Dexrazoxane as a viable microsporidia control agent in *Anopheles gambiae*

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Abstract

Microsporidia have long been proposed as biological agents for controlling disease vectors and the parasites they transmit. However, their study in vector biology has been constrained due to challenges in manipulating microsporidia within hosts. In this study, we investigated the effect of Dexrazoxane, a candidate drug against microsporidiosis, on the establishment and development of *Vavraia culicis* infection in its natural host, the mosquito *Anopheles gambiae*, the main malaria vector. Our findings show that Dexrazoxane significantly reduces spore load, particularly in mosquitoes reared individually, without affecting the overall infection success of the parasite. This result aligns with studies in *Caenorhabditis elegans*, where Dexrazoxane inhibited new spore production without hindering initial spore integration into the host gut cells. Dexrazoxane's DNA topoisomerase II inhibitor mechanism likely explains its impact on mosquito development, as larvae exposed to the drug failed to emerge as adults. These findings highlight Dexrazoxane's potential as a viable tool for controlling microsporidia in adult mosquitoes and hope to enhance the study of mosquito-microsporidia interactions. Further research is required to explore its broader application in vector-borne disease control, including malaria.

Introduction

Microsporidia are common obligate parasites of many animals, and their prevalence has been increasingly reported over the past decades across a wide range of taxa, sparking a surge in microsporidia studies [1–7]. While research in human-infecting microsporidia mainly focuses on clearing them (i.e., microsporidiosis) [8], in invertebrates such as insects, their study is more nuanced, as evidence suggests microsporidia may serve as biological control agents for certain pests [9] and vector species, and the diseases they transmit [10–12].

Nearly half of the microsporidia species that infect insects have specialised in targeting dipterans, such as flies and mosquitoes, which are vectors for several human diseases. Among those infecting mosquitoes, *Vavraia culicis* [13] is one of the most well-studied microsporidia species [14–23]. This generalist intracellular parasite is known to infect several mosquito genera, including *Anopheles gambiae*, the major vector of malaria [18]. Infection typically occurs during the larval stage when mosquito larvae ingest spores along with food in water bodies. Once the spores penetrate the host's gut wall, they develop and produce new spores. The limited immune response of *An. gambiae* to this particular microsporidian species [24] allows for a steady progression of the infection, characterised by continuous consumption of host resources and accumulation of spores in the body cavity, leading to costs in host longevity [18, 25] and fecundity [14].

Although *An. gambiae* incurs significant costs from *V. culicis* infection, it comes with a benefit in protection against co-infection with *Plasmodium spp.* by negatively affecting oocyst development [26,

27]. These findings led the community to hypothesise that microsporidian parasites, such as the one in

this study, could be used as biological control agents for mosquitoes and their parasites by reducing the

mosquito lifespan and parasite transmission to mammal hosts, respectively [11, 26, 27]. However, the

mechanisms underlying this interaction remain unclear, partly due to the technical challenges of

manipulating microsporidian infections within mosquito hosts.

Hence, to facilitate the control and study of microsporidians within the mosquito host, a candidate

chemical, Dexrazoxane, recently shown to have anti-microsporidian activity in *Caenorhabditis elegans*

[29] was tested. Due to its antimitotic and redox activity, Dexrazoxane has been extensively used as a

protector agent against the cardiotoxicity of chemotherapy in humans [30]. Its mechanism of action is

similar to ethylenediaminetetraacetic acid (EDTA) but diffuses into cells more readily [31], where it

binds to iron and lowers oxidative stress formation [32]. However, its role against microsporidia has

only recently been studied. For instance, in *C. elegans*, Dexrazoxane is known to inhibit *Nematocida*

parisii development after it has established itself in the gut cells [29]. While it does not prevent initial

infection, it suppresses the production of new spores, thereby hindering disease progression and

transmission. Given its promising effects on microsporidia infection in *C. elegans*, we aimed to test

whether these effects extend to microsporidian infections in mosquitoes, using *An. gambiae* and *V.*

culicis as the experimental system. Our findings suggest that Dexrazoxane's effect might be more

general than previously thought, extending to mosquito-microsporidia interactions. This result may

enhance our understanding of microsporidia interactions with their host and other parasites and could

potentially inform and fuel future vector-borne disease control strategies.

Materials and Methods

Experimental system

We used the *An. gambiae s.s* Kisumu strain [33], which had been maintained at our standard laboratory

conditions (about 600 individuals per cage, constant access to 6% sucrose solution, $26 \pm 1^{\circ}$C, $70 \pm 5\%$

relative humidity and 12 h light/dark) for many years before the experiments.

The microsporidian *V. culicis floridensis* was provided by J.J Becnel (USDA, Gainesville, FL, USA).

Originally discovered in *Ae. albopictus,* this parasite was later found to parasitise several mosquito

genera, including *Aedes*, *Culex*, *Anopheles*, *Culiseta*, *Ochlerotatus* and *Orthopodomyia* [20, 34, 35].

We maintained it by alternately infecting *Aedes aegypti* and *An. gambiae* to ensure its status as a

generalist parasite, as described previously [18].

V. culicis is an obligatory, intracellular parasite. Mosquitoes are infected when, as larvae, they ingest

the spores along with their food. After several rounds of replication, the parasite produces new

infectious spores that spread to gut and fat body cells. The spores are released to larval sites when

infected larvae die, when adults die on the surface of water, or when eggs covered with spores are laid

onto the water's surface.

Rearing and maintenance of experimental mosquitoes

Freshly hatched (0-3 hours old) *An. gambiae* larvae were individually placed into 12-well culture plates,

each containing 3 ml of deionized water. The larvae received Tetramin Baby® fish food every day

according to their age (0.04, 0.06, 0.08, 0.16, 0.32 and 0.6 mg/larva respectively on days 0, 1, 2, 3, 4

and 5 or older [36]).

In a first experiment, we assessed the impact of Dexrazoxane on larval development and spore growth.

Two-day-old larvae were exposed to either 0 or 10,000 spores of *V. culicis*, and received one of our

three Dexrazoxane doses (0 µM, 30 µM and 60 µM) at one of three administration times (four, five, or

six days after egg hatching), in a fully combinatorial design. Survival and pupation of the larvae were

tracked daily.

In a second experiment, we assessed the impact of Dexrazoxane on spore growth at the adult stage and

if being reared individually as opposed to collectively could influence this growth. Two-day-old larvae

were exposed to either 0 or 10,000 spores of *V. culicis*. Pupae were transferred to individual 50 ml

falcon tubes with approximately 10 ml of deionised water [37]. Female adults were moved to 150 ml

cups (5.5 cm Ø x 10 cm) covered with a net or cages, depending on their treatment. The cups contained

a 50 mm Ø petri dish (to prevent mosquitoes from drowning) floating on the surface of 50 ml deionised

water and a 10 x 7 cm rectangular filter paper to maintain humidity. The mosquitoes were provided

with cotton balls soaked in a 6% sucrose solution or a 6% sucrose solution supplemented with 60 µM

of Dexrazoxane, which was replaced every other day.

On the 8th day, alive individuals were collected in a 2 ml Eppendorf tube containing 0.1 ml of deionised

water and stored at -20º C for spore load quantification. A stainless-steel bead (Ø 5 mm) was added to

each tube and the samples were homogenised using a Qiagen TissueLyser LT at a frequency of 30 Hz

for two minutes. The number of spores in each sample was then counted using a haemocytometer with

0.1 µl of the sample under a phase-contrast microscope (400x magnification).

Statistical analysis

All analyses were conducted with the R software [38] version 4.4.1, using the packages DHARMa [39],

car [40], glmmTMB [41], emmeans [42] and multcomp [43]. Significance was assessed with the

"Anova" function of the "car" package [40]. We used a type III ANOVA in the case of a significant

interaction and a type II ANOVA otherwise. When relevant, we performed post-hoc multiple
comparisons with the package "emmeans", using the default Tukey adjustment.

The presence of spores was coded as a binary response (presence or absence of detectable spores) and
analysed with a generalised linear model with a binomial distribution of errors, where the explanatory
variables were diet (with or without Dexrazoxane), type or rearing (individual or collective) and their
interaction. Because we counted the number of spores in a haemocytometer containing 0.1 µl of the
sample (i.e. 1/1000 of the total volume), the detection threshold was estimated to be 1000 spores.

Spore load of mosquitoes with positive spore counts was analysed with a generalised linear model with
a negative binomial distribution of errors. The explanatory variables were diet (with or without
Dexrazoxane), type or rearing (individual or collective), and their interaction.

Results

Larval exposure to Dexrazoxane delayed larval development and blocked pupation, independently of
the dose, with none of the treated larvae surviving past pupation (as opposed to Dexrazoxane-unexposed
larvae). That led us to discard the use of this chemical during the larval stage.

The proportion of individuals harbouring spores was lower for individuals reared in cups than for
individuals reared in cages (0.65 vs 0.85, χ^2 = 12.25, df = 1, p < 0.001, **Fig. 1a**) but was not affected by
the consumption of Dexrazoxane (0.75 vs 0.75, χ^2 = 0.00, df = 1, p = 0.945) or their interaction (χ^2 =
0.28, df = 1, p = 0.595). In contrast, the spore load of mosquitoes with positive counts was reduced by
half in individuals consuming Dexrazoxane, but only when were reared in cups (4.9×10^4 vs 1.2×10^5
in cups, and 9.5×10^4 vs 6.7×10^4 in cages, interaction type of rearing * diet: χ^2 = 10.35, df = 1, p =
0.001, **Fig. 1b**). For further details on the statistical analysis see **Table 1**.

152　**Table 1. Effect of exposure to Dexrazoxane on infection.** Statistical models carried on to test for
153　effect of exposure to Dexrazoxane, rearing condition (individual in cup, or collective in cage) and their
154　interaction on the presence of spores (Model 1a) and number of spores (Model 1b). See Materials and
155　methods for more details on the experimental design and statistical tests. Values in bold are statistically
156　significant.

Tested effect			
Model 1a – Presence of spores	df	χ^2	p
Exposure to Dexrazoxane	1	0.00	0.945
Rearing (Cup/Cage)	1	12.25	**<0.001**
Exposure to Dexrazoxane: Rearing (Cup/Cage)	1	0.28	0.595
Model 1b – Spore load	df	χ^2	p
Exposure to Dexrazoxane	1	2.13	0.144
Rearing (Cup/Cage)	1	4.68	**0.030**
Exposure to Dexrazoxane: Rearing (Cup/Cage)	1	10.35	**0.001**

Figure 1. Parasite load of infected mosquitoes after consuming a specific diet for seven days. The proportion of mosquitoes reared individually or collectively that had detectable spores for **(a)** each Dexrazoxane diet and their spore load **(b)**. The sample sizes were 70, 60, 60 and 60 for **(a)**, and 60, 50, 38 and 40 for **(b)**. Error bars show the 95% confidence intervals of the means. Letters indicate statistically significant differences from multiple comparisons. See Table 1 for further details on the statistical analyses.

Discussion

Our study investigates the effect of Dexrazoxane on the infection by the microsporidian *V. culicis* in the mosquito host *An. gambiae*. The findings extend the current knowledge of microsporidia-mosquito interactions and suggest potential strategies for controlling microsporidia infections and mosquito-borne diseases like malaria.

The primary finding from this study is that Dexrazoxane significantly reduces the spore load of infected mosquitoes but only under certain rearing conditions, mainly when mosquitoes are raised individually in cups **(Fig. 1b)**. Moreover, although the number of spores produced drops, the number of successful infections did not differ, compared to unexposed **(Fig. 1a)**. This result aligns with previous research on its inhibitory effects on microsporidian development in *C. elegans* [29], which demonstrated Dexrazoxane primarily reduces new spore production, rather than affecting the initial spore integration into the host gut wall. This reduction in spore numbers came at no cost to host survival.

The reduced spore load in mosquitoes raised individually in cups highlights a potential indirect effect of Dexrazoxane on the mosquito immune response or microsporidian metabolism. While any hypothesis cannot be ruled out, this inhibition is likely due to the drug's mechanism of action. For human purposes, Dexrazoxane primarily functions through iron chelation, where it blocks circulating iron from being used to fuel oxidative stress and tissue damage. Although this mechanism has been shown to affect *Plasmodium falciparium* development *in vitro* [44], the same cannot be said for microsporidia. In fact, *C. elegans* infection with the microsporidia *N. parisii* can be blocked using Dexrazoxane but cannot be reverted through iron supplementation [29]. This result suggests inhibition in microsporidian infections might not happen via iron chelation but instead through another mechanism of action. Interestingly, Dexrazoxane also acts as a DNA topoisomerase II inhibitor, which likely explains why larvae exposed to the drug failed to emerge as adults (see Materials and Methods). In a study on *Ae. aegypti*, larvae treated with Dexrazoxane exhibited morphological defects and low survival during both larval and pupal stages [45]. Therefore, it is plausible that *V. culicis* relies on the same protein or a structurally similar one and inhibiting this protein during the parasite growth phase might block spore production and infection progression.

It is important to note that the context of mosquito hosts introduces additional complexity compared to previous studies performed *in vitro* or with *C. elegans* [29, 44]. Dexrazoxane's reduction in spore load, especially in isolated conditions, suggests that other factors, such as social interactions and spore transmission, may influence its efficacy. These factors should be carefully controlled and studied in future research, along with determining the optimal dosage. Despite these complexities, the results of this study open the door for further investigation into microsporidia as potential biological control agents, particularly for *V. culicis*. Moreover, the strategic timing of the suppression of the infection

progression, without completely clearing it, offers several advantages for studying interactions with vector-borne parasites, such as *Plasmodium.* We believe this approach will allow researchers to investigate the mechanisms behind microsporidia-driven inhibition, for instance, whether it is driven by collateral immunity or resource competition. A further understanding of these mechanisms will allow us to better prepare and design vector control strategies, which are essential today.

References

[1]   Weidner E, Overstreet R. Microsporidia have a peculiar outer membrane with exterior cytoplasmic proteins. *Acta Parasitol* 2021; 1–6.

[2]   Becnel JJ, Weiss LM. *Microsporidia: pathogens of opportunity*. John Wiley & Sons Incorporated, 2014.

[3]   Wadi L, Reinke AW. Evolution of microsporidia: An extremely successful group of eukaryotic intracellular parasites. *PLoS Pathog* 2020; 16: e1008276.

[4]   Bojko J, Reinke AW, Stentiford GD, et al. Microsporidia: a new taxonomic, evolutionary, and ecological synthesis. *Trends Parasitol* 2022; 38: 642–659.

[5]   Franzen C. Microsporidia: a review of 150 years of research. *Open Parasitol J*; 2.

[6]   Han B, Takvorian PM, Weiss LM. Invasion of host cells by microsporidia. *Front Microbiol* 2020; 11: 172.

[7]   Seatamanoch N, Kongdachalert S, Sunantaraporn S, et al. Microsporidia, a highly adaptive organism and its host expansion to humans. *Front Cell Infect Microbiol* 2022; 12: 924007.

[8]   Han B, Pan G, Weiss LM. Microsporidiosis in humans. *Clin Microbiol Rev* 2021; 34: e00010-20.

[9]   Bjørnson S, Oi D. Microsporidia biological control agents and pathogens of beneficial insects. *Microsporidia Pathog Oppor* 2014; 635–670.

[10]   Becnel JJ, Andreadis TG. Microsporidia in insects. *Microsporidia Pathog Oppor* 2014; 521–570.

[11]   Bukhari T, Pevsner R, Herren JK. Microsporidia: a promising vector control tool for residual malaria transmission. *Front Trop Dis* 2022; 3: 957109.

[12]   Koella JC, Lorenz L, Bargielowski I. Microsporidians as evolution-proof agents of malaria control? *Adv Parasitol* 2009; 68: 315–327.

[13]   Vavra J, Becnel JJ. Vavraia culicis (Weiser, 1947) Weiser, 1977 revisited: cytological

characterisation of a Vavraia culicis-like microsporidium isolated from mosquitoes in Florida
and the establishment of Vavraia culicis floridensis subsp. n. *Folia Parasitol (Praha)* 2007;
54: 259.

[14] Silva LM, Koella JC. Complex interactions in the life cycle of a simple parasite shape the
evolution of virulence. *bioRxiv* 2024; 2024.01.28.577571.

[15] Vyas-Patel N. The Suppression of Plasmodium berghei in Anopheles coluzzii infected later
with Vavraia culicis. *bioRxiv* 2023; 2002–2023.

[16] Rivero A, Agnew P, Bedhomme S, et al. Resource depletion in Aedes aegypti mosquitoes
infected by the microsporidia Vavraia culicis. *Parasitology* 2007; 134: 1355–1362.

[17] Silva LM, Koella JC. Virulence evolution: thinking outside of the host. *bioRxiv* 2024; 2005–
2024.

[18] Zeferino TG, Koella JC. Host-specific effects of a generalist parasite of mosquitoes. *Sci Rep*
2024; 14: 18365.

[19] Agnew P, Bedhomme S, Haussy C, et al. Age and size at maturity of the mosquito Culex
pipiens infected by the microsporidian parasite Vavraia culicis. *Proc R Soc London Ser B Biol*
Sci 1999; 266: 947–952.

[20] Kelly JF, Anthony DW, Dillard CR. A laboratory evaluation of the microsporidian Vavraia
culicis as an agent for mosquito control. *J Invertebr Pathol* 1981; 37: 117–122.

[21] Desjardins CA, Sanscrainte ND, Goldberg JM, et al. Contrasting host–pathogen interactions
and genome evolution in two generalist and specialist microsporidian pathogens of
mosquitoes. *Nat Commun* 2015; 6: 7121.

[22] Biron DG, Agnew P, Marche L, et al. Proteome of Aedes aegypti larvae in response to
infection by the intracellular parasite Vavraia culicis. *Int J Parasitol* 2005; 35: 1385–1397.

[23] Sy VE, Agnew P, Sidobre C, et al. Reduced survival and reproductive success generates
selection pressure for the dengue mosquito Aedes aegypti to evolve resistance against infection
by the microsporidian parasite Vavraia culicis. *Evol Appl* 2014; 7: 468–479.

[24] Silva LM. Selection on time to parasite transmission shapes the host Anopheles gambiae
transcriptional response and suggests immune evasion. *bioRxiv* 2024; 2009–2024.

[25] Agnew P, Berticat C, Bedhomme S, et al. Parasitism increases and decreases the costs of
insecticide resistance in mosquitoes. *Evolution (N Y)* 2004; 58: 579–586.

[26] Bargielowski I, Koella JC. A possible mechanism for the suppression of Plasmodium berghei

development in the mosquito Anopheles gambiae by the microsporidian Vavraia culicis. *PLoS*
One 2009; 4: e4676.

[27]   Lorenz LM, Koella JC. The microsporidian parasite Vavraia culicis as a potential late life–
263   acting control agent of malaria. *Evol Appl* 2011; 4: 783–790.

264   [28]   Herren JK, Mbaisi L, Mararo E, et al. A microsporidian impairs Plasmodium falciparum
265   transmission in Anopheles arabiensis mosquitoes. *Nat Commun* 2020; 11: 1–10.

266   [29]   Murareanu BM, Antao N V, Zhao W, et al. High-throughput small molecule screen identifies
267   inhibitors of microsporidia invasion and proliferation in C. elegans. *Nat Commun* 2022; 13:
268   5653.

269   [30]   Vejpongsa P, Yeh ETH. Prevention of anthracycline-induced cardiotoxicity: challenges and
270   opportunities. *J Am Coll Cardiol* 2014; 64: 938–945.

271   [31]   Hasinoff BB, Kuschak TI, Yalowich JC, et al. A QSAR study comparing the cytotoxicity and
272   DNA topoisomerase II inhibitory effects of bisdioxopiperazine analogs of ICRF-187
273   (dexrazoxane). *Biochem Pharmacol* 1995; 50: 953–958.

274   [32]   Eneh C, Lekkala MR. Dexrazoxane.

275   [33]   Vulule JM, Beach RF, Atieli FK, et al. Reduced susceptibility of Anopheles gambiae to
276   permethrin associated with the use of permethrin-impregnated bednets and curtains in Kenya.
277   *Med Vet Entomol* 1994; 8: 71–75.

278   [34]   Andreadis TG. Microsporidian parasites of mosquitoes. *J Am Mosq Control Assoc* 2007; 23:
279   3–29.

280   [35]   Fukuda T, Willis OR, Barnard DR. Parasites of the Asian tiger mosquito and other container-
281   inhabiting mosquitoes (Diptera: Culicidae) in northcentral Florida. *J Med Entomol* 1997; 34:
282   226–233.

283   [36]   Kulma K, Saddler A, Koella JC. Effects of age and larval nutrition on phenotypic expression
284   of insecticide-resistance in Anopheles mosquitoes. *PLoS One* 2013; 8: e58322.

285   [37]   Dao A, Kassogue Y, Adamou A, et al. Reproduction-longevity trade-off in Anopheles gambiae
286   (Diptera: Culicidae). *J Med Entomol* 2010; 47: 769–777.

287   [38]   Team RC. RA language and environment for statistical computing, R Foundation for
288   Statistical. *Computing*.

[39]   Hartig F, Hartig MF. Package 'DHARMa'. *R Packag*.

[40]   Fox J, Weisberg S, Adler D, et al. Package 'car'. *Vienna R Found Stat Comput*; 16.

[41] Brooks ME, Kristensen K, van Benthem KJ, et al. glmmTMB balances speed and flexibility
among packages for zero-inflated generalized linear mixed modeling. *R J* 2017; 9: 378–400.

[42] Lenth R, Singmann H, Love J, et al. Package 'emmeans'. *R Packag version*; 1.

[43] Hothorn T, Bretz F, Westfall P, et al. Package 'multcomp'. *Simultaneous inference Gen*
Parametr Model Proj Stat Comput Vienna, Austria.

[44] Loyevsky M, Sacci Jr JB, Boehme P, et al. Plasmodium falciparumandPlasmodium yoelii:
Effect of the Iron Chelation Prodrug Dexrazoxane onin VitroCultures. *Exp Parasitol* 1999; 91:
105–114.

[45] Santos D, Gomes HF, Ribeiro L, et al. Inhibition of Aedes aegypti DNA topoisomerase II by
etoposide: Impact on survival and morphology of larvae and pupae. *Comp Biochem Physiol*
Part C Toxicol Pharmacol 2021; 247: 109066.

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Author contributions

307 LMS and TGZ designed and conducted the experiments, analysed the data and wrote the manuscript.

Data availability

309 All data generated or analysed during this study are included as Supplementary Information files.

Additional Information

311 The authors declare no competing interests.

Figure legends

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