MOSQUITO HOST RANGE AND SPECIFICITY OF EDHAZARDIA AEDIS (MICROSPORA: CULICOSPORIDAE)

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ABSTRACT. Edhazardia aedis was transmitted horizontally to its natural host, Aedes aegypti, and to 6 alternate hosts: Ae. albopictus, Ae. triseriatus, Ae. taeniorhynchus, Anopheles quadrimaculatus, Orthopodomyia signifera and Toxorhynchites rutilus rutilus. The microsporidium produced both binucleate and uninucleate spores in all susceptible hosts. Transovarial transmission, however, was only successful in Ae. aegypti. Therefore, while E. aedis can infect a variety of mosquito species from diverse genera, it is specific for its natural host, Ae. aegypti. Five other mosquito species were not susceptible to E. aedis.

INTRODUCTION

Edhazardia aedis (Kudo) was originally observed and described from Aedes aegypti (Linn.) in Puerto Rico (Kudo 1930) and reisolated in Thailand (Hembree 1979). This microsporidium has a complex life cycle termed the "parentalhost filial-host alternation pathway" (Becnel et al. 1989): part of this pathway involves transovarial transmission from infected adult females to progeny via a binucleate spore (Hembree and Ryan 1982). Uninucleate, pyriform spores are formed in the fat body of these progeny, and this process is normally fatal. Spores released from these dead individuals into the aquatic environment are infectious per os to Ae. aegypti larvae yielding infected adults to complete development. Because of these and other life cycle features, E. aedis has potential as a biocontrol agent for container-inhabiting mosquitoes (Becnel 1990).

Hembree (1982) reported that Ae. aegypti and Ae. taeniorhynchus (Wied.) were susceptible to E. aedis; Culex quinquefasciatus Say and Anopheles stephensi Liston were not susceptible. Aedes taeniorhynchus was found to be "almost as infectious" to E. aedis as Ae. aegypti, but no details were given on other host-parasite interactions. The purpose of the present study was to further define the mosquito host range of E. aedis and to examine the interactions of the parasite with the susceptible alternate hosts.

MATERIALS AND METHODS

The isolate of *E. aedis* used originated from Thailand and has been maintained since 1981 in laboratory colonies of *Ae. aegypti* according to the methods of Hembree and Ryan (1982). Larvae from infected eggs were reared at 27°C for 7–9 days, and patently infected larvae were isolated for transmission experiments. A portion of these larvae were triturated in a glass tissue grinder, and large particulate matter was removed from this extract by forcing it through cotton in a syringe. The inoculum was washed and centrifuged 3 times according to the protocol of Undeen and Maddox (1973), held at 20°C and used within 24 h. Spore concentrations were determined with a hemocytometer prior to feeding.

Twelve mosquito species representing 7 genera were bioassayed for susceptibility to E. aedis (Table 1). Mosquitoes for bioassays either were obtained from existing laboratory colonies or were field-collected from the environs of Gainesville, Florida.

Second instar mosquito larvae (\sim 48 h old) were exposed in groups of 100 in 100 ml of a spore suspension for 24 h. A range of doses between 1 \times 10² and 1 \times 10⁵ spore/larva was used to calculate estimates of the IC₅₀ and LC₅₀ dosages. For species that were not susceptible to E. aedis in this range of doses, a maximum challenge of 1×10^6 spores/larva was administered. All treatment groups were then transferred to pans with 500 ml of water and fed according to standard rearing protocols. Second instar larvae (72 h old) of the predacious mosquito, Toxorhynchites rutilus rutilus (Coq.), were isolated and fed one infected 4th instar Ae. aegypti larva (one larva $= 3 \times 10^5$ spores) and healthy larvae thereafter. Second instar larvae of Ae. aegvpti were exposed with each test to verify spore viability. Control groups were used in all tests and handled in a similar manner but without the addition of spores.

Individuals from the exposed groups were sampled approximately 1 h post-exposure, and their gut contents were examined with phase microscopy for the presence and condition of spores. Mortality was recorded when pupation began, and a sample of the surviving individuals was smeared on microscope slides and stained with a Giemsa-stain solution; percentage of infection was determined microscopically. The IC_{50} and LC_{50} dosages were calculated when possible by probit analysis.

							Transe transm	ovarial nission
		No. exposed		95% confidence		95% confidence	No.	No.
Mosquito host	Source	(replications)	IC_{50}	interval	LC ₅₀ 1	interval	exam.	pos.
tedes aegypti	C3	10,000 (10)	39	29-49	1,270	560-2,400	22	22
tedes albopictus	U	300 (3)	84	68-126	3,800	975-29,423	17	0
Aedes taeniorhynchus	U	300 (3)	3,080	1,758-5,145	29,600	18,675-52,962	15	0
Aedes triseriatus	U	300 (3)	29.7	2-100	2,280	1,600-3,500	30	0
Anopheles quadrimaculatus	с С	400 (4)	2.5	1	58,500	16,000-200,000	20	0
Anopheles albimanus	U	300 (3)	0	I	. 1	I	N/A	N/A
Culex nigripalpus	FC ³	300 (1)	0	I	I	I	N/A	N/A
Culex aunquefasciatus	U	400 (4)	0	I	I	I	N/A	N/A
Culiseta melanura	FC	250 (2)	0	I	I	I	N/A	N/A
Orthopodomyia signifera	FC	50 (1)	pos.	ł	I	1	N/A	N/A
Uranotaenia sapphirina	FC	22 (2)	0	1	I	I	N/A	N/A
Toxorhynchites rutilus rutilus	c	74 (2)	pos.	I	1	1	×	0

Table 1. Mosquito host range of Edhazardia aedis.

¹ Spores per individual. ² Colonized. ³ Field-collected.

JOURNAL OF THE AMERICAN MOSQUITO CONTROL ASSOCIATION

Vol. 9, No. 3

Two methods were required to determine if the parasite could complete its life cycle in an alternate mosquito host. One was to examine infected female adults (after oviposition) for the presence of binucleate spores. If binucleate spores were observed, progeny from these infected females were examined for the presence of infection and or pyriform spores. The other method was to hold infected individuals with extended development (either larvae, pupae or adults) and examine them for spores (binucleate and pyriform spores) at various times post-exposure. If pyriform spores were produced in the alternate host, these were bioassayed in *Ae. aegypti* larvae to determine viability.

RESULTS

Uninucleate spores of *E. aedis* were observed among the gut contents of individuals from each species of mosquito tested. Ungerminated spores were easily recognized by their highly refractile appearance as demonstrated in the natural host, *Ae. aegypti* (Fig. 1), and in *Cx. quinquefasciatus* (Fig. 2). Spores of *E. aedis* readily germinated in all mosquitoes tested and were identified by their dark color and clearly defined spore wall (Figs. 1 and 2). In *Ae. aegypti*, the everted polar tube penetrated the peritrophic membrane (Fig. 1), but this was not observed in *Cx. quinquefasciatus* (Fig. 2).

Edhazardia aedis was transmitted to 7 mosquito species representing 4 genera: Ae. aegypti, Ae. albopictus (Skuse), Ae. triseriatus (Say), Ae. taeniorhynchus, An. quadrimaculatus Say, Orthopodomyia signifera (Coq.) and Tx. rutilus rutilus. Estimates for the IC₅₀ and LC₅₀ are given in Table 1. There was no evidence of infection in the other mosquito species tested.

Developmental stages of E. aedis in the alternate mosquito hosts appeared as previously described (Becnel et al. 1989) beginning with uninucleate stages (gamonts and gametes), followed by plasmogamy and nuclear association to form diplokaryotic meronts. These stages underwent sporulation (sporogony plus sporogenesis) to form binucleate spores. Binucleate spores were found in larvae, pupae and adults of each mosquito host that became infected with E. aedis (Figs. 3 and 4). In the alternate hosts, these spores were determined to be functionally mature by their normal germination and eversion of the polar tube (Fig. 4).

Edhazardia aedis was transmitted transovarially and therefore completed its life cycle in all families of Ae. aegypti examined (n = 22). No infections were found in the F₁ progeny of individual families from infected females (positive for binucleate spores) of Ae. albopictus (n = 17), Ae. triseriatus (n = 30), Ae. taeniorhynchus (n = 15), An. quadrimaculatus (n = 20) or Tx. rutilus rutilus (n = 8). Transovarial transmission in Or. signifera could not be determined.

Uninucleate, pyriform spores were found in horizontally (orally) infected Ae. aegypti, Ae. albopictus, Ae. triseriatus, Ae. taeniorhynchus and Tx. rutilus rutilus with extended development (>12 days). Regardless of the host in which they were formed, these uninucleate spores were infectious per os to Ae. aegypti larvae in which they initiated the normal developmental sequence leading to the production of binucleate spores.

DISCUSSION

Edhazardia aedis was transmitted horizontally to its natural host, Ae. aegypti, and 6 alternate hosts: Ae. albopictus, Ae. triseriatus, Ae. taeniorhynchus, An. quadrimaculatus, Or. signifera and Tx. rutilus rutilus. The dosage of E. aedis which produced mortality when exposed to 2nd instar Ae. aegypti (LC₅₀ = 1,270 spores/larva) was approximately 30 times the dose required to infect $(IC_{50} = 39 \text{ spores/larva})$. These dosages are similar to those reported by Hembree (1982). The estimated IC₅₀ and LC₅₀ for the alternate hosts varied greatly with either large or incalculable confidence intervals (Table 1), which indicated that individual responses are not predictable for the unnatural hosts. Anopheles quadrimaculatus was the most susceptible species to E. aedis (IC_{50} = 2.5) but also required the highest dose for mortality (LC₅₀ = 58,500). The other species responded similarly to Ae. aegypti except for Ae. taeniorhynchus, which was considerably less susceptible. This differed from Hembree (1982) who found similar IC₅₀s for 24-h-old Ae. aegypti and Ae. taeniorhynchus larvae exposed to E. aedis. This discrepancy could be due to the age of larvae used for the comparison or to strain differences between the mosquito species.

Normal vegetative growth and development occurred in all of the susceptible species, and *E. aedis* did sporulate to form binucleate spores in larvae and adults. In its natural host, *Ae. aegypti*, binucleate spores were formed in oenocytes that invaded the ovaries (Becnel et al. 1989) and consistently resulted in transovarial transmission to larval progeny (Table 1). However, *E. aedis* was not transovarially transmitted to the filial generations in any of the susceptible, alternate mosquito hosts examined, even though mature binucleate spores were present and presumably germinated in these species.

In what was described as "deviations from the parental-host filial-host alternation," Becnel et al. (1989) described 2 alternate pathways by which *E. aedis* completes its developmental cycle in *Ae*.



Figs. 1–4. Spores of *Edhazardia aedis*. Fig. 1. Uninucleate spores within the peritrophic membrane of *Aedes aegypti*. Fig. 2. Uninucleate spores within the peritrophic membrane of *Culex quinquefasciatus*. Fig. 3. Binucleate spores from adult *Ae. triseriatus*. Fig. 4. Germinating binucleate spore from adult *Ae. albopictus*. G, germinated spore; UG, ungerminated spore; PM, peritrophic membrane; PT, polar tube; IS, immature spore. All fresh preparations and $1,000 \times$.

aegypti. One of these involves individuals infected per os as larvae whose longevity extended up to 2 wk. In these individuals, the entire developmental cycle of E. aedis was often completed by producing binucleate spores which presumably germinated (autoinfection) to initiate the sequences leading to the production of uninucleate spores. This alternate pathway may serve a supportive role for parasite maintenance within Ae. aegypti populations under certain conditions. In the current study, a similar pathway was found in each of the alternate mosquito hosts in which E. aedis completed its developmental cycle in those infected individuals with an extended development time. This ability of E. aedis to complete its developmental cycle in alternate hosts might also serve a supportive role to the primary parental-host filial-host pathway in Ae. aegypti. While the species assemblage used in this study is not usually associated with the natural host, it demonstrates that alternate hosts in the natural system may provide reservoirs for maintaining and amplifying the parasite when *Ae. aegypti* numbers are low. Maintenance of *E. aedis* within living hosts seems necessary, especially due to the fragile nature of the uninucleate spores, which would not be expected to persist for long periods once released into the environment (Undeen and Becnel 1992).

Andreadis (1989) demonstrated that 4 alternate mosquito hosts were susceptible to oral infection with *Amblyospora connecticus* Andreadis from the natural host *Ae. cantator* (Coq.); however, binucleate spores were produced only in female *Ae. epactius* Dyar and Knab. These binucleate spores did not invade or germinate in ovarian tissues, and *A. connecticus* could not complete its life cycle via transovarial transmission. Andreadis concluded that A. connecticus is specific for Ae. cantator and that host specificity operates at 3 different levels: larval infectivity. sporulation and spore germination with ovarian infection. While the host specificity of E. aedis was clearly operational at the larval infectivity level, specificity was not related to the sporulation or germination of binucleate spores. Both binucleate and uninucleate spores of E. aedis were produced in each of the alternate mosquito hosts tested, and germination of binucleate spores seems likely as this is assumed to be a prerequisite for the production of uninucleate spores (Becnel et al. 1989). While E. aedis and A. connecticus are similar in that transovarial transmission is not successful in alternate hosts. E. aedis differs in the ability to complete its developmental cycle in alternate hosts in the absence of transovarial transmission. Therefore, host specificity of E. aedis cannot be defined by developmental parameters, such as sporulation, germination or completion of the developmental cycle, but rather by the successful transovarial transmission of the parasite to the next generation.

Uninucleate spores germinated in the guts of all experimental mosquito larvae, and germination was not correlated with the susceptibility of the host to E. aedis. Uninucleate spores of E. *aedis* also germinated in the guts of a variety of nontarget aquatic organisms but did not produce infections (Becnel 1992). Based on previous studies with Nosema algerae Vavra and Undeen, Undeen (1976) speculated that the lack of detectable infections was due to a gut barrier preventing the infective germ from entering the hosts rather than some physiological incompatibility preventing parasite development. Our observations of the polar tube penetrating the peritrophic membrane of a susceptible host, Ae. aegypti (Fig. 1), and not penetrating the membrane of a refractory host, Cx. quinquefasciatus (Fig. 2), support this hypothesis.

The environmental safety of a microbial control agent is evaluated, in part, on knowledge of its host range. Emphasis has been placed on selecting candidates with both a wide host range for target pests and a minimal risk to nontarget organisms (Brooks 1988). For microsporidia with complex host-parasite relationships, there is a clear difference between the host range of a particular species and its specificity. This study has demonstrated that E. aedis can infect a variety of mosquito species representing diverse genera and, therefore, has a broad host range. However, E. aedis is specific for its natural host, Ae. aegypti, based on its ability to complete its life cycle via transovarial transmission, which does not occur in the alternate mosquito hosts tested. The criteria needed to establish host limits of certain microsporidia must therefore consider the differences between host range and host specificity. This has important implications for the taxonomic determination of specific microsporidia as well as identifying suitable target hosts for biological control projects.

Because of its specificity for Ae. aegypti, E. aedis has its greatest potential as a biocontrol agent for its natural host, Ae. aegypti. Since both horizontal and transovarial transmission are necessary for maintenance of parasites like E. aedis in mosquito populations (Andreadis and Hall 1979), it is unlikely that E. aedis could persist in an alternate mosquito host without a functional transovarial transmission sequence.

ACKNOWLEDGMENTS

Our sincere appreciation goes to Genie White for her excellent technical assistance with the statistical analyses. We also thank T. G. Andreadis, T. Fukuda and A. H. Undeen for their helpful suggestions and comments on an earlier draft of the manuscript.

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