, 1993).

#### **DISCUSSION**

## Euvarroa wongsirii and E. sinhai

Both species are mesostigmatids in the family Varroidae, (Table I) and were first found in southeast Asia (Delfinado and Baker 1974, Delfinado-Baker 1984). Morphology and biological characters confirm that *E. wongsirii* is a separate species found on the small dwarf honey bee, *A. andreniformis*(Lekprayoon and Tangkanasing 1991, 1993, Morin and Otis 1993). The general morphology and chaetotaxy of *E. wongsirii* are very similar to *E. sinhai* but some characters are different. The characters that readily distinguish the female of *E. wongsirii* from *E. sinhai* are a broadly triangular-shaped body, wider, posteriorly, bearing 47-54 slender, long, lanceolate setae. The female of *E. sinhai* has a pear-shaped body, bearing 39-40 very long,

lanceolate setae. E. wongsirii is restricted to its original host, A. andreniformis and is limited to Southeast Asia, and was discovered in Thailand where its host is native. E. sinhaii has been reported from A. mellifera in India (Kapil and Aggarwal 1985) and A. andreniformis in Malaysia (Morin and Otis 1993), and is easily found on drone brood of its native host, A. florea, in Thailand. E. wongsirii is not a common species since its host, A. andreniformis, is rare, at least in Thailand. Euvarroa species are brood mites parasitizing their respective host species, but they can live phoretically in bee colonies both with and without drones. Koeniger et al. (1993) and Lekprayoon et al. (1994) reported that A. florea and A. andreniformis show aggressive cleaning behavior toward foreign mites and kill them (see Table 2). These observations are the basis of our knowledge of their ecology and nature. The relationship between mites and bees can be used to evaluate the hypothesis of a coevolutionary relationship among mite taxa specific to their own hosts (OConnor 1988).

# Varroa jacobsoni and Tropilaelaps clareae

Of the many extraordinary mites associated with honey bees, the brood mites V. jacobsoni and T. clareae are the most important and familiar haemophagous parasites. These two species of mesostigmatid mites evolved in southeast Asia as obligatory parasites of Apis. jacol ni was originally associated with A. cerana and T. clareae with A. dorsata (De Jong et al. 1982, Woyke 1984, Eickwort 1988). Delfinado-Baker's (1982) record for T. clareae from A. cerana established that the mite is a parasite of A. cerana (Wongsiri and Tangkanasing 1989). The original host of T. clareae and T. koenigerum is A. dorsata, but T. koenigerum remains restricted to A. dorsata in Sri Lanka only, not in Southeast Asia (Delfinado-Baker and Baker 1982). When beekeepers brought A. mellifera to Southeast Asia, V. jacobsoni and T. clareae became devastating pests (Eickwort 1988, Wongsiri et al. (1987).

Nymphal and adult female mites feed on the hemolymph of bee larvae and pupae. From a world

Table 1. Taxon and ecological diversity of honey bees (Apis) and their parasitic mites in Southeast Asia

Hosts	Mites (Taxon)	Family	Ecology	Range			
	Parasitiformes - Mesostigmata						
Apis andreniformis	Euvarroa wongsirii Varroidae		obligatory, parasitic	Southeast Asia			
	E. sinhai	Varroidae	incidental, parasitic?	Southeast Asia			
Apis florea	E. sinhai	Varroidae	obligatory, parasitic	Asia			
	Varroa jacobsoni	Varroidae	incidental, phoretic, parasitic?	Asia			
	Tropilaelaps clareae	Laelapidae	incidental, phoretic, parasitic?	Asia			
Apis dorsata	T. clareae	Laelapidae	obligatory, parasitic	Asia			
	T. koenigerum	Laelapidae	obligatory, parasitic	Sri Lanka			
Apis cerana	V. jacobsoni	Varroidae	obligatory, parasitic	Asia			
	T. clareae	Laelapidae	incidental, phoretic	Asia			
Apis koschevnikovi	V. rinderer*	Varroidae	obligatory, parasitic	Borneo, Indonesia			
Apis mellifera	V. jacobsoni	Varroidae	obligatory, parasitic	Asia, Africa, America, Europe			
	E. sinhai	Varroidae	incidental, phoretic	Asia			
	T. clareae	Laelapidae	obligatory, parasitic	Asia			
	Acariformes - Astigmata						
Apis mellifera	Acarapis woodi	Tarsonemidae	obligatory, parasitic	almost cosmopolitan, including			
	A. dorsalis	Tarsonemidae	obligatory, parasitic	Asia, but is not yet found in Thailand.			
	A. externus	Tarsonemidae	obligatory, parasitic				
pis cerana	A. woodi	Tarsonemidae	obligatory, parasitic				

<sup>\*</sup>de Guzman and Delfinado-Baker, 1996.

standpoint, *V. jacobsoni* is the most destructive and, hence, is of the greatest economic importance to beekeepers. However, *T. clareae* has come to be a more serious pest of A. mellifera than *V. jacobsoni* in Southeast Asia. When the two species are found in the same *A. mellifera* colony, *T. clareae* outnumber *V. jacobsoni* by several times (Burget et al. 1983, Wongsiri 1989). That appears to be explained by the rate of reproduction of *T. clareae*, which is much more rapid than that of *V. jacobsoni*. Adult females of *V. jacobsoni* feed on the hemolymph of their adult phoretic hosts and can survive low brood periods

during the rainy season and then increase reproduction again when the drone brood is present. In contrast, phoretic *T. clareae* do not feed on the adult host and can survive only 3-4 days without access to brood (Woyke 1984, Rinderer et al. 1994). Thus, *T. clareae* spends only short periods of time on adult honey bees. Perhaps a mechanism exists that allows mites to attach to foraging bees and shift to bees from different colonies at foraging sites. There may be physiological responses that permit *T. clareae* to attach to migrating *A. dorsata* and survive over the long periods required to follow such migrations

Hosts	A. andreniformis	A. florea	A. dorsata	A. cerana	A. mellifera	reference
Parasitic Mite	es on Adult Workers					
E. sinhai		70% (24hrs)		8%	63%	Koeniger et al. 1993
T. clareae	5% (12 hrs)	8% (12 hrs)	80%	31%	69%	Rinderer et al. 1994, Lekprayoon et al. 1994
V. jacobsoni			,	63%(1 hr)	73%(1 hr)	Wongiri et al. 1993

Table 2. Comparison of survival rates of parasitic mites in cage experiments

(Ruttner 1988, Rinderer et al. 1994). The survival of mites in our study, whether on A. mellifera or A. dorsata, is about one day longer than previously reported (Woyke 1984, Rinderer et al. 1994). Perhaps still longer periods of survival occur under other conditions. In any event, it is clear that the highly pestiferous T. clareae could easily survive even the longest of international airline flights. The recent discovery of the mite in Africa (Kumar et al. 1993), suggests that such an event has occurred.

When V. jacobsoni and T. clareae reproduce on their own original hosts they feed only on drone broad (Wongsiri and Tangkanasing 1989, Rath 1993) and are seldom found in worker brood cells. We found a resistance mechanism in Asian honey bees, A. cerana and the giant honey bee, A. dorsata, to their ectoparasitic mites (Peng et al. 1987, Wongsiri et al. 1987, Wongsiri 1988, Wongsiri and Tangkanasing 1989). Trouiller et al. (1993) showed that the kairomones from the cuticle of drone larvae were much more attractive to V. jacobsoni than kairomones of worker larvae. This could explain the preference of these mites for only the drone brood. The fact that neither of these mites are significant pests of their own native hosts may be an evolved mutualism and coevolution (OConnor 1988), that results in the mites killing the drones but not the workers. This may even be advantageous, since the tropical climates of Southeast Asia lack the cold temperature that kill drones. Drones are killed in the A. mellifera colonies in temperate climates by cold weather.

We also observed the cleaning behavior of A. mellifera, A. cerana, A. dorsata, A. florea and A. andreniformis toward different kinds of mites (Table 2) (Wongsiri and Tangkanasing 1989, Koeniger et al. 1993, Rinderer et al. 1994, Lekprayoon et al. 1994). These reports show that most mites are restricted to their own hosts except for A. mellifera, which is susceptible to V. jacobsoni and T. clareae. Perhaps the kairomones from

foreign bees are more attractive than their own species. However, A. mellifera also has a much lower frequency of cleaning behavior against mites than the other bees.

Population genetic analysis and subspecific hybridization between populations of A. mellifera have been reported by Rinderer and Oldroyd (1993). DNA analysis of A. cerana (Sylvester and Wongsiri 1993) and A. florea (Oldroyd et al. 1994) but no analysis of genetic variation in most of the parasitic bee mites has been reported. Transposable elements in Drosophila genes transferred by the mite Proctolaelaps regalis has been studied by Houck et al. (1991). Such transfer might be a long-term solution to the control of external parasitic mites in A. mellifera. The most promising avenue is the transfer of the behavioral and physiological resistance adaptations of Asian honey bees to A. mellifera. There are two major obstacles to such a transfer. First, we have no knowledge of the genetic basis in Asian honey bees of mite resistance. Obtaining such understanding will be labor intensive. Second, the technology to transfer these traits in honey bees is not yet available.

#### **SUMMARY**

The biodiversity of parasitic honey bee mites in Southeast Asia is an undervalued and neglected resource, both from a biologic and economic view, especially as it is related to honey bees as pollinators. On a global scale, the rate of species loss due to human disturbance is at least 1,000 times greater than the natural background rate. Southeast Asia's commercial forests were once of great economic importance, and would not now be depleted if we knew enough about the million or more wild bee colonies lost due to fire and the timber industry. All A. andreniformis, A. florea, A. dorsata, A. cerana and A. koschevnikovi live in the forest. Due to a lack of proper knowledge, the

alarmingly high losses of A. andreniformis due to human disturbance, and of biodiversity at all levels make it imperative to study these bees.

Southeast Asia is the center of honey bee-mite diversity. All six of their host species in the genus Apis are found here. Curiously, the most studied and economically exploited honey bee, A. mellifera, is an "exotic" species recently imported to Southeast Asia. The ranges for A. cerana, A. dorsata, A. florea, A. andreniformis and A. koschevnikovi are only partially known. A long evolutionary history of honey bees in Southeast Asia favor conditions for the evolution of the diverse set of diseases, pests, and parasitic mites of honey bees. Thus, Southeast Asia is an ideal place for studies of the biodiversity and control of these problems under conditions that existed before their appearance elsewhere in the world.

In conclusion, the association between parasitic mites and honey bees offers great opportunities for evolutionary studies. Many of the examples cited as mechanisms of resistance demonstrate the scientific rewards to be gained through the collaborative study by acarologists with bee scientists. These observations form the basis for our knowledge of the ecology and nature of the relationship between mites and bees. Similarly, taxonomists should be encouraged to discover the new species of bee mites and formulate hypotheses of evolutionary relationships among mite taxa and bees.

### **ACKNOWLEDGMENTS**

We thank the Research Affairs Division, Chulalongkron University for funding this research. Sincere thanks are extended to Prof. Dr. Pensri Tangkanasing of the Bee Biology Research Unit, Chulalongkron University and Dr. Mercedes Delfinado-Baker, Systematic Entomology Laboratory, ARS-USDA, Beltsville, Maryland for reviewing and discussing this manuscript.

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