1 *Wolbachia* inhibits ovarian formation and increases blood avidity in female *Aedes aegypti* – a
2 novel time-specific reproductive alteration
3

4 Meng-Jia Lau[1*]; Perran A. Ross[1]; Nancy M. Endersby-Harshman[1]; Qiong Yang[1]; Ary A.
5 Hoffmann[1]
6
7 [1] Pest and Environmental Adaptation Research Group, Bio21 Institute and the School of
8 BioSciences, The University of Melbourne, Parkville, Victoria, Australia
9 * Corresponding author, email: mengjia.liu@unimelb.edu.au; ORCID: 0000-0003-2943-7022

Abstract

Wolbachia, a gram-negative endosymbiotic bacterium widespread in arthropods, is well-
known for changing the reproduction of its host in ways that increase its rate of spread, but
there are also costs to hosts that can reduce this. Here we investigated a novel reproductive
alteration of *Wolbachia* *w*AlbB on its host *Aedes aegypti*, based on studies of mosquito life
history traits, ovarian dissections and gene expression assays. We found that an extended
period of the larval stage as well as the egg stage as previously shown can increase the
proportion of *Wolbachia*-infected females that become infertile; the impact of *Wolbachia* on
infertility therefore accumulated before pupation. We found that ovarian formation was
blocked in infertile females, even though infected females had relatively lower *Wolbachia*
densities than fertile infected females. Infertile females also showed a higher frequency of
blood feeding following a prior blood meal, indicating that they do not enter a complete
gonotrophic cycle. Treatments leading to infertility decreased expression of genes related to
reproduction. Our results demonstrate effects related to the causes and consequences of
infertile *w*AlbB-infected *Ae. aegypti* females with implications for *Wolbachia* releases and
have evolutionary implications for *Wolbachia* infections in novel hosts.
27

28 **Keywords:** *Wolbachia*; reproductive alteration; *Aedes aegypti*; ovarian formation

Introduction

Wolbachia, a Rickettsia-like maternally inherited endosymbiotic bacterium widespread in
insects [1], can alter insect reproduction in multiple ways to enhance its transmission [2].
Specific alterations include parthenogenesis – infected females can produce female offspring

without mating; feminization – infected genetic males are transformed into functional

females and produce female offspring; male killing – infected immature males die while

females survive, and cytoplasmic incompatibility (CI) – females infected with the

predominant *Wolbachia* strain obtain a fertility advantage in reproduction because males

infected with *Wolbachia* can cause female embryonic lethality when they mate with

uninfected females or females infected with a different *Wolbachia* strain [3, 4]. Aside from

these alterations that enhance the spread of *Wolbachia*, there can also be costs to the infection

that slow its spread [5-7]. These effects of *Wolbachia* are normally based on investigations of

natural *Wolbachia* strains that have existed in their hosts for many years, but are also

frequently found in artificial infections [8].

In the last decade, artificially introduced *Wolbachia* have been used in reducing the

transmission of arboviral diseases, with typical examples being the use of *Wolbachia w*Mel

[9] and *w*AlbB [10] in *Aedes aegypti* mosquitoes. Studies on native hosts tend to indicate

only a weak fitness cost of *Wolbachia* [6], but *Wolbachia* can have diverse fitness costs on its

new introduced host *Ae. aegypti,* such as life-shortening, heat sensitivity, quiescent egg

viability and reduced blood feeding success [11-14]. The effects of *Wolbachia* in native and

introduced hosts points to the possibility of a mutualistic relationship between *Wolbachia*

developing over time as a consequence of co-evolution and adaptation. In fact, it is

acknowledged that sometimes *Wolbachia* can act as both parasite and mutualist [5, 15, 16].

As a result, *Wolbachia* field release and establishment in new populations can be useful for

studying evolutionary changes in *Wolbachia* given that the invasion history of the population

is known [17-19].

In our previous study of the effects of *Wolbachia w*Mel and *w*AlbB on *Ae. aegypti* eggs, we

discovered a novel reproductive effect induced by *w*AlbB, in which infected females that

hatched from quiescent eggs lost their fertility [20]. Several other reproductive fitness costs

of *Wolbachia*, including those on the number and viability of eggs, can be attributed to the

consequence of nutritional competition between *Wolbachia* and its host [21] and represent

quantitative effects. However, in our study *Wolbachia*-infected females lost their fertility

entirely, indicating a qualitative effect difficult to attribute to nutritional competition [20], so

we suspect the involvement of other mechanisms.

In this paper, we investigate the causes and consequences of *Ae. aegypti* female infertility induced by *Wolbachia w*AlbB infection after egg quiescence. Firstly, we tested if the effects of *Wolbachia* on female infertility are permanent and dissected female ovaries. Secondly, we performed a larval starvation experiment combined with real-time PCR assays to understand the influence of *Wolbachia* on female development during different stages that could lead to infertility effects. Finally, we tested the blood avidity of fertile and infertile mosquitoes to see the impacts of the loss of ovaries on the rate of female blood feeding. Our study highlights the novel nature of reproductive effects altered by *Wolbachia w*AlbB in its new host *Ae. aegypti*, with implications for invasion of *Wolbachia* into uninfected populations and the evolution of mutualism more generally.

Materials and Methods

Mosquito populations

We used uninfected and *Wolbachia w*AlbB-infected *Aedes aegypti*. The uninfected population was derived from eggs collected in 2019 from regions in Cairns, Queensland, Australia where *Ae. aegypti* were not infected with *Wolbachia*. The *w*AlbB-infected population was generated by microinjection [22] and infected females were crossed to males from the uninfected population regularly [23, 24] to maintain a similar genetic background between populations. In following experiments, uninfected females were hatched from eggs stored for less than one month at 26 ± 1 °C as long-term storage does not influence the female fertility of uninfected populations [20]. However for consistency, uninfected females had the same rearing and feeding conditions as *w*AlbB-infected fertile and infertile females to which they were compared. *w*AlbB-infected females from non-stored eggs were hatched from freshly-laid eggs that had been dried and preserved for one week at 26 ± 1 °C. The process of mosquito rearing and "isolation and identification" is referred to Supplementary material 1.

Infertility across gonotrophic cycles

We hatched a mixture (approximately 1 : 1) of *w*AlbB-infected eggs that had been stored for two or four months, and did an "isolation and identification" for 200 females, fertile and infertile females were then provided with a second blood meal and 20 from each group were randomly selected for oviposition in 70 mL cups individually for another week to test if their fertility status was maintained.

Infertile female dissection and *Wolbachia* density

To examine ovarian development, fertile and infertile females after "isolation and identification" were dissected in saline (0.9% sodium chloride) and their ovaries were observed under a compound light microscope (Motic B1 series, Australian Instrument Services Pty. Ltd., Australia). The development of ovaries was further examined by dissecting females 4 ± 1 days post-emergence (before their first blood meal). Then we used an NIS Elements BR imaging microscope (Nikon Instruments, Japan) for photography and measurement.

We also measured *Wolbachia* density of fertile and infertile females after "isolation and identification", for 16 individuals per group. Screening based on real-time PCR assays followed methods described previously [25]. For each replicate, all samples were set up in a 384-well plate with white wells and measured by primers *aeg* and *wMwA* [26], two consistent replicates ($\Delta Ct < 1$) were obtained and their values were averaged for the density analysis.

Impact of larval stage extension on mosquito infertility

Based on results obtained from experiments described in previous sections, we suspected that mechanical changes causing female infertility occurred before metamorphosis. To further investigate whether the impact of *Wolbachia* infection on female infertility was only mediated through the egg stage or whether the prepupal stage also had an impact, we deprived 2nd instar *w*AlbB-infected larvae of food for two weeks before feeding them again until pupation then followed with an "isolation and identification". In non-starved controls, *w*AlbB-infected larvae were provided with food *ad libitum*. Each starvation treatment was performed with both stored eggs (stored at $26 \pm 1\square$ for 10 weeks) and non-stored eggs (stored at $26 \pm 1\square$ for 1 week) for a total of four treatments. For the group where infected females were neither stored nor starved, we had two replicates, while for the other three experimental groups we had three replicates, with each replicate containing 30 individual females. Before the blood meal, 15 individual females from each treatment 4 ± 1 days post-emergence were screened for *Wolbachia* infection and *Wolbachia* density, using screening methods for fertile and infertile females identical to those described in the previous section. After "isolation and identification", females were also dissected to confirm their fertility status by checking for the presence of ovaries. We also performed all four treatments with uninfected *Ae. aegypti* (with the same egg storage and larval starvation conditions) and

dissected 15 individuals 4 ± 1 days post-emergence from each replicate to check for existence

of ovaries.

Gene expression assays

We selected three genes related to mosquito reproduction and tested their expression levels in

females at different developmental stages: gene *vitellogenin receptor* (*vgr*), *ecdysone*

receptor (*ecr*) and *eggshell organizing factor* (*eof*) (Supplementary material 2). Gene

expression level was quantified relative to a control gene, *RPS17*, which encodes ribosomal

protein S17 [27]. We tested gene expression in female pupae 1 ± 0.5 days post-pupation and

in female adults 1 ± 0.5 days post-emergence in *w*AlbB-infected females that been stored as

eggs for 14 weeks as well as both *w*AlbB-infected and uninfected females that had not been

stored. For the *w*AlbB-infected females that had been stored as eggs, we also tested gene

expression levels in fertile and infertile females after "isolation and identification". Mosquito

samples were given a second blood meal and stored in RNA*later*® (Sigma Aldrich Cat No.

R0901-100ML) on the third day after, followed with the reverse transcription and real-time

PCA assays (Supplementary material 2)

Blood avidity of female mosquitoes

Normal fully engorged fertile females are reluctant to feed within a gonotrophic cycle. In this

experiment, we test the blood avidity of uninfected, *w*AlbB-infected fertile and infertile

females. After "isolation and identification", we measured blood avidity of female

mosquitoes by providing fertile and infertile females a third blood meal three days after they

had become engorged in the second blood feeding, when blood was almost digested. The

same volunteer who provided previous blood meals placed their forearm into a BugDorm

cage and left it there for 15 minutes. Fully engorged females were considered as having

successfully fed after this feeding attempt. Replicates were processed through different

batches of eggs and three replicates were completed with 20–30 individual females from each

replicate. We also randomly weighed 20 or to the maximum of females that had fully fed and

20 that had not been provided with a blood meal from all three replicates to estimate their

blood meal weight.

Statistics

We used R v. 3.6.0 with R studio v. 1.1.453 to conduct data analyses and visualizations [28].

After real-time PCR measurement, we used the $2^{-\Delta\Delta Ct}$ method to compare gene expression

levels [29], using either uninfected females from pupae or young adult stage as controls for

the same stage, or fertile females as the control group of infertile females after blood feeding.

The values of $2^{-\Delta\Delta Ct}$ were log-transformed for ANOVA analysis. For *Wolbachia* density we

used the $2^{\Delta Ct}$ method [30] to calculate relatively density before log transformation for

ANOVA analysis. We also use Tukey's honest significant difference (HSD) tests [31] for

further pairwise comparisons. For the proportion data from larval starvation and blood avidity

experiments we analysed data with binomial logistic regression models [32]. In the blood

avidity test, student's t-tests were used to test for changes of mosquito weight after blood

feeding.

Results

Consistency of infertility across gonotrophic cycles and ovarian dissection

We did an "isolation and identification" for *w*AlbB-infected females and provided a second

blood meal to fertile and infertile females. We confirmed that all females we scored as fertile

or infertile after the first gonotrophic cycle maintained their phenotype in the second

gonotrophic cycle. We then dissected and compared the ovarian appearance of fertile and

infertile females and discovered that ovaries cannot be observed in infertile females under a

microscope (Figure 1). Only in very rare cases (around one in 20) can immature ovarian

structure be seen, but with a narrow width similar to that of a Malpighian tube

(Supplementary material 3).

Figure 1. Appearance of internal organs in *w*AlbB-infected *Aedes aegypti* mosquitoes. (a)

Infertile females lack observable ovaries, while ovaries are present in (b) fertile females. The

background of the photographs was removed for clarity, with the original photographs

presented in the Supplementary material 3.

After "isolation and identification", we tested 16 fertile and infertile females and found that

fertile females had higher densities of *Wolbachia* (Figure 2: ANOVA: $F_{1,30}$ = 8.632, P =

0.006).

Figure 2. Box plots of relative *Wolbachia* density of fertile and infertile *w*AlbB-infected

females after isolation and identification (one week after blood meal).

Impact of larval stage extension on mosquito infertility

To test if there is any impact of larval stage extension on infertility in *w*AlbB-infected

females, we starved mosquito larvae to increase the duration of the larval stage from

approximately one week to three weeks for mosquitoes that had or had not been stored as

eggs. We found significant effects of both storage and starvation treatment, with an

interaction between these terms (Figure 3A: storage: $F_{1,7}$ = 256.304, P < 0.001; starvation:

$F_{1,7}$ = 32.412, P < 0.001; interaction: $F_{1,7}$ = 21.515, P = 0.002). Infertility rates associated

with *w*AlbB infection increased further with extension of the larval stage over and above that

seen with stored eggs by 15%, and the extension of the larval stage by starvation also induced

infertility of around 10% by itself. For uninfected *Ae. aegypti* treated in the same manner, all

females that were dissected contained ovaries, indicating the essential role of *Wolbachia*

*w*AlbB in inducing female infertility during larval starvation and egg storage.

Figure 3. (A) Boxplots showing the proportion of female infertility for *w*AlbB-infected

mosquitoes when stored or not stored as eggs and when larvae had been starved or not

starved. Values above boxplots represent corresponding averaged percentages. (B) Boxplots

of relative *Wolbachia* density in 4 ± 1 days post-emergence females.

We also screened for *Wolbachia* in all the treatments to confirm *Wolbachia* infection status.

We failed to detect an infection in five out of 60 individuals and these were excluded from

the density analysis (Supplementary material 4). *Wolbachia* density was significantly

influenced by larval starvation but not egg quiescence, with a significant interaction (Figure

3B: ANOVA: starvation: $F_{1,51} = 7.145$, P = 0.010; egg quiescence: $F_{1,51} = 2.475$, P = 0.122,

interaction: $F_{1,51} = 8.396$, P = 0.006). Egg quiescence decreased *Wolbachia* density, but

density increased with larval starvation, potentially cancelling out this effect [33].

Gene expression assays

We selected three genes related to reproductive development and tested their expression level

at different developmental stages. For comparisons before females took a blood meal (pupae

and early adult stage), we considered the uninfected mosquitoes as a control. For the pupal

stage, we found significant differences among the three groups of mosquitoes, but no

interaction with the genes (Figure 4A, ANOVA: mosquito group: $F_{2,81} = 26.717$, p < 0.001;

genes: $F_{2,81} = 1.044$, p = 0.357; interaction: $F_{4,81} = 0.423$, p = 0.792). Specifically, in Tukey's

HSD test, *w*AlbB-infected females that been stored as eggs had lower expression levels of all

of the three genes than the expression levels in uninfected females but only not gene *ecr* had

lower expression level in *w*AlbB-infected females that had not been stored.

For young adults 1± 0.5 days post-emergence, significant differences of gene expression

levels were found among mosquito groups, but there was no interaction with the genes

(Figure 4B, mosquito group: $F_{2,81} = 15.641$, $p < 0.001$; genes: $F_{2,81} = 0.383$, $p = 0.683$;

interaction: $F_{4,81} = 7.100$, $p = 0.792$). However, while differences among mosquito groups

were similar across the three genes, only differences for *vgr* expression were significant

between *w*AlbB-infected females that been stored and the other two mosquito groups, with

some individuals from stored eggs often having very low expression levels and these females

are probably infertile. The *w*AlbB infection may have increased expression of this gene,

indicating that there may also have some impacts of *Wolbachia* on the reproduction of non-

stored females (Figure 4B).

Figure 4. Boxplots of relative expression of reproduction-related genes *ecdysone receptor*

(*ecr*), *eggshell organizing factor* (*eof*) and *vitellogenin receptor* (*vgr*) in female (A) pupae

and (B) young adults (1 ± 0.5 days post-emergence). Values with the same letter are not

significantly different based on Tukey's HSD tests.

After "isolation and identification", female mosquitoes were provided a second blood meal

and were screened for gene expression levels three days later. Significant differences were

found between fertile and infertile females, and also between genes selected and their

interaction (Figure 5, mosquito group: $F_{1,54} = 235.696$, $p < 0.001$; genes: $F_{2,54} = 68.553$, $p <$

0.001; interaction: $F_{2,54} = 68.553$, $p < 0.001$). Specifically, all three genes showed the same

trend and there were significant differences between fertile and infertile females for

expression of *eof* and *vgr*, and especially for *vgr* whose expression was negligible in the

infertile mosquitoes.

Figure 5. Boxplots of relative expression of reproduction-related genes *ecdysone receptor*

(*ecr*), *eggshell organizing factor* (*eof*) and *vitellogenin receptor* (*vgr*) in fertile and infertile

females one week after their second blood meal. Values with the same letter are not

significantly different according to Tukey's HSD tests.

Blood avidity

After "isolation and identification", we provided female mosquitoes a second blood meal

following a successively blood feeding attempt on the third day after to test their blood

avidity. We found that infertile females had a higher proportion feeding compared to fertile

and uninfected females (Figure 6A, $F_{2,6} = 61.037$, $p < 0.001$). We also compared the weight

of fully fed and unfed females and noted significant differences between mosquito groups,

feeding status and their interaction (Figure 6B, ANOVA: mosquito group: $F_{1,73} = 29.06$, p <

0.001; feeding: $F_{1,73} = 322.89$, $p < 0.001$; interaction: $F_{2,73} = 15.26$, $p < 0.001$). Specifically,

significant weight increases were recorded for both *w*AlbB-infected fertile females ($t_{35} =$

9.627, $p < 0.001$) and *w*AlbB-infected infertile females ($t_{38} = 15.658$, $p < 0.001$), but the

weight of fully fed fertile females was significantly lower compared to infertile females ($t_{35} =$

-4.645, $p < 0.001$), while there was no significant difference between unfed fertile and

infertile females ($t_{38} = -1.096$, $p = 0.280$), suggesting that infertile females took in a larger

amount of blood during successive feeding.

Figure 6. Boxplots of (A) female proportion that blood fed three days after they had been

fully fed; (B) the weights of fed and unfed females.

Discussion

Wolbachia is a well-known bacterium that can alter the reproduction of its host to benefit its

spread. In our previous study, *Wolbachia w*AlbB was found to cause *Aedes aegypti* females

to become infertile following egg storage [20], inhibiting their ability to transmit vertically.

In this paper, we investigated this phenomenon and obtained the following results: 1) female

infertility is permanent; the development of ovaries has been interrupted by *Wolbachia*, no

mature ovaries can be observed and low expression levels of reproductive genes in infertile

females; 2) the effect accumulates across pre-pupal stages, in that the frequency of female

infertility depends on the duration before metamorphosis (including egg and larval stages);

and 3) infertile females maintain other female characteristics such as mating and blood

feeding, but they do not enter a gonotrophic cycle and blood feeding occurs more frequently.

We therefore confirmed that *Wolbachia w*AlbB altered the reproductive process in its new

host *Ae. aegypti*.

The effect of *Wolbachia* on female infertility is different to other fitness impacts of

Wolbachia and the four typical reproductive alterations (parthenogenesis, feminization, male

killing and cytoplasmic incompatibility) in which *Wolbachia* alters host reproduction in ways

that increase its vertical transmission success [4, 34]. Feminization found in lepidopteran

insects that have female-heterogametic chromosomes [35, 36] can also act on the

development of reproductive organs [37]. Unlike Lepidoptera, *Ae. aegypti* mosquitoes have a

heterogametic sex determining region [38], which results in maleness being expressed in the

heterozygous form [39]. However, feminization converts genetic males into functional

females, while the infertility effect converts functional females into non-functional females

without male characteristics, suggesting different *Wolbachia*-related mechanisms [40, 41].

The development of mosquito ovaries can be divided into multiple stages [42], including

previtellogenic stages when follicles remain dormant before a blood meal, and vitellogenic

stages, when ovarian development can be observed within 30 minutes after females engorge

[43]. The feeding process stimulates the expression of a steroid hormone, 20-

hydroxyecdysone (20E), which plays a critical role in reproductive regulation by interacting

with its intracellular receptor, and consists of two protein subunits: ecdysteroid receptor (EcR)

and ultraspirale [44, 45]. EcR can bind to specific DNA sequences and regulates the

expression of related genes such as vitellogenin (*Vg*) in the mosquito fat body, which encodes

a yolk protein precursor (YPP) [46-49]. The YPP is then secreted to the haemolymph and

internalized through related receptor VgR in developing oocytes [50, 51]. At a late stage of

egg production, the gene *eof* is expressed, and encodes an essential protein for eggshell

formation [52]. In this study, we tested the expression level of *vgr, ecr* and *eof* at the female

pupal, young adult before blood feed and old adult after blood feed stage to compare

expression between stored and non-stored/fertile and infertile females. Our data show that

differences in gene expression were substantial at each stage. For pupae, large variations of

expression levels were found among individuals, especially for gene *vgr*, probably indicates

its expression focus on pupae of a certain age, and *w*AlbB-infected *Ae. aegypti* females that

had been stored long-term as eggs had lower expression levels of the three reproductive

genes compared to uninfected females, and the expression level of gene *vgr* was significantly

lower than both uninfected and *w*AlbB-infected normal females at young adult stage,

suggesting the formation and development of oocytes was impacted, which is also supported

by the very low relative expression of *vgr* in infertile females when compared with fertile

females at the third day after their blood feed, and the inability to observe mature ovaries in

infertile females. What is more, we also found substantial lower expression level of genes *ecr*

and *eof* in *w*AlbB-infected female pupae hatched from long-stored eggs, while this difference

was not found in young adults. As *ecr* is also expressed in other organs as it encodes the

receptor of a hormone [53], the down-regulated of *ecr* may indicate that egg quiescence have

substantial impact on the pupation process of *w*AlbB-infected females. However, interactions

between *Wolbachia* and specific reproductive signals and mechanisms are poorly understood

and require further investigation such as transcriptome and pathway analysis [54, 55].

The novel reproductive alteration induced by *Wolbachia w*AlbB could slow the spread of the

infection in mosquito populations. This may occur not only when the egg desiccation period

is extended during dry and warm seasons [56, 57], but also when the larval period is extended

due to poor provision of food [58, 59], for females are less likely to produce offspring. These

effects can reduce the efficiency of *Wolbachia* invasion which aims to inhibit the

transmission of arboviruses [8, 24, 60]. While *w*AlbB has established successfully in

Malaysia which is warm and humid year-round precluding extended egg desiccation periods

and promoting fast development of larvae [10, 61], our discovery provides guidance for

future releases in other climates and highlights the importance of future monitoring in release

areas [20].

Our observation that infertile females show increased blood avidity highlights a potential risk

of increased nuisance biting following a *Wolbachia* release program, which may lead to

community discontentment. Normal *Ae. aegypti* females ingest human blood to obtain unique

nutrients for egg development [62, 63]. However, while infertile females lack ovaries, they

still behave like normal females in terms of blood feeding. Previous research found that

multiple feeding within a gonotrophic cycle is caused by nutritional reserve depletion or

feeding interruption [64]. We tested blood feeding behaviour after the first gonotrophic cycle

of "isolation and identification", which can also reduce the influence of nutrition reserves as

multiple blood feeding may rescue the effect of nutritional deprivation [65, 66]. We selected

a three-day interval which is the period mosquitoes need for blood digestion to test the

frequency of refeeding, and found that a large proportion of infertile females will refeed,

indicating they do not follow a gonotrophic cycle. This behavioural change of *Wolbachia*-

infected infertile females has significant implications for vector control population release

strategies [8, 67]; in particular it would be interesting to test whether the ability of these

infertile females to carry and transmit arbovirus has changed as the density of their

Wolbachia infection may be lower than that of fertile females [68, 69]. Similar concerns have

been raised about increased biting frequencies of irradiated female mosquitoes in sterile

insect technique programs [70], however the prevalence of infertile females is likely to be

much higher in *Wolbachia* mosquito populations compared to irradiated females that are only

released accidentally.

The reproductive alterations of *Wolbachia* discovered in this study have significant

evolutionary implications. So far our work relates to *Ae. aegypti*, where the *Wolbachia*

*w*AlbB was artificially introduced recently from a close relative of *Ae. aegypti*, *Aedes*

albopictus, in order to reduce the transmission of arboviral diseases [71]. A *Wolbachia* strain

may induce similar fitness costs in a new host when compared to its native host, such in the

case of life-shortening of *w*MelPop which is expressed in both *Drosophila melanogaster* and

Ae. aegypti [11], and remains stably expressed following long-term laboratory culture [72].

We do not yet know if there are similar reproductive effects of *Wolbachia* *w*AlbB expressed

in its native host, *Ae. albopictus*, while previous study indicating that low temperature

adaptation [33, 73], a different effect of *w*AlbB, might be associated. On the other hand, we

have also discovered a smaller loss of female infertility in *w*Mel infected *Ae. aegypti* [20],

though the native host of *w*Mel, *D. melanogaster,* does not enter egg quiescence, reduced

fecundity of *w*Mel-infected females was discovered under dormancy conditions [74].

Although other *Wolbachia* strains remain to be investigated, it is also possible that *Ae.*

aegypti has mechanisms underlying ovarian formation that can be interrupted by

endosymbionts. However, it is unclear why the frequency of infertile females increases when

the density of *Wolbachia* decreases as the result of increased periods of egg storage or larval

development time, and why infertile females have lower *Wolbachia* densities than fertile

females in the same treatment. There may be cumulative effects of *Wolbachia* not evident

from density measures at early stages of development contributing to failure of ovarian

formation. One possible hypothesis is that *Wolbachia* competes for some essential but sparse

nutrients with its host *Ae. aegypti* at an early life stage, which requires further investigation.

In conclusion, we further investigate a novel reproductive alteration of *Wolbachia* that we

discovered previously [20]. We confirmed that *Ae. aegypti* females infected with *Wolbachia*

*w*AlbB can become infertile when they are unable to form functional ovaries during

metamorphosis, but these females retain other female characteristics, leading to an increased

biting frequency. Our study provides significant guidance for future *Wolbachia* releases and

has important evolutionary implications for understanding the reproductive alteration of

Wolbachia, especially in novel hosts.

Funding

This research was funded by the National Health and Medical Research Council (1132412,

1118640, www.nhmrc.gov.au). The funders had no role in the study design, data collection

and analysis, decision to publish, or preparation of the manuscript.

Acknowledgment

We thank Véronique Paris for her assistance in measuring mosquito weights.

1.   Zug R., Hammerstein P. 2012 Still a host of hosts for *Wolbachia*: analysis of recent
data suggests that 40% of terrestrial arthropod species are infected. *PLoS One* **7**(6), e38544.
(doi:10.1371/journal.pone.0038544).
2.   Stouthamer R., Breeuwer J.A., Hurst G.D. 1999 *Wolbachia pipientis*: microbial
manipulator of arthropod reproduction. *Annu Rev Microbiol* **53**, 71-102.
(doi:10.1146/annurev.micro.53.1.71).
3.   Werren J.H., Baldo L., Clark M.E. 2008 *Wolbachia*: master manipulators of
invertebrate biology. *Nat Rev Microbiol* **6**(10), 741.
4.   O'Neill S.L., Hoffmann A., Werren J. 1997 *Influential passengers: inherited*
microorganisms and arthropod reproduction, Oxford University Press.
5.   Zug R., Hammerstein P. 2015 Bad guys turned nice? A critical assessment of
Wolbachia mutualisms in arthropod hosts. *Biol Rev Camb Philos Soc* **90**(1), 89-111.
(doi:10.1111/brv.12098).
6.   Hoffmann A.A., Turelli M., Harshman L.G. 1990 Factors affecting the distribution of
cytoplasmic incompatibility in *Drosophila simulans*. *Genetics* **126**(4), 933-948.
7.   Perrot‐Minnot M.J., Cheval B., Migeon A., Navajas M. 2002 Contrasting effects of
Wolbachia on cytoplasmic incompatibility and fecundity in the haplodiploid mite
Tetranychus urticae. *J Evol Biol* **15**(5), 808-817.
8.   Ross P.A., Turelli M., Hoffmann A.A. 2019 Evolutionary ecology of *Wolbachia*
releases for disease control. *Annu Rev Genet* **53**, 93-116. (doi:10.1146/annurev-genet-
112618-043609).
9.   Ryan P.A., Turley A.P., Wilson G., Hurst T.P., Retzki K., Brown-Kenyon J., Hodgson L.,
Kenny N., Cook H., Montgomery B.L. 2019 Establishment of *w*Mel *Wolbachia* in *Aedes*
aegypti mosquitoes and reduction of local dengue transmission in Cairns and surrounding
locations in northern Queensland, Australia. *Gates Open Res* **3**, 1547.
10.   Nazni W.A., Hoffmann A.A., NoorAfizah A., Cheong Y.L., Mancini M.V., Golding N.,
Kamarul G.M., Arif M.A., Thohir H., NurSyamimi H. 2019 Establishment of *Wolbachia* strain
*w*AlbB in Malaysian populations of *Aedes aegypti* for dengue control. *Curr Biol* **29**(24), 4241-
4248.
11.   McMeniman C.J., Lane R.V., Cass B.N., Fong A.W.C., Sidhu M., Wang Y.-F., O'Neill S.L.
2009 Stable introduction of a life-shortening *Wolbachia* infection into the mosquito *Aedes*
aegypti. *Science* **323**(5910), 141-144. (doi:10.1126/science.1165326).
12.   Turley A.P., Moreira L.A., O'Neill S.L., McGraw E.A. 2009 *Wolbachia* infection reduces
blood-feeding success in the dengue fever mosquito, *Aedes aegypti*. *PLoS Negl Trop Dis* **3**(9),
e516.
13.   Yeap H.L., Mee P., Walker T., Weeks A.R., O'Neill S.L., Johnson P., Ritchie S.A.,
Richardson K.M., Doig C., Endersby N.M., et al. 2011 Dynamics of the "Popcorn" *Wolbachia*

infection in outbred *Aedes aegypti* informs prospects for mosquito vector control. *Genetics*
187(2), 583-595. (doi:10.1534/genetics.110.122390).
14. Ware-Gilmore F., Sgrò C.M., Xi Z., Dutra H.L.C., Jones M.J., Shea K., Hall M.D.,
Thomas M.B., McGraw E.A. 2021 Microbes increase thermal sensitivity in the mosquito
Aedes aegypti, with the potential to change disease distributions. *PLoS Negl Trop Dis* **15**(7),
e0009548. (doi:10.1371/journal.pntd.0009548).
15. Fenn K., Blaxter M. 2006 *Wolbachia* genomes: revealing the biology of parasitism
and mutualism. *Trends Parasitol* **22**(2), 60-65. (doi:10.1016/j.pt.2005.12.012).
16. Weeks A.R., Turelli M., Harcombe W.R., Reynolds K.T., Hoffmann A.A. 2007 From
parasite to mutualist: rapid evolution of *Wolbachia* in natural populations of *Drosophila*.
PLoS Biol **5**(5), e114.
17. Lau M.J., Schmidt T.L., Yang Q., Chung J., Sankey L., Ross P.A., Hoffmann A.A. 2021
Genetic stability of *Aedes aegypti* populations following invasion by wMel *Wolbachia*. *BMC*
Genomics **22**(1), 894. (doi:10.1186/s12864-021-08200-1).
18. Carrington L.B., Hoffmann A.A., Weeks A.R. 2010 Monitoring long-term evolutionary
changes following *Wolbachia* introduction into a novel host: the *Wolbachia* popcorn
infection in *Drosophila simulans*. *Proc Biol Sci* **277**(1690), 2059-2068.
19. Ford S.A., Allen S.L., Ohm J.R., Sigle L.T., Sebastian A., Albert I., Chenoweth S.F.,
McGraw E.A. 2019 Selection on *Aedes aegypti* alters *Wolbachia*-mediated dengue virus
blocking and fitness. *Nat Microbiol* **4**(11), 1832-1839. (doi:10.1038/s41564-019-0533-3).
20. Lau M.J., Ross P.A., Hoffmann A.A. 2021 Infertility and fecundity loss of *Wolbachia*-
infected *Aedes aegypti* hatched from quiescent eggs is expected to alter invasion dynamics.
PLoS Negl Trop Dis **15**(2), e0009179. (doi:10.1371/journal.pntd.0009179).
21. Allman M.J., Fraser J.E., Ritchie S.A., Joubert D.A., Simmons C.P., Flores H.A. 2020
Wolbachia's deleterious impact on *Aedes aegypti* egg development: The potential role of
nutritional parasitism. *Insects* **11**(11), 735. (doi:10.3390/insects11110735).
22. Xi Z.Y., Khoo C.C.H., Dobson S.L. 2005 *Wolbachia* establishment and invasion in an
Aedes aegypti laboratory population. *Science* **310**(5746), 326-328.
(doi:10.1126/science.1117607).
23. Garcia G.d.A., Sylvestre G., Aguiar R., da Costa G.B., Martins A.J., Lima J.B.P.,
Petersen M.T., Lourenço-de-Oliveira R., Shadbolt M.F., Rašić G., et al. 2019 Matching the
genetics of released and local *Aedes aegypti* populations is critical to assure *Wolbachia*
invasion. *PLoS Negl Trop Dis* **13**(1), e0007023. (doi:10.1371/journal.pntd.0007023).
24. Hoffmann A.A., Montgomery B., Popovici J., Iturbe-Ormaetxe I., Johnson P., Muzzi F.,
Greenfield M., Durkan M., Leong Y., Dong Y. 2011 Successful establishment of *Wolbachia* in
Aedes populations to suppress dengue transmission. *Nature* **476**(7361), 454-457.
25. Lee S.F., White V.L., Weeks A.R., Hoffmann A.A., Endersby N.M. 2012 High-
throughput PCR assays to monitor *Wolbachia* infection in the dengue mosquito (*Aedes*
aegypti) and *Drosophila simulans*. *Appl Environ Microbiol* **78**(13), 4740-4743.
26. Lau M.-J., Hoffmann A.A., Endersby-Harshman N.M. 2021 A diagnostic primer pair to
distinguish between wMel and wAlbB *Wolbachia* infections. *PLoS One* **16**(9), e0257781.
(doi:10.1371/journal.pone.0257781).
27. Dzaki N., Ramli K.N., Azlan A., Ishak I.H., Azzam G. 2017 Evaluation of reference
genes at different developmental stages for quantitative real-time PCR in *Aedes aegypti*. *Sci*
Rep **7**, 43618. (doi:10.1038/srep43618).
28. Team R.C. 2019 R: A language and environment for statistical computing. R
Foundation for Statistical Computing, Vienna, Austria. (

29.   Livak K.J., Schmittgen T.D. 2001 Analysis of relative gene expression data using real-
time quantitative PCR and the 2(-delta delta C(T)) method. *Methods* **25**(4), 402-408.
(doi:10.1006/meth.2001.1262).
30.   Unckless R.L., Boelio L.M., Herren J.K., Jaenike J. 2009 *Wolbachia* as populations
within individual insects: causes and consequences of density variation in natural
populations. *Proc Biol Sci* **276**(1668), 2805-2811. (doi:10.1098/rspb.2009.0287).
31.   Tukey J.W. 1949 Comparing individual means in the analysis of variance. *Biometrics*
5(2), 99-114.
32.   Hastie T.J. 2017 Generalized additive models. In *Statistical models in S* (ed. Science
S.), pp. 249-307, Routledge.
33.   Lau M.J., Ross P.A., Endersby-Harshman N.M., Hoffmann A.A. 2020 Impacts of low
temperatures on *Wolbachia* (Rickettsiales: Rickettsiaceae)-infected *Aedes aegypti* (Diptera:
Culicidae). *J Med Entomol* **57**(5), 1567-1574. (doi:10.1093/jme/tjaa074).
34.   Werren J.H. 1997 Biology of *Wolbachia*. *Annu Rev Entomol* **42**, 587-609.
35.   Kageyama D., Nishimura G., Hoshizaki S., Ishikawa Y. 2002 Feminizing *Wolbachia* in
an insect, *Ostrinia furnacalis* (Lepidoptera: Crambidae). *Heredity (Edinb)* **88**(6), 444-449.
(doi:10.1038/sj.hdy.6800077).
36.   Hiroki M., Kato Y., Kamito T., Miura K. 2002 Feminization of genetic males by a
symbiotic bacterium in a butterfly, *Eurema hecabe* (Lepidoptera: Pieridae).
Naturwissenschaften **89**(4), 167-170. (doi:10.1007/s00114-002-0303-5).
37.   Narita S., Kageyama D., Nomura M., Fukatsu T. 2007 Unexpected mechanism of
symbiont-induced reversal of insect sex: feminizing *Wolbachia* continuously acts on the
butterfly *Eurema hecabe* during larval development. *Appl Environ Microbiol* **73**(13), 4332-
4341. (doi:10.1128/aem.00145-07).
38.   Hall A.B., Timoshevskiy V.A., Sharakhova M.V., Jiang X., Basu S., Anderson M.A., Hu
W., Sharakhov I.V., Adelman Z.N., Tu Z. 2014 Insights into the preservation of the
homomorphic sex-determining chromosome of *Aedes aegypti* from the discovery of a male-
biased gene tightly linked to the M-locus. *Genome biology evolution* **6**(1), 179-191.
39.   Lutrat C., Olmo R.P., Baldet T., Bouyer J., Marois E. 2022 Transgenic expression of Nix
converts genetic females into males and allows automated sex sorting in *Aedes albopictus*.
Commun Biol **5**(1), 210. (doi:10.1038/s42003-022-03165-7).
40.   Rigaud T., Juchault P., Mocquard J.P. 1997 The evolution of sex determination in
isopod crustaceans. *Bioessays* **19**(5), 409-416.
41.   Bandi C., Dunn A.M., Hurst G.D., Rigaud T. 2001 Inherited microorganisms, sex-
specific virulence and reproductive parasitism. *Trends Parasitol* **17**(2), 88-94.
(doi:10.1016/s1471-4922(00)01812-2).
42.   Raikhel A.S., Kokoza V.A., Zhu J., Martin D., Wang S.F., Li C., Sun G., Ahmed A.,
Dittmer N., Attardo G. 2002 Molecular biology of mosquito vitellogenesis: from basic studies
to genetic engineering of antipathogen immunity. *Insect Biochem Mol Biol* **32**(10), 1275-
1286. (doi:10.1016/s0965-1748(02)00090-5).
43.   Clements A., Boocock M. 1984 Ovarian development in mosquitoes: stages of
growth and arrest, and follicular resorption. *Physiol Entomol* **9**(1), 1-8.
44.   Yao T.P., Segraves W.A., Oro A.E., McKeown M., Evans R.M. 1992 *Drosophila*
ultraspiracle modulates ecdysone receptor function via heterodimer formation. *Cell* **71**(1),
63-72. (doi:10.1016/0092-8674(92)90266-f).

45.   Hodin J., Riddiford L.M. 1998 The ecdysone receptor and ultraspiracle regulate the
timing and progression of ovarian morphogenesis during *Drosophila metamorphosis*. *Dev*
Genes Evol **208**(6), 304-317. (doi:10.1007/s004270050186).
46.   Martín D., Wang S.F., Raikhel A.S. 2001 The vitellogenin gene of the mosquito *Aedes*
aegypti is a direct target of ecdysteroid receptor. *Mol Cell Endocrinol* **173**(1-2), 75-86.
(doi:10.1016/s0303-7207(00)00413-5).
47.   Wang S.F., Miura K., Miksicek R.J., Segraves W.A., Raikhel A.S. 1998 DNA binding and
transactivation characteristics of the mosquito ecdysone receptor-Ultraspiracle complex. *J*
Biol Chem **273**(42), 27531-27540. (doi:10.1074/jbc.273.42.27531).
48.   Roy S.G., Raikhel A.S. 2011 The small GTPase Rheb is a key component linking amino
acid signaling and TOR in the nutritional pathway that controls mosquito egg development.
Insect Biochem Mol Biol **41**(1), 62-69. (doi:10.1016/j.ibmb.2010.10.001).
49.   Raikhel A.S., Lea A.O. 1983 Previtellogenic development and vitellogenin synthesis in
the fat body of a mosquito: an ultrastructural and immunocytochemical study. *Tissue Cell*
15(2), 281-299. (doi:10.1016/0040-8166(83)90023-x).
50.   Sappington T.W., Raikhel A.S. 1998 Molecular characteristics of insect vitellogenins
and vitellogenin receptors. *Insect Biochem Mol Biol* **28**(5-6), 277-300. (doi:10.1016/s0965-
1748(97)00110-0).
51.   Tufail M., Takeda M. 2009 Insect vitellogenin/lipophorin receptors: molecular
structures, role in oogenesis, and regulatory mechanisms. *J Insect Physiol* **55**(2), 87-103.
(doi:10.1016/j.jinsphys.2008.11.007).
52.   Isoe J., Koch L.E., Isoe Y.E., Rascón A.A., Jr., Brown H.E., Massani B.B., Miesfeld R.L.
2019 Identification and characterization of a mosquito-specific eggshell organizing factor in
Aedes aegypti mosquitoes. *PLoS Biol* **17**(1), e3000068. (doi:10.1371/journal.pbio.3000068).
53.   Koelle M.R., Talbot W.S., Segraves W.A., Bender M.T., Cherbas P., Hogness D.S. 1991
The *Drosophila EcR* gene encodes an ecdysone receptor, a new member of the steroid
receptor superfamily. *Cell* **67**(1), 59-77. (doi:10.1016/0092-8674(91)90572-g).
54.   Belles X., Piulachs M.D. 2015 Ecdysone signalling and ovarian development in insects:
from stem cells to ovarian follicle formation. *Biochim Biophys Acta* **1849**(2), 181-186.
(doi:10.1016/j.bbagrm.2014.05.025).
55.   Dansereau D.A., Lasko P. 2008 The development of germline stem cells in *Drosophila*.
Methods Mol Biol **450**, 3-26. (doi:10.1007/978-1-60327-214-8_1).
56.   Juliano S.A., O'Meara G.F., Morrill J.R., Cutwa M.M. 2002 Desiccation and thermal
tolerance of eggs and the coexistence of competing mosquitoes. *Oecologia* **130**(3), 458-469.
57.   Faull K.J., Williams C.R. 2015 Intraspecific variation in desiccation survival time of
Aedes aegypti (L.) mosquito eggs of Australian origin. *J Vector Ecol* **40**(2), 292-300.
(doi:10.1111/jvec.12167).
58.   Ross P.A., Endersby N.M., Hoffmann A.A. 2016 Costs of three *Wolbachia* infections
on the survival of *Aedes aegypti* larvae under starvation conditions. *PLoS Negl Trop Dis* **10**(1),
e0004320.
59.   Barrera R., Medialdea V. 1996 Development time and resistance to starvation of
mosquito larvae. *J Nat Hist* **30**(3), 447-458.
60.   Koh C., Audsley M.D., Di Giallonardo F., Kerton E.J., Young P.R., Holmes E.C., McGraw
E.A. 2019 Sustained *Wolbachia*-mediated blocking of dengue virus isolates following serial
passage in *Aedes aegypti* cell culture. *Virus Evol* **5**(1), vez012.
61.   Ahmad N.A., Mancini M.V., Ant T.H., Martinez J., Kamarul G.M.R., Nazni W.A.,
Hoffmann A.A., Sinkins S.P. 2021 *Wolbachia* strain *w*AlbB maintains high density and dengue

inhibition following introduction into a field population of *Aedes aegypti. Philos Trans R Soc*
Lond B Biol Sci **376**(1818), 20190809. (doi:10.1098/rstb.2019.0809).
62. Gulia-Nuss M., Elliot A., Brown M.R., Strand M.R. 2015 Multiple factors contribute to
anautogenous reproduction by the mosquito *Aedes aegypti. J Insect Physiol* **82**, 8-16.
(doi:10.1016/j.jinsphys.2015.08.001).
63. Scott T.W., Naksathit A., Day J.F., Kittayapong P., Edman J.D. 1997 A fitness
advantage for *Aedes aegypti* and the viruses it transmits when females feed only on human
blood. *Am J Trop Med Hyg* **57**(2), 235-239. (doi:10.4269/ajtmh.1997.57.235).
64. Farjana T., Tuno N. 2013 Multiple blood feeding and host-seeking behavior in *Aedes*
aegypti and *Aedes albopictus* (Diptera: Culicidae). *J Med Entomol* **50**(4), 838-846.
(doi:10.1603/me12146).
65. Koella J.C., Sørensen F.L., Anderson R.A. 1998 The malaria parasite, *Plasmodium*
falciparum, increases the frequency of multiple feeding of its mosquito vector, *Anopheles*
gambiae. Proc Biol Sci **265**(1398), 763-768. (doi:10.1098/rspb.1998.0358).
66. Reyes-Villanueva F. 2004 Egg development may require multiple bloodmeals among
small *Aedes aegypti* (Diptera: Culicidae) field collected in northeastern Mexico. *Fla Entomol*
87(4), 630-632.
67. Almeida L., Duprez M.G., Privat Y., Vauchelet N. 2019 Mosquito population control
strategies for fighting against arboviruses. *Math Biosci Eng* **16**(6), 6274-6297.
(doi:10.3934/mbe.2019313).
68. Johnson K.N. 2015 The impact of *Wolbachia* on virus infection in mosquitoes. *Viruses*
7(11), 5705-5717. (doi:10.3390/v7112903).
69. Terradas G., McGraw E.A. 2017 *Wolbachia*-mediated virus blocking in the mosquito
vector *Aedes aegypti. Curr Opin Insect Sci* **22**, 37-44. (doi:10.1016/j.cois.2017.05.005).
70. Moretti R., Lampazzi E., Damiani C., Fabbri G., Lombardi G., Pioli C., Desiderio A.,
Serrao A., Calvitti M. 2022 Increased biting rate and decreased *Wolbachia* density in
irradiated Aedes mosquitoes. *Parasit Vectors* **15**(1), 67. (doi:10.1186/s13071-022-05188-9).
71. Xi Z., Dean J.L., Khoo C., Dobson S.L. 2005 Generation of a novel *Wolbachia* infection
in *Aedes albopictus* (Asian tiger mosquito) via embryonic microinjection. *Insect Biochem Mol*
Biol **35**(8), 903-910.
72. Ross P.A., Axford J.K., Callahan A.G., Richardson K.M., Hoffmann A.A. 2020 Persistent
deleterious effects of a deleterious *Wolbachia* infection. *PLoS Negl Trop Dis* **14**(4),
e0008204-e0008204. (doi:10.1371/journal.pntd.0008204).
73. Reinhold J.M., Lazzari C.R., Lahondère C. 2018 Effects of the environmental
temperature on *Aedes aegypti* and *Aedes albopictus* mosquitoes: a review. *Insects* **9**(4), 158.
74. Kriesner P., Conner W.R., Weeks A.R., Turelli M., Hoffmann A.A. 2016 Persistence of
a *Wolbachia* infection frequency cline in *Drosophila melanogaster* and the possible role of
reproductive dormancy. *Evolution* **70**(5), 979-997. (doi:10.1111/evo.12923).
631