1 Localisation and tissue tropism of the symbiont Microsporidia MB in the germ line

2 and somatic tissues of Anopheles arabiensis

- 3 Edward E. Makhulu^{†a}, Thomas O. Onchuru^{†a,b}, Joseph Gichuhi^a, Fidel G. Otieno^a, Anne
- 4 W. Wairimu^a, Joseph .N. Muthoni^a, Lizette Koekoemoer^c and Jeremy K. Herren^{1#}
- 5
- ⁶ ^aInternational Centre of Insect Physiology and Ecology (ICIPE), Kasarani, Nairobi, Kenya.
- ⁷ ^bDepartment of Physical and Biological Sciences, Bomet University College, Bomet,
- 8 Kenya.
- 9 ^cWits Research Institute for Malaria, University of the Witwatersrand, South Africa.
- 10
- 11 Short title: Tissue tropism of *Microsporidia MB* in *An. arabiensis*
- 12
- 13 **#Corresponding author:** Jeremy K. Herren, <u>iherren@icipe.org</u>

14 Abstract

The Anopheles symbiont, Microsporidia MB, is maternally inherited and has a strong malaria transmission-blocking phenotype in Anopheles arabiensis. Microsporidia MB is also vertically transmitted, sexually transmitted and avirulent. These characteristics are expected to promote its spread through mosquito populations, enhancing the potential of *Microsporidia MB* as a candidate for the development of a symbiont-mediated malaria transmission blocking strategy. We found that the patterns of *Microsporidia MB* localisation over the development of *An. arabiensis* indicate accumulation in tissues 22 linked to its transmission, specifically the male and female gonadal tissues. Transovarial vertical transmission of Microsporidia MB occurs in the female An. arabiensis ovary when 23 24 *Microsporidia MB* becomes localised to the cytoplasm of the developing oocyte. In male An. arabiensis, Microsporidia MB is localised in the testis and vas deferens. Notably, a 25 high intensity of *Microsporidia MB* can also be observed in the *An. arabiensis* adult but 26 27 not larval gut. The levels of *Microsporidia MB* found in the female ovary are linked to the 28 progression of oogenesis, increasing after blood feeding initiates the development of 29 eggs. The levels of *Microsporiodia MB* in the male and female gonadal and gut tissue do 30 not increase as mosquitoes age. Altogether, the high specificity of Microsporidia MB tissue localisation patterns and changes in infection prevalence and intensity suggest 31 32 adaptation to maximise transmission and avirulence in Anopheles arabiensis.

33

34 Importance

35 *Microsporidia MB* is a symbiont with strong malaria transmission-blocking phenotype in Anopheles arabiensis. It spreads in mosquito populations through mother-to-offspring 36 37 and sexual transmission. The ability of *Microsporidia MB* to block *Plasmodium* 38 transmission together with its ability to spread within Anopheles populations and its 39 avirulence to the host makes it a very attractive candidate for developing a key strategy 40 to stop malaria transmissions. Here, we report the basis of Microsporidia MB 41 transmission. We find that Microsporidia MB accumulates in Anopheles arabiensis tissues 42 linked to its sexual and vertical transmission. Its prevalence and intensity in the tissues 43 over the mosquito life cycle suggest adaptation to maximise transmission and avirulence

in *Anopheles arabiensis*. These findings provide the foundation for understanding the
 factors that affect *Microsporidia MB* transmission efficiency. This will contribute to the
 establishment of strategies to maximize *Microsporidia MB* transmission for *Anopheles* mosquito population replacement and malaria transmission blocking.

48

49 **KEYWORDS:** *Microsporidia MB*, Malaria vectors, symbiotic microbes

50

51 Introduction

52 The malaria disease burden remains a major impediment to good health and economic development in many regions of sub-Saharan Africa (SSA). In 2021, a total of 247 million 53 54 cases were reported that resulted in 619,000 deaths, a strong indication that current 55 control measures and their deployment levels are insufficient (1). Large-scale insecticide 56 treated net (ITN) distribution campaigns have had a significant impact on reducing the number of malaria cases (2). However, many malaria vectors have now developed 57 58 resistance to the insecticides used in ITNs (3) and they are increasingly biting outdoors, 59 where nets offer no protection (4). In addition, many malaria control efforts were 60 significantly disrupted by the COVID-19 pandemic (5) and the recent invasion of An. 61 stephensi across Africa is leading to a significant rise in malaria transmission (6-8). 62 Altogether, these factors threaten to reverse the gains achieved for malaria reduction and indicate an urgent need for new strategies to control Anopheles mosquito populations or 63 64 their capacity to transmit *Plasmodium* parasites.

65 A novel and potentially transformative method of controlling vector-borne disease 66 involves the use of transmission blocking symbionts. For dengue, a control strategy based on the transmission blocking symbiont Wolbachia has been highly effective and 67 controlled field trials are currently implemented in over 13 countries (9). A similar strategy, 68 69 based on a *Plasmodium* transmission-blocking symbiont in *Anopheles* mosquitoes could 70 be transformative for controlling malaria. The microsporidian symbiont, *Microsporidia MB*, 71 is naturally found in anopheline mosquito populations and has been shown to block 72 Plasmodium development (10). In addition, Microsporidia MB is both vertically and 73 sexually transmitted (11). In conjunction with the *Plasmodium* blocking, the high efficiency 74 of vertical and sexual transmission of *Microsporidia MB*, which could facilitate the spread 75 and maintenance of *Microsporidia MB* in *Anopheles* mosquito populations, has led to the 76 suggestion that this symbiont could be deployed as a new tool for malaria control (12).

77 The success of a *Microsporidia MB*-based malaria control strategy will depend on efficient 78 vertical and horizontal transmission of the symbiont, which would enable the spread and 79 maintainance of Microsporidia MB in Anopheles vector populations. For symbionts that 80 are vertically and sexually transmitted, avirulence towards the host can be expected (13). 81 Vertically and sexually transmitted symbionts can be selected for their ability to enhance 82 host fitness (14, 15). The tissue level localisation pattern and intensity of infection have 83 been shown to play an important role in symbiont transmission and host fitness effects. 84 Currently, little is known regarding Microsporidia MB localization patterns in infected 85 mosquitoes. Here, we investigated the tissue-level localisation and changes of 86 Microsporidia MB infection intensity across development of An. arabiensis. We found that 87 Microsporidia MB was predominately found in the reproductive organs of An. arabiensis

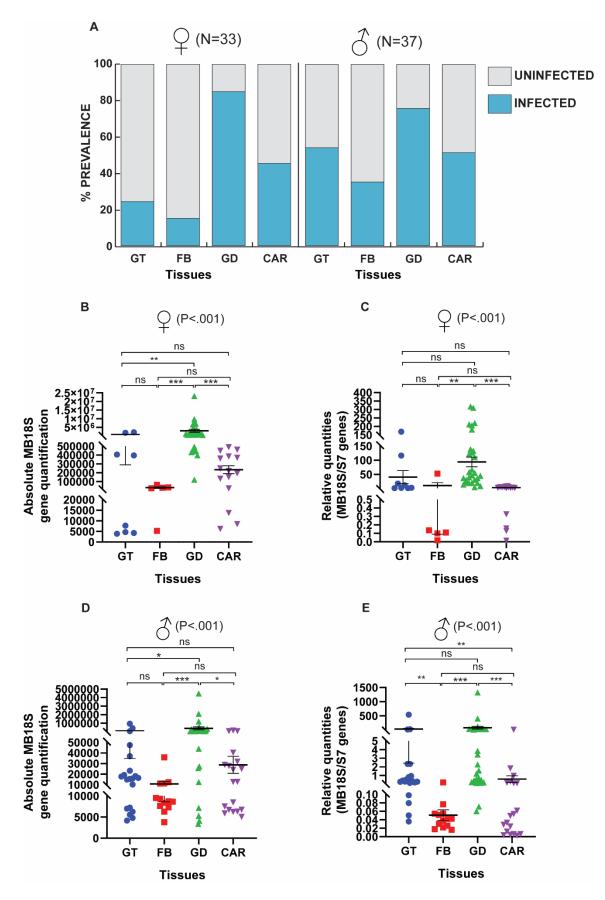
88 males and females. Additionally, we observed *Microsporidia MB* inside developing 89 oocytes in the An. arabiensis ovaries, indicating transovarial vertical transmission. 90 Interestingly, the intensity of *Microsporidia MB* infection in the female ovaries increased 91 The prevalence and intensity of *Microsporidia MB* infection in the after blood feeding. 92 female ovaries was found to decrease as An. arabiensis mosquitoes aged. In male An. 93 arabiensis, Microsporidia MB is found in the testis and vas deferens offering further 94 confirmation of the basis of male to female sexual transmission as previously reported 95 (11). *Microsporidia MB* is also in found in the *An. arabiensis* adult intestine at moderate 96 prevalences. Notably, *Microsporidia MB* is always absent from the larval intestine but 97 present in the larval body.

98

99 **Results**

The *An. arabiensis* male and female gonads are the primary site of *Microsporidia MB* infection in *An. arabiensis*

102 The prevalence and intensity of *Microsporidia MB* was investigated in the gonads, gut fat-103 body and carcass of seven-day-old adult An. arabiensis by gPCR (Figure 1). We show 104 that An. arabiensis male and female gonads had the highest Microsporidia MB prevalence 105 (Figure 1A) and the highest absolute intensity of *Microsporidia MB* infection in comparison 106 to other tissues and the carcass (Figures 1B-1D). In female and male An. arabiensis, the 107 gonads also had higher mean *Microsporidia MB* intensities relative to host gene copy 108 number (relative Microsporidia MB intensity) than fat bodies and carcasses. These 109 findings indicate that the primary site of *Microsporidia MB* infection in both male and 110 female adult An. arabiensis was the gonadal tissue. This is in line with a study that 111 investigated *Microsporidia MB* in only male *An. arabiensis* (11). A moderate prevalence 112 of Microsporidia MB infection was observed in the An. arabiensis male gut (54%, Figure 113 1A). In addition, the mean relative *Microsporidia MB* intensity in the male gut was 114 significantly higher than the fat body and carcass (Figure 1E), indicating that 115 Microsporidia MB density can reach high levels in this tissue. In females, the relative 116 intensity of *Microsporidia MB* in the gut was not significantly different from the relative 117 intensity in the gonads (Figure 1C). Altogether, these findings suggest that the gut is likely 118 to be the secondary site of infection. In both sexes, the prevalence and intensity of 119 *Microsporidia MB* was lowest in the fat body (Figure 1A-E).



121 Figure 1. *Microsporidia MB* infection prevalence and intensity in adult *An. arabiensis* tissues. (A)

The prevalence of *Microsporidia MB* is highest in the gonads of male and female *An. arabiensis* in comparison to other somatic tissues and the carcass. The absolute *Microsporidia MB* intensity was higher in female (**B**) (Dunn's post hoc test, GT vs GD P =0.008, FB vs GD P<0.001, CAR vs GD P<0.001) and male (**C**) (Dunn's post hoc test, GT vs GD P =0.019, FB vs GD P<0.001, CAR vs GD P =0.011) gonads relative to somatic tissues. (**D**) The relative intensity of Microsporidia MB in the female *An. arabiensis* midgut is not significantly different from the gonads. (**E**) The relative *Microsporidia MB* intensity determined by qPCR was significantly higher in the male gut in comparison to the fat body and carcass.

129 (Abbreviations: **GT** – Gut, **FB** – Fat bodies, **GD** – Gonads, **CAR** – Carcass)

130 *Microsporidia MB* cells are identified by microscopy in oocytes in female and testis

131 and gut in male *An. arabiensis*

132 To better understand where *Microsporidia MB* is located inside the male and female *An*. arabiensis gonads and guts, we visualised these tissues using Fluorescent In-Situ 133 134 Hybridization (FISH) and confocal microscopy. The visualisation of Microsporidia MB 135 confirmed that male and females An. arabiensis gonads were the primary site of infection 136 (Figure 2A-2B). In the An. arabiensis female gonad, Microsporidia MB is observed in all stages of egg chamber development as well as the germ-line stem cells (Figure 2A). As 137 138 the oocytes enter vitellogenesis, higher intensities of *Microsporidia MB* are observed. In 139 the male gonad, *Microsporidia MB* is observed in the testis and vas deferens, in close 140 proximity to developing sperm cells (Figure 2B). In the An. arabiensis male gut, 141 Microsporidia MB is occasionally observed at high infection levels in a small number of 142 cells in the midgut and hindgut regions (Figure 2C).

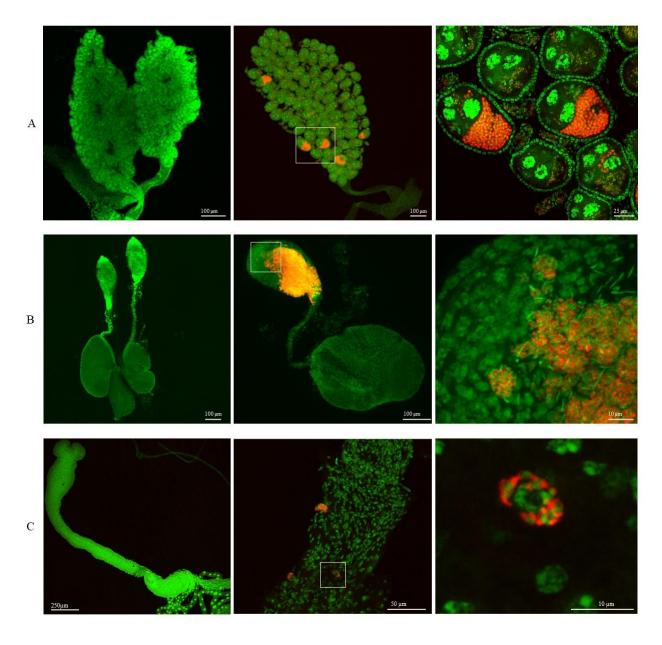


Figure 2. *Microsporidia MB* localisation to cells in the male and female gonads and male gut. *Microsporidia MB* infection is observed by FISH and confocal fluorescence microscopy in the gonads and gut of *An. arabiensis* mosquitoes. (A) *Microsporidia MB* in the female *An. arabiensis* gonads. A high density of *Microsporidia MB* is observed in vitellogenic oocytes. (B) *Microsporidia MB* in the male *An. arabiensis* gonad. A high density of *Microsporidia MB* is observed in parts of the testis. (C) *Microsporidia MB* is observed in the *An. arabiensis* male midgut cells.

150 *Microsporidia MB* is not found in the larval gut and is higher in female adult 151 mosquitoes

152 To better understand where *Microsporidia MB* is located during the larval stage of *An* 153 arabiensis, L4 larval stages were dissected to separate the gut from the carcass, with 154 carcasses containing all the remaining larval tissue, including the hemolymph. We 155 observed that none of the gut samples were infected with *Microsporidia MB* (Figure 3A). 156 However, larval carcasses were found to have a high prevalence of *Microsporidia MB* infection. The absolute intensity of Microsporidia MB infection in An. arabiensis larvae 157 158 was significantly less than that observed in adult female but not male mosquitoes (Figure 159 3B).

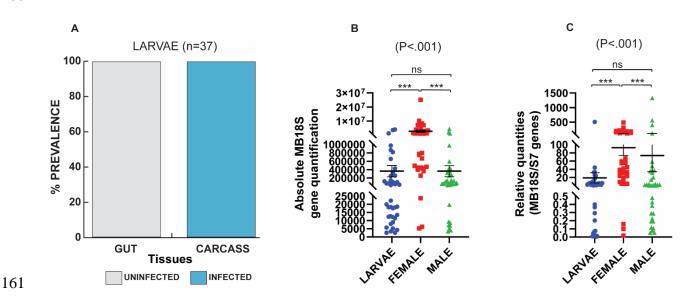
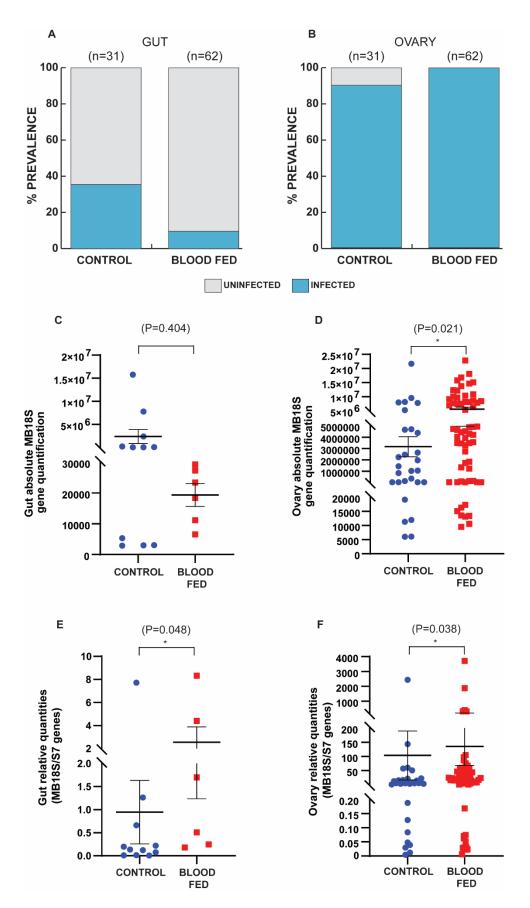


Figure 3. Localisation of *Microsporidia MB* in larval tissues and comparison of its density between larvae and adult mosquito stages. (A) *Microsporidia MB* is present in the carcass of dissected L4 larvae but not in the gut. *Microsporidia MB* infection intensity is only higher in *An. arabiensis* adult females relative to L4 larvae both in an absolute quantification (P<0.001) (B) and a relative quantification (P<0.001) (C) upon conducting a pairwise comparisons using Dunn's test.

167 Blood meal affects the intensity of *Microsporidia MB* in the ovaries

168 In An. arabiensis, oogenesis begins after female mosquitoes eclose but is arrested until 169 females take a blood meal. To determine if the intensity of *Microsporidia MB* in An. 170 arabiensis is affected by the onset of egg development and other physiological changes 171 associated with taking a blood meal, we compared the prevalence and intensity of 172 Microsporidia MB infection in the guts and gonads of An. arabiensis females that had fed 173 on a blood meal with those that had not fed on a blood meal. In the An. arabiensis female 174 gut, the prevalence of *Microsporidia MB* was slightly lower in blood fed mosquitoes 175 relative to non-blood fed mosquitoes (Figure 4A). While there was a significantly higher 176 relative intensity of *Microsporidia MB* in female An. arabiensis guts that had been blood 177 fed, the absolute levels of *Microsporidia MB* were not significantly different between the 178 two groups (Figure 4C and 4E). This discrepancy is likely the result of the turnover (and 179 shedding) of gut epithelial cells that is known to occur after blood feeding, which could 180 decrease the amount of host DNA in this tissue (16,17). The prevalence of *Microsporidia* 181 *MB* infection was high in ovaries regardless of the blood feeding status (Figure 4B). 182 However, we observed significantly higher relative and absolute intensities of 183 Microsporidia MB in gonads of blood fed An. arabiensis (Figure 4D and 4F). These 184 observations suggests that the amount and density of *Microsporidia MB* in the female 185 gonad increases as the gonotrophic cycle progresses and egg development is initiated.



188 Figure 4. Microsporidia MB prevalence and intensity in female An. arabiensis gonads is affected by 189 blood feeding. (A) The prevalence of Microsporidia MB infection is slightly lower in the guts of blood fed 190 An. arabiensis females. (B) The prevalence of Microsporidia MB infection is high in blood fed and non-191 blood fed An. arabiensis female gonads. (C) The absolute intensity of Microsporidia MB infection is not 192 significantly different between the guts of blood fed and non-blood fed An. arabiensis females (two-tailed 193 Mann–Whitney test, P = 0.404, error bars reflect SEM). (D) The absolute intensity of Microsporidia MB 194 infection is significantly higher in the gonads of blood fed An. arabiensis females (two-tailed Mann-Whitney 195 test, P = 0.021, error bars reflect SEM). (E) The relative intensity of Microsporidia MB infection is 196 significantly higher in the guts of blood fed An. arabiensis females (two-tailed Mann–Whitney test, P = 0.048, 197 error bars reflect SEM). (F) The relative intensity of Microsporidia MB infection is higher in the gonads of 198 blood fed An. arabiensis females (two-tailed Mann–Whitney test, P = 0.038, error bars reflect SEM).

199

Age affects the intensity of *Microsporidia MB* density in *An. arabiensis* gonads and guts

202 To investigate the effect of aging on the prevalence and intensity of *Microsporidia MB* in An. arabiensis gonadal and gut tissue, male and female An. arabiensis were aged for 2, 203 204 7 and 14 days prior to dissection of tissues and quantification of *Microsporidia MB* by 205 qPCR. The prevalence of *Microsporidia MB* in the guts of female An. arabiensis was 206 slightly higher in the 14-day old mosquitoes relative to 2- and 7-day old mosquitoes 207 (Figure 4A). The prevalence of *Microsporidia MB* in the guts of male *An. arabiensis* did 208 not change as mosquitoes were aged (Figure 4B). In both males and females, there was 209 a general trend towards a decrease in the prevalence of *Microsporidia MB* in gonads as 210 mosquitoes were aged (Figure 4A-4B). There was a slight increase in the relative 211 intensity, but not absolute intensity of Microsporidia MB in the guts of female An.

212 arabiensis between 2- and 14 days (Figure 4C-4D). The intensity of Microsporidia MB in 213 the guts of male An. arabiensis was apparently not affected by aging (Figure 4G-4H). The 214 relative intensity of Microsporidia MB in female An. arabiensis gonads increased between 215 day 2 and 7, however, the absolute levels remained constant (Figure 4E-4F). This 216 suggests that the increase in relative *Microsporidia MB* intensity is a consequence of host 217 DNA content decreasing. Notably, the absolute, but not relative, intensity of *Microsporidia* 218 MB was lower in the gonads of 14 day old females relative to 2 and 7 day old females 219 (Figure 4E-4F). The relative intensity of *Microsporidia MB* in the *An. arabiensis* male 220 gonad increased between 2 and 7 days of aging (Figure 4I). The absolute intensity of 221 Microsporidia MB in the male gonad was not significantly different across all ages (Figure 222 4J).

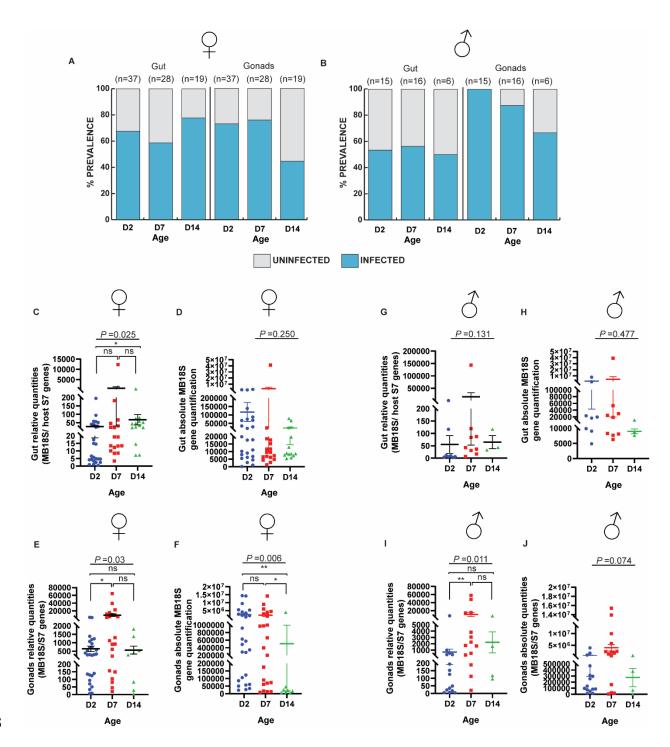


Figure 5. The intensity of *Microsporidia MB* in the *An. arabiensis* gonad and gut is affected by age. (A) The prevalence of *Microsporidia MB* in the guts of female *An. arabiensis* was slightly higher at 14 days relative to seven and two days. A decrease in the prevalence of *Microsporidia MB* is observed in the gonads of 14-day old female *An. arabiensis*. (B) The prevalence of *Microsporidia MB* in the guts of male *An.*

228 arabiensis did not change as mosquitoes were aged. A decrease in the prevalence of Microsporidia MB is 229 observed in the gonads of 14-day old male An. arabiensis. (C-D) An increase in the relative intensity (Dunn's 230 post hoc test, P =0.022), but not absolute intensity, of Microsporidia MB is observed in the guts of female 231 An. arabiensis between 2- and 14 days. (E-F) The relative intensity (Dunn's post hoc test, P =0.046), but 232 not absolute intensity, of Microsporidia MB in female An. arabiensis gonads increased between day two 233 and seven. The absolute intensity of Microsporidia MB was lower in the gonads of 14-day old females 234 relative to (Dunn's post hoc test, P =0.005) and seven (Dunn's post hoc test, P =0.022) females. (G-H) The 235 intensity of Microsporidia MB in the guts of male An. arabiensis does not change over aging. (I-J) The 236 relative intensity of Microsporidia MB in the An. arabiensis male gonad increased between two and seven 237 days of aging (Dunn's post hoc test, P =0.008). The absolute intensity of Microsporidia MB in the male 238 gonad was not significantly different across all ages.

239 **Discussion**

240 The results demonstrate that the gonads of male and female An. arabiensis are the primary sites of Microsporidia MB infection. This finding indicates that maternal and 241 242 sexual transmission are likely to be the most important *Microsporidia MB* infection routes. 243 Both maternal and sexual transmission require that An. arabiensis hosts do not incur 244 excessive fitness costs when infected with Microsporidia MB (13). In line with this, 245 Microsporidia MB does not have a major effect on the fitness of larvae or adult female 246 mosquitoes (10). It remains unresolved how *Microsporidia MB* can reach very high 247 intensities, up to 2.5x10⁷ copies of the *Microsporida MB* 18S gene per mosquito, without 248 negatively impacting host fitness.

Other insect endosymbionts, including *Wolbachia*, have intensity-dependent effects on host fitness (18). It is therefore possible that very high intensity *Microsporidia MB* infections do have a fitness cost, but this would need further investigation. The link

252 between Wolbachia intensity and host fitness is complex and, in many cases, fitness is 253 not affected by high endosymbiont intensities (19). It is likely that the pattern of 254 endosymbiont localisation in host tissues has important fitness consequences for 255 endosymbionts such as Wolbachia and Microsporidia MB. A strain of Wolbachia, 256 *w*MelPop, that over proliferates and decreases host fitness has been found at high 257 intensities in the central nervous system and muscles of its insect hosts (20, 21). It is 258 possible that endosymbionts mitigate effects on host fitness by limiting most of the 259 infection and proliferation to certain tissues.

Our findings indicate that *Microsporidia MB* is primarily found in gonadal tissue and this is suggestive adaptation to minimise host fitness costs without compromising transmission. Fluorescence confocal microscopy revealed that *Microsporidia MB* was present in oocytes and nurse cells across all stages of oogenesis. These findings clearly demonstrate that *Microsporidia MB* maternal transmission is transovarial, which explains the need for higher *Microsporidia MB* intensities to be maintained in the female gonads.

In the male germline, *Microsporidia MB* is primarily localised to the testis and vas deferens. The testis is the site of sperm production, comprising of a proximal end with stem cell divisions, spermatocysts in different stages of development and a distal spermatozoa reservoir (22). In *An. arabiensis*, we observed *Microsporidia MB* cells as clusters in the spermatocyst and spermatozoa reservoir regions of the testis and more rarely in the stem cell region. These findings suggest that *Microsporidia MB* is packaged with spermatozoa at their early development stages. 273 We observed very low levels of *Microsporidia MB* in the *An. arabiensis* adult fat body. 274 Pathogenic Microsporidians are known to infect and proliferate in the insect fat body (22). 275 It is likely that as the primary energy storage tissue, fat bodies provide the nutrients 276 required for this proliferation. The low levels of *Microsporidia MB* in the adult fat body of 277 An. arabiensis could be part of the explanation for its avirulence in An. arabiensis. The 278 moderate levels of *Microsporidia MB* in the carcass samples, which contained all the 279 tissues apart from gut, gonads and fat body, suggest that we cannot exclude there being 280 another *Microsporidia MB* infection reservoir in adult *An. arabiensis*.

281 In the adult An. arabiensis gut, there were a small proportion of samples which had very 282 high intensities of *Microsporidia MB*. The significance of these high intensity gut infections 283 is yet to be resolved. Fluorescence confocal microscopy revealed that *Microsporidia MB* 284 was found at very high densities in a small subset of cells in the adult male mid and 285 hindgut. The shape and positioning of these cells suggests that they could be intestinal 286 stem cells (17). It is notable that in contrast to adults, *Microsporidia MB* was never 287 observed in the larval gut. A possible explanation for this is that presence in the larval gut 288 could have prohibitive fitness cost since larvae feed continuously. In addition, the total 289 intensity of *Microsporidia MB* is lower in larvae relative to adult females.

We also investigated how localisation and *Microsporidia MB* intensity changes over the life cycle of *An. arabiensis*. In *An. arabiensis*, oogenesis starts when mosquitoes elcose, but is then paused until females take a blood meal (24). Once females take a blood meal, several physiological changes occur that enable the blood to be digested an ultimately to meet nutritional demands of egg production (25). Since there are major changes occurring in the gut and gonads of females post blood feeding, we investigated the 296 changes in the levels of *Microsporidia MB* in these tissues. We observed that the levels 297 of *Microsporidia MB* in the gut did not change in response to blood feeding; the relative 298 intensity of *Microsporidia MB* did increase in the gut after blood feeding but the absolute 299 intensity appeared to be constant, which suggests a decrease in host nuclear gene copy 300 number. This decrease is likely to be linked to the process of gut epithelium renewal (17). 301 In contrast, we observed a marked increase in the intensity of *Microsporidia MB* in the 302 female gonad after blood feeding. Notably, Microsporidia MB is observed at high 303 intensities in the later (vitellogenic) stages of oogenic development, which develop after 304 the blood meal induced initiation of oogenesis (24). Therefore, the proliferation of 305 Microsporidia MB in vitellogenic oocytes is a possible explanation for the increase in 306 intensity of *Microsporidia MB* after blood feeding.

307 We did not observe Microsporidia MB levels increasing as male or female guts and 308 gonads as An. arabiensis mosquitoes are aged. This pattern of growth is consistent with 309 symbionts that regulate proliferation to mitigate fitness costs (26). While there was an 310 increase in the relative intensity of *Microsporida MB* in male and female gonads between 311 day 2 and day 7, this is likely to be the consequence of a decrease in host nuclear gene 312 copy number. Notably, there was a significant decrease in the prevalence and absolute 313 intensity of *Microsporidia MB* in female gonads between 7 and 14 days of aging. This may 314 be indicating a loss of *Microsporidia MB* from germline cells after numerous rounds of 315 division and an inability for these cell to be re-infected. However, the aged An. arabiensis 316 female mosquitoes had not been blood fed and therefore had not completed any 317 gonotrophic cycles which could be expected to lower the rate of stem cell division (27).

Overall, our findings suggest that *Microsporidia MB* localisation patterns may serve to maximise transmission success while minimising negative effects on host fitness. *Microsporidia MB* has proliferation and localisation patterns that are consistent with a coevolved symbiont that exploits sexual and vertical transmission routes. A better understanding of changes in localisation and symbiont intensity across the *An. arabiensis* lifecycle will contribute to improving *Microsporidia MB* infected mosquito rearing and aid in the formulation of strategies to disseminate *Microsporidia MB*.

325 Materials and methods

326 Mosquito sample collection

Wild-caught gravid adult female *Anopheles gambiae s.l.*, were collected using mouth aspiration. Collected mosquitoes were identified morphologically to the species level and sub-species identification of *Anopheles gambiae s.l.*, was carried out by PCR (28). Sampling was conducted from one site in Kenya, the Ahero irrigation scheme (– 34.9190W, –0.1661N). The mosquitoes used in this study were collected between September 2021 and October 2021. Sampled mosquitoes were transported to the *icipe* -Duduville campus in Nairobi and maintained on 6% glucose and water.

334 Mosquito processing and development of isofemale lines

Wild collected mosquitoes were maintained in an insectary at 27 ± 2.5 °C, humidity 60– 80% and 12-h day and 12-h night cycles and induced to oviposit in individual microcentrifuge tubes containing a wet 1 cm × 1 cm Whatman filter paper. Eggs from each female were counted under a compound microscope using a paint brush and then 339 dispensed into water troughs for larval development at 30.5°C and 30–40% humidity. 340 Tetramin[™] baby fish food was used to feed developing larvae. Upon laying eggs, the 341 G₀ females were screened for presence of *Microsporidia MB* by PCR. The larval offspring 342 of *Microsporidia MB* positive field-caught female mosquitoes were transferred into larval rearing troughs for further development. Upon reaching the L4 larval stage, representative 343 344 samples from each positive isofemale line were screened to determine the Microsporidia 345 *MB* infection status of the line. Were mosquitos were maintained as adults, this was done 346 in an insectary at 30°C, with a relative humidity of 75% and 12-h day and 12-h night cycles 347 and a feeder with 6% glucose solution.

348 Larval and adult dissection

349 An. arabiensis G1 larvae and adult mosquitoes were dissected using forceps under the 350 Zeiss Stemi 2000-C stereomicroscope to obtain G1 L4 larval and adult Anopheles tissues. 351 During the dissection of the L4 larvae, alive samples were placed on a drop of 1X 352 Phosphate Buffered Saline (1X PBS). The larvae were restrained at the junction between 353 the head and thorax with a pair of forceps. A dissecting needle pin was then used to probe 354 the intersection between the second and last abdomen segments. The siphon and saddle 355 of the larvae were held and pulled gently to obtain the gut of the larvae using a new pair 356 of forceps. The carcass and the gut were placed in separate 1.5 micro-centrifuge tubes 357 containing 20µl of 1X PBS and 0.5mm zirconium beads for DNA extractions. For adults, 358 G1 mosquito samples were first anaesthetised for a few minutes until immobilised by 359 aspirating them in 1.5 micro-centrifuge tubes and placing them in ice. Upon immobilisation, the mosquitoes were placed in a drop of 1X PBS on a glass slide. The 360 361 junction between the last and second-last abdomen was probed using a dissecting

needle. The last segment was gently pulled to release the gonads and the gut. The gut
and gonads were separated and placed in separate 1.5 micro-centrifuge tubes containing
20µl of 1X PBS and 0.5mm zirconium beads ready for nucleic acid extractions. Similarly,
the fat bodies and carcass (all remaining tissue) and were also separated and placed in
separate 1.5 micro-centrifuge tubes as described above. Forceps were sterilised after
every dissection to prevent contamination.

368

369 Mosquito blood feeding

370 Two to three-day-old *Microsporidia MB* positive and negative *An. arabiensis* mosquitoes 371 maintained at a temperature of 30°C, and a relative humidity of 75 % were starved for 2 372 hours without water or glucose in preparation for blood feeding. Membrane feeding was 373 conducted according to (29). Breifly, bovine blood was transferred into a Hemotek® 374 feeding apparatus (Hemotek, UK) with a Parafilm-A membrane set at. The Hemotek 375 feeding apparatus was placed on top of a cage of starved An. arabiensis mosquitoes and 376 feeding was allowed to take place for 1hr. Fully engorged An. arabiensis were transferred 377 to new cages, maintained for five days post-infection, and later dissected to obtain the 378 gut and ovaries which were screened for *Microsporidia MB* presence and levels.

379 Flourescent *in-situ* hybridisation (FISH) localisation

Dissected tissues were fixed overnight in 4% Paraformaldehyde (PFA) at 4 °C. After fixation, the samples were rehydrated in 50% ethanol for 30 minutes and transferred to 1X PBS for another 30 minutes. FISH was then conducted to localize *Microsporia MB* within the tissues. Hybridization was done by incubating the tissues in 100µl of hybridization buffer overnight at 50°C. The FISH hybridization mix contained hybridization buffer (dH2O, 5M NaCl, 1M Tris/HCl [pH=8], and 10% SDS), *Microsporidia MB* specific
CY5 probe (10), 0.5µM final concentration, and SYTOX Green general DNA staining.
After staining, the samples were washed twice with 100µl of prewarmed wash buffer
(dH2O, 5M NaCl, 1M Tris/HCl [pH=8], 0.5M EDTA, and 10% SDS). The tissues were then
placed on a slide and visualized immediately using a Leica SP5 confocal microscope
(Leica Microsystems, USA). The images were analyzed with the ImageJ 1.50i software
package (30).

392

393 DNA extraction and molecular detection and quantification of *Microsporidia MB*

394 DNA was extracted using the ammonium acetate protein precipitation method (10). 395 Extracted DNA samples were screened to determine their *Microsporidia MB* infection 396 status using the Microsporidia MB specific primers (MB18SF: CGCCGG 397 CCGTGAAAAATTTA and MB18SR: CCTTGGACGTG GGAGCTATC) targeting the 398 Microsporidia MB 18S rRNA region (10). The PCR reaction used in detection comprised 399 a 10 µl reaction consisting of 2 µl HOTFirepol Blend Master mix Ready-To-Load (Solis 400 Biodyne, Estonia), 0.5 µl of 5 pmol/µl forward and reverse primers, 2 µl of the DNA 401 template, and 5 µl nuclease-free PCR water was undertaken. The thermocycling 402 conditions employed were initial denaturation at 95°C for 15 min, followed by 35 cycles of 403 denaturation at 95°C for 1 min, annealing at 62°C for 90 s, and extension at 72°C for a 404 further 60 s. Final elongation was done at 72°C for 5 min. Samples positive for 405 Microsporidia MB were subjected to relative and absolute qPCR analysis to quantify 406 infection levels (10, 31). The qPCR analysis involved the MB18SF/ MB18SR primers, 407 normalised with ribosomal the Anopheles S7 (primers, S7F: gene

408 TCCTGGAGCTGGAGATGAAC and S7R: GACGGGTCTGTACCTTCTGG) as the 409 reference host gene. The 10µl PCR reaction consisted of 2µl HOT FIREPol® EvaGreen® 410 HRM no ROX Mix Solis qPCR Master mix (Solis Biodyne, Estonia), 0.5 µl of 5 pmol/µl 411 forward and reverse primers, 2 µl of the DNA template from Microsporidia MB positive 412 samples, and 5 µl nuclease-free PCR. The thermocycling conditions employed were an 413 initial denaturation at 95°C for 15 min, followed by 35 cycles of denaturation at 95°C for 1 414 min, annealing at 62°C for 90 s, and extension at 72°C for a further 60 s. Finally, melting 415 curves were generated by melting analysis with temperature ranges from 65°C to 95°C. 416 The PCR and qPCR was carried out on a MIC qPCR cycler (BioMolecular Systems, Australia). Each sample was confirmed to have the characteristic melt curve associated 417 418 with the *Microsporidia MB* MB18SF/ MB18SR primers.

419

420 Data analysis

421 All the data was tested for nomarlity using the Shapiro-Wilk test. For non-normal

422 unpaired data, a two tailed Mann–Whitney U test was used to determine the difference

423 between the groups. In cases with more than two data groups, we used Kruskal-Wallis

424 H test to determine the significance of differences between the groups. If significant, we

425 carried out a Dunn's post hoc test. All statistical analyses were preformed using

426 Graphpad Prizm version 6.0c software or R (version 8.0.2). P-values of *p < 0.05,

427 **p < 0.01, ***p < 0.001 and ****p < 0.0001 were deemed to be statistically significant.

428 Figures were created and/or edited using Adobe Illustrator (version 23.0.1).

429

430 **References**

431	1.	World Health Organization. 2022. The World Malaria Report 2022.
432	2.	Bhatt S, Weiss DJ, Cameron E, Bisanzio D, Mappin B, Dalrymple U, Battle KE,
433		Moyes CL, Henry A, Eckhoff PA, Wenger EA, Briët O, Penny MA, Smith TA,
434		Bennett A, Yukich J, Eisele TP, Griffin JT, Fergus CA, Lynch M, Lindgren F,
435		Cohen JM, Murray CLJ, Smith DL, Hay SI, Cibulskis RE, Gething PW. 2015. The
436		effect of malaria control on Plasmodium falciparum in Africa between 2000 and
437		2015. Nature 526:207–211.
438	3.	World Health Organization. 2018. Global report on insecticide resistance in
439		malaria vectors: 2010–2016.World Health Organization.
440	4.	Kabbale FG, Akol AM, Kaddu JB, Onapa AW. 2013. Biting patterns and
441		seasonality of Anopheles gambiae sensu lato and Anopheles funestus
442		mosquitoes in Kamuli District, Uganda. Parasites and Vectors 6:1–9.
443	5.	World Health Organization. 2021. The World Malaria Report 2021.
444	6.	Tadesse FG, Ashine T, Teka H, Esayas E, Messenger LA, Chali W, Meerstein-
445		Kessel L, Walker T, Wolde Behaksra S, Lanke K, Heutink R, Jeffries CL,
446		Mekonnen DA, Hailemeskel E, Tebeje SK, Tafesse T, Gashaw A, Tsegaye T,
447		Emiru T, Simon K, Bogale EA, Yohannes G, Kedir S, Shumie G, Sabir SA,
448		Mumba P, Dengela D, Kolaczinski JH, Wilson A, Churcher TS, Chibsa S, Murphy
449		M, Balkew M, Irish S, Drakeley C, Gadisa E, Bousema T. 2021. Anopheles
450		stephensi mosquitoes as vectors of Plasmodium vivax and falciparum, Horn of
451		Africa, 2019. Emerg Infect Dis 27:603.
452	7.	Ochomo EO, Milanoi S, Abong'o B, Onyango B, Muchoki M, Omoke D, Olanga E,
453		Njoroge L, Juma E, Otieno JD MD. 2023. Molecular surveillance leads to the first

454 detection of *Anopheles stephensi* in Kenya.

- 455 8. Mnzava, A., Monroe, A. C., & Okumu F. 2022. *Anopheles stephensi* in Africa
 456 requires a more integrated response. Malar J 21:1–6.
- 457 9. Utarini A, Indriani C, Ahmad RA, Tantowijoyo W, Arguni E, Ansari MR, Supriyati
- 458 E, Wardana DS, Meitika Y, Ernesia I, Nurhayati I, Prabowo E, Andari B, Green
- 459 BR, Hodgson L, Cutcher Z, Rancès E, Ryan PA, O'Neill SL, Dufault SM,
- 460 Tanamas SK, Jewell NP, Anders KL, Simmons CP. 2021. Efficacy of Wolbachia-
- 461 infected mosquito deployments for the control of Dengue. N Engl J Med
- 462 **384:2177-2186**.
- 463 10. Herren JK, Mbaisi L, Mararo E, Makhulu EE, Mobegi VA, Butungi H, Mancini MV,
- 464 Oundo JW, Teal ET, Pinaud S, Lawniczak MKN, Jabara J, Nattoh G, Sinkins SP.
- 465 2020. A microsporidian impairs *Plasmodium falciparum* transmission in

466 Anopheles arabiensis mosquitoes. Nat Commun 11:2187.

- 467 11. Nattoh G, Maina T, Makhulu EE, Mbaisi L, Mararo E, Otieno FG, Bukhari T,
- 468 Onchuru TO, Teal E, Paredes J, Bargul JL, Mburu DM, Onyango EA, Magoma G,
- 469 Sinkins SP, Herren JK. 2021. Horizontal transmission of the symbiont
- 470 *Microsporidia MB* in *Anopheles arabiensis*. Front Microbiol 12:647183.
- 471 12. Bukhari, T., Pevsner, R., and Herren JK. 2022. Microsporidia: a promising vector
- 472 control tool for residual malaria transmission. Front Trop Dis 3:66.
- 473 13. Lipsitch M, Siller S, Nowak MA. 1996. The evolution of virulence in pathogens
- 474 with vertical and horizontal transmission. Evolution (N Y) 50:1729–1741.
- 475 14. Knell RJ, Webberley KM. 2004. Sexually transmitted diseases of insects:
- 476 Distribution, evolution, ecology and host behaviour. Biol Rev Camb Philos Soc.

- 477 15. Moran NA. 2006. Symbiosis. Curr Biol 16:866–871.
- 478 16. Cui Y, Franz AW. 2020. Heterogeneity of midgut cells and their differential
- 479 responses to blood meal ingestion by the mosquito, Aedes aegypti. Insect
- 480 Biochem Mol Biol 127:103496.
- 481 17. Hixson B, Taracena ML, Buchon N. 2021. Midgut epithelial dynamics are central
- 482 to mosquitoes' physiology and fitness, and to the transmission of vector-borne
 483 disease. Front Cell Infect Microbiol 180.
- 484 18. Chrostek E, Marialva MSP, Esteves SS, Weinert LA, Martinez J, Jiggins FM,
- 485 Teixeira L. 2013. *Wolbachia* variants induce differential protection to viruses in
- 486 Drosophila melanogaster. A phenotypic and phylogenomic analysis. PLoS Genet
 487 9:e1003896.
- 488 19. López-Madrigal S, Duarte EH. 2019. Titer regulation in arthropod- *Wolbachia*489 symbioses. FEMS Microbiol Lett 366.
- 490 20. Min KT, Benzer S. 1997. Wolbachia, normally a symbiont of Drosophila, can be
- 491 virulent, causing degeneration and early death. Proc Natl Acad Sci U S A
- 492 **94:10792-10796**.
- Albertson R, Casper-Lindley C, Cao J, Tram U, Sullivan W. 2009. Symmetric and
 asymmetric mitotic segregation patterns influence *Wolbachia* distribution in host
 somatic tissue. J Cell Sci 122:4570–4583.
- 496 22. Huho BJ, Ng'habi KR, Killeen GF, Nkwengulila G, Knols BGJ, Ferguson HM.
- 497 2006. A reliable morphological method to assess the age of male *Anopheles*498 *gambiae*. Malar J 5:1–11.
- 499 23. Becnel JJ, Andreadis TG. 2014. Microsporidia in insects. Microsporidia Pathog

500 Oppor First Ed 521–570.

- 501 24. Mitchell SN, Catteruccia F. 2017. Anopheline reproductive biology: Impacts on
 502 vectorial capacity and potential avenues for malaria control. Cold Spring Harb
 503 Perspect Med.
- 504 25. Attardo GM, Hansen IA, Raikhel AS. 2005. Nutritional regulation of vitellogenesis
- in mosquitoes: Implications for anautogeny. Insect Biochem Mol Biol 35:661-675.
- 506 26. Herren JK, Paredes JC, Schüpfer F, Arafah K, Bulet P, Lemaitre B. 2014. Insect 507 endosymbiont proliferation is limited by lipid availability. Elife 3:e02964.
- 508 27. Valzania L, Mattee MT, Strand MR, Brown MR. 2019. Blood feeding activates the
- 509 vitellogenic stage of oogenesis in the mosquito *Aedes aegypti* through inhibition of
- 510 glycogen synthase kinase 3 by the insulin and TOR pathways. Dev Biol 454:85–
- 511 95.
- 512 28. Santolamazza F, Mancini E, Simard F, Qi Y, Tu Z, Della Torre A. 2008. Insertion
- 513 polymorphisms of SINE200 retrotransposons within speciation islands of
- 514 Anopheles gambiae molecular forms. Malar J 7:1–10.
- 515 29. Kajla MK, Barrett-Wilt GA, Paskewitz SM. 2019. Bacteria: A novel source for 516 potent mosquito feeding-deterrents. Sci Adv 5:eaau6141.
- 517 30. Schneider CA, Rasband WS, Eliceiri KW. 2012. NIH Image to ImageJ: 25 years of
 518 image analysis. Nat Methods 9:671-675.
- 519 31. Herren JK, Lemaitre B. 2011. Spiroplasma and host immunity: Activation of
- 520 humoral immune responses increases endosymbiont load and susceptibility to
- 521 certain Gram-negative bacterial pathogens in *Drosophila melanogaster*. Cell

522 Microbiol 13:1385-1396.

523

524 Acknowledgement

- 525 The study was supported by Open philanthropy (grant no.). The funders had no role in
- 526 study design, data collection, and analysis, decision to publish, or preparation of the
- 527 manuscript. We also thank the support provided by the *icipe* insectary team (Milcah
- 528 Gitau, Jeniffer Thiong'o, Peris Wambui, David Alila, and Charles Amara), field work
- 529 team (Gerald Ronoh and Robinson Kisero) and the project administrator Faith Kyengo.
- 530

531 Data accessibility

- 532 The following data will be submitted to the Dryad repository: *Microsporidia MB*
- 533 localization and prevalence in adult tissues, prevalence and density in larvae vs adults,
- 534 effect of bloodfeeding on prevalence and density of *Microsporidia MB*, density of
- 535 *Microsporidia MB* in the *An. arabiensis* gonad and gut across different ages.
- 536

537 Author contributions

- 538 EEM Conceptualization, data curation, validation, visualization, formal analysis,
- 539 investigation, methodology, writing original draft, writing review & editing.
- 540 TOO Data curation, validation, visualization, formal analysis, investigation,
- 541 methodology, supervision, writing original draft, writing review & editing.
- 542 JG, FGO, AWW, JNM Investigation, writing review & editing.
- 543 LK Conceptualization, supervision, writing review & editing.
- 544 JKH Conceptualization, data curation, formal analysis, funding acquisition,

545 methodology, supervision, validation, visualization, writing – original draft, writing –

546 review & editing.