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Modeling the potential of wAu-Wolbachia strain invasion in mosquitoes to control Aedes-borne arboviral infections

Samson T. Ogunlade^{1,2™}, Adeshina I. Adekunle¹, Michael T. Meehan¹, Diana P. Rojas³ & Emma S. McBryde¹

Arboviral infections such as dengue, Zika and chikungunya are fast spreading diseases that pose significant health problems globally. In order to control these infections, an intracellular bacterium called *Wolbachia* has been introduced into wild-type mosquito populations in the hopes of replacing the vector transmitting agent, *Aedes aegypti* with one that is incapable of transmission. In this study, we developed a *Wolbachia* transmission model for the novel wAu strain which possesses several favourable traits (e.g., enhanced viral blockage and maintenance at higher temperature) but not cyctoplasmic incompatibility (CI)—when a *Wolbachia*-infected male mosquito mates with an uninfected female mosquito, producing no viable offspring. This model describes the competitive dynamics between wAu-*Wolbachia*-infected and uninfected mosquitoes and the role of imperfect maternal transmission. By analysing the system via computing the basic reproduction number(s) and stability properties, the potential of the wAu strain as a viable strategy to control arboviral infections is established. The results of this work show that enhanced maintenance of *Wolbachia* infected mosquito invasion. This study will support future arboviral control programs, that rely on the introduction of new *Wolbachia* variants.

Arthropod-borne viruses, or arboviruses, are viruses that are transmitted via blood feeding arthropods¹. Arboviral infections such as dengue, Zika and chikungunya are fast spreading diseases that pose significant health problems globally^{2–5}. These viral infections, in particular dengue, are transmitted mainly by *Aedes aegypti* and sometimes by *Aedes albopictus* (Asian Tiger) female mosquitoes when taking a blood meal from the host^{6,7}. Approximately 390 million dengue infections are estimated to occur worldwide annually, putting 40% of the total human population at risk⁸. Dengue infection is the most geographically wide-spread of the arboviral infections^{3,8}. It has different severity levels which are classified according to disease progression from dengue without warning signs to dengue with warning signs and then severe dengue⁹. Clinical manifestation includes sudden high-grade fever, headache, nausea, arthralgia, eye pain, muscle ache and rash in some cases¹⁰. Presently, there is no specific universal treatment for dengue infections: the vaccine envelopment targets young populations; the efficacy of the only vaccine licensed depends on prior immunity to at least one serotype of dengue; and it provides heterogeneous protection against the different serotypes^{11,12}.

Other arboviral infections such as Zika, chikungunya and yellow fever are also of global health concern¹³. These arboviral infections have occurred simultaneously with dengue^{13,14}. Some of these infections share many similar clinical manifestations with dengue infection and also allow arboviral coinfection such as dengue and chikungunya¹⁵, chikungunya and Zika¹⁶ and yellow fever and chikungunya¹⁷. Although, there are no specific treatments for Zika and chikungunya viral infections, these infections can be managed by supportive treatment of symptomatic individuals and adequate rest. This treatment includes fluid intake and administering drugs such

¹Australian Institute of Tropical Health and Medicine, James Cook University, Townsville, QLD, Australia. ²College of Medicine and Dentistry, James Cook University, Townsville, QLD, Australia. ³College of Public Health, Medical and Veterinary Sciences, Division of Tropical Health and Medicine, James Cook University, Townsville, QLD, Australia. [∞]email: samson.oqunlade@my.jcu.edu.au

Features	wAu	wMel	wMelPop	wAlbA	wAlbB
Viral blockage	High ²³	Medium ^{40,41}	High ^{29,41–43}	Medium ²³	High ⁴⁴
Maternal transmission	High ²³	High ³⁰	High ^{32,45}	High ²³	High ⁴⁶
Loss of Wolbachia infection at higher hemperature	Low ²³	High ²³	High ²³	Medium ²³	Medium ²³
Fitness cost	Medium ²³	Medium ²³	High ^{47,48}	High ²³	Medium ²²
Cytoplasmic incompatibility	None ²³	High ³⁰	High ^{32,45}	High ²³	High ⁴⁶

Table 1. Characteristics of different *Wolbachia* strains in *Aedes* mosquitoes: as defined in²², the percentages (%) of the effects of these features are: High \longrightarrow above 90, Medium \longrightarrow 20 to 90, Low \longrightarrow less than 20 and None \longrightarrow 0, (features not detected).

as acetaminophen to suppress pain and fever 18,19 . However the prevention strategy for yellow fever infection is available i.e. vaccination 20,21 .

To control these infections, an intracellular bacterium called *Wolbachia* can be used to suppress transmission in arthropods such as mosquitoes and flies^{22–25}. *Wolbachia* infection inhibits arboviral transmission in mosquitoes via four mechanisms: immune priming—preactivation of the mosquito immune system; induction of the phenoloxidase cascade—triggers immune response to viruses; competition of intracellular resources—inducing authophagy; and induction of microRNA-dependent immune pathways—essential for gene regulation and stability, immune defense, ageing and organ differentiation²⁶. This endosymbiotic bacterium which exists naturally in more than 50% of all insect species can be found within the cytoplasm of the cells of their hosts^{25,27,28}. Whilst *Wolbachia* is not naturally present in *Aedes aegypti*, it can be introduced via stable transinfections using microinjections^{29,30}.

The *Wolbachia*-based control strategy is carried out by infecting mosquitoes with a strain of *Wolbachia* and then releasing them into wild mosquito populations in the hopes of replacing the vector transmitting agent *Aedes aegypti* with one that is incapable of transmission^{29–31}. Infecting an *Aedes* mosquito with *Wolbachia* can change some of the *Aedes* characteristic features. In practice, *Wolbachia* can reduce the life-span of mosquitoes by half producing a deleterous fitness effect³². Another feature is cytoplasmic incompatibility (CI)^{22,33–35} which occurs when a *Wolbachia* infected male mates with an incompatible female mosquito (usually *Wolbachia* uninfected) producing no offspring³⁶. Other features of *Wolbachia* which serve as liabilities in mosquitoes include: imperfect maternal transmission (IMT)^{30,37} and loss of *Wolbachia* infection (LWI). LWI impedes the establishment of *Wolbachia*-infected mosquitoes and is a result of mosquito vulnerability to high temperature^{38,39}.

However, a novel strain of *Wolbachia*: *w*Au, has shown to produce high viral blockage whilst maintaining *Wolbachia* infection in *Aedes* mosquitoes at higher temperature²³. Moreover, *w*Au allows superinfection to occur when *w*Au and other strains of *Wolbachia* co-exist in the vector host²³. Despite these favourable features, *w*Au does not induce CI²³. Although CI absence does not establish *Wolbachia* infected mosquitoes, the effect could be outweighed by LWI and IMT³⁷.

The difference in the common Wolbachia strain features are described in Table 1 below.

In general, the introduction of mathematical models to understand infection dynamics of diseases has long been helpful in the area of disease control⁴⁹. A number mathematical models of *Wolbachia* dynamics in a mosquito population have been formulated^{37,50–58}. Some of these models introduced *Wolbachia* strain(s) into a mosquito population and classified them into age-sturctured Wolbachia-infected and -uninfected mosquito compartments 37,53,54,57. Ndii et al.53, formulated a mathematical model for the Wolbachia interaction between the immature stages (aquatic stage), adult male and female mosquito populations to investigate the persistence of mosquitoes infected with Wolbachia when competing with the uninfected ones. They derived the steady state solutions and showed that parameters such as maternal transmission, reproductive, death and maturation rates drive the persistence of the Wolbachia-infected mosquito population. A similar model developed by Xue et al. considered the Wolbachia-induced fitness change and the CI effect⁵⁷. They showed that if the basic reproduction number (R_0) of the Wolbachia-infected mosquitoes is less than one, an endemic Wolbachia infection can still occur via backward bifurcation if a sufficient number of the mosquitoes are introduced into the population. A mathematical model of Wolbachia to control dengue fever transmission⁵² was developed by Hughes et al. The model showed that the use of Wolbachia has high potential to control dengue where the R_0 due to Wolbachiainfected Aedes mosquitoes is not too large in endemic areas. Another study of a Wolbachia invasive model incorporated IMT and LWI and showed that CI does not guarantee the establishment of Wolbachia-infected mosquitoes as the disadvantages derived from IMT and LWI in the production of Wolbachia-infected mosquitoes could outweigh CI³⁷.

Additionally, a study conducted by O'Reilly et al combining multiple modeling methods, was used to estimate the burden of dengue and map its distribution across Indonesia⁵⁹. They predicted that there was a reduction in dengue transmission after a nationwide release of wMel-Wolbachia-infected mosquitoes. In addition, they predicted about 86% of the estimated 7.8 million annual cases of symptomatic dengue in Indonesia could be averted following a complete nationwide rollout of Wolbachia-infected mosquitoes. Recently, a modeling study presented a dengue transmission model in the presence of female wild-type and wMelPop Wolbachia-infected Aedes aegypti mosquitoes. They concluded that although the wMelPop strain reduces the lifespan of infected mosquitoes, which could be challenging to achieve replacement of wild-type mosquitoes, its optimal release ensured the replacement of wild-type mosquitoes and also reduced dengue burden in the human population⁵¹. A mosquito-Wolbachia model was developed by Xue et al, to compare the potential effectiveness of two Wolbachia

strains (*w*Mel and *w*AlbB) to control arboviral spread⁶⁰. They observed that each of the two different strains of *Wolbachia* can effectively decrease the rate of arboviral transmission.

Here, we develop a general *Wolbachia* model capable of faithfully replicating all of the strain features described in Table 1. The general transmission model is an extention of the *Wolbachia* transmission model introduced in Adekunle et al.³⁷, which described the competitive dynamics between (*w*Mel-like) *Wolbachia*-infected and uninfected mosquitoes. Despite the non-induction of CI in *w*Au-*Wolbachia*-infected mosquitoes, *w*Au infection is retained and able to block viral transmission efficiently compared to other strains even at high temperature. Therefore, we incorporated this feature to determine if the advantages (*Wolbachia* retainment) of the *w*Au strain outweigh the ineffectiveness of CI. This feature has not been considered in previous models. Furthermore, we incorporate imperfect maternal transmission into the model. By analysing the system via computing the basic reproduction number(s) and investigating the stability properties of the equilibrium points, the potential of the *w*Au strain as a viable strategy to control *Aedes*-borne infections can be established. The aim of this modeling approach is to support future *Aedes*-borne viral control programs, particularly with the introduction of new *Wolbachia* variants.

Methods

Model formation. Here, we investigate a modified *Wolbachia* transmission model studied in Adekunle et al.³⁷, focusing on a novel *Wolbachia* strain, wAu, which has high retainment, high viral blockage and does not induce CI. The mosquito population is subdivided into two groups: the uninfected mosquitoes (.)_w and the *Wolbachia* infected mosquitoes (.)_w. The term (.) can be aquatic/immature (eggs, larvae and pupae) A, male M or female F mosquitoes. In addition, we denote the aquatic/immature stages, mature male and mature female uninfected mosquitoes as A_w , M_w , F_w respectively. As in Adekunle et al.³⁷ the model also incorporates the IMT of wAu-wOlbachia.

There are four possible mosquitoes' mating pairs: F_uM_u , F_uM_w , F_wM_u and F_wM_w . As Wolbachia infection is maternally transmitted, F_uM_u and F_uM_w will produce uninfected offspring while F_wM_u and F_wM_w will typically produce infected offspring. However if there is imperfect maternal transmission, the two latter strategies could produce some proportions of uninfected offspring²³.

To mathematically write the system of differential equations governing the *Wolbachia* transmission dynamics, we express the feasible mating strategies of uninfected and *Wolbachia* infected mosquito populations together with their per capita egg laying rates as Eqs. (1)–(6):

$$\frac{dA_u}{dt} = \left[\frac{\rho_{uu}(F_u M_u + (1 - \phi)F_u M_w) + \rho_{ww}((1 - \delta)F_w M_u + (1 - \nu)F_w M_w)}{M} \right] \left(1 - \frac{A}{K} \right)$$

$$= (\tau + u + v + A) A$$
(1)

$$\frac{dF_u}{dt} = (1 - \psi)\tau_u A_u + \sigma F_w - \mu_u F_u, \tag{2}$$

$$\frac{dM_u}{dt} = \psi \tau_u A_u + \sigma M_w - \mu_u M_u \tag{3}$$

$$\frac{dA_w}{dt} = \left[\frac{\rho_{ww}(\nu F_w M_w + \delta F_w M_u)}{M}\right] \left(1 - \frac{A}{K}\right) - (\tau_w + \mu_{Aw}) A_w,\tag{4}$$

$$\frac{dF_w}{dt} = (1 - \psi)\tau_w A_w - \sigma F_w - \mu_w F_w,\tag{5}$$

$$\frac{dM_w}{dt} = \psi \tau_w A_w - \sigma M_w - \mu_w M_w,\tag{6}$$

where $F = F_u + F_w$, $M = M_u + M_w$, $A = A_u + A_w$.

Here, ϕ represents the CI effect which can be either 0 if there is no CI, or 1 if CI is present. σ is the effect of LWI, such that it can either be 0, if there is no *Wolbachia* loss or greater than zero otherwise. In Adekunle et al. ³⁷ where CI is assumed and LWI is considered, these quantities are set to $\phi=1$ and $\sigma\geq0$. In our modified model, considering different strains with the exception of wAu strain, $\phi=1$ and σ could vary from values greater than zero onwards. However, for the wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-w

To mathematically express the above schematics, we have that, the feasible mating strategies of uninfected and *Wolbachia* infected mosquito populations together with their per capita egg laying rates are given by the following differential system:

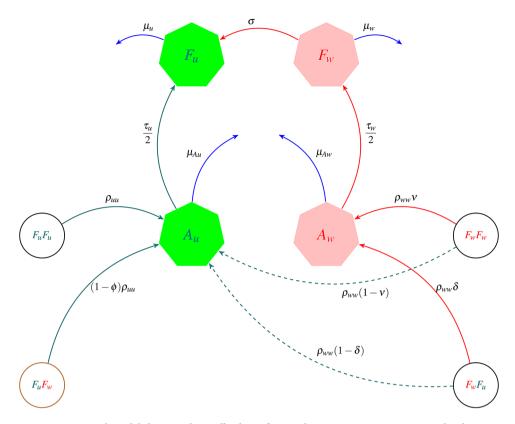


Figure 1. General model showing the *Wolbachia* infection dynamics in mosquitoes as M has been set equal to F. The green and pink compartmental polygons represent wild-type and *Wolbachia*-infected mosquitoes respectively. A_u and F_u represent the aquatic (eggs, larvae and pupae) and adult female mosquitoes for the uninfected mosquito population respectively while A_w and F_w represent their *Wolbachia* infected counterparts. The teal and red arrows illustrate the population progression of uninfected and *Wolbachia*-infected mosquitoes respectively. The four circles (three black and one brown) represent the mosquito mating strategies. The effect of cytoplasmic incompatibility (ϕ) , i.e. for wAu and wMel strains, $\phi = 0$ and $\phi = 1$ respectively, is illustrated by the brown-circled F_u F_w . The dashed lines represent the proportion of uninfected offspring caused by imperfect maternal transmission (IMT). The blue lines depict mosquito mortality. If there is loss of *Wolbachia* infection (LWI), $\sigma > 0$. But if there is no LWI as in wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-wAu-

$$\frac{dA_u}{dt} = \left[\frac{\rho_{uu}(F_u^2 + (1 - \phi)F_uF_w) + \rho_{ww}((1 - \nu)F_w^2 + (1 - \delta)F_wF_u)}{F}\right] \left(1 - \frac{A}{K}\right) - (\tau_u + \mu_{Au})A_u, \quad (7)$$

$$\frac{dF_u}{dt} = \frac{\tau_u}{2} A_u + \sigma F_w - \mu_u F_u,\tag{8}$$

$$\frac{dA_w}{dt} = \left[\frac{\rho_{ww}(\nu F_w^2 + \delta F_w F_u)}{F}\right] \left(1 - \frac{A}{K}\right) - (\tau_w + \mu_{Aw}) A_w,\tag{9}$$

$$\frac{dF_w}{dt} = \frac{\tau_w}{2} A_w - \sigma F_w - \mu_w F_w,\tag{10}$$

where $F = F_u + F_w$ and $A = A_u + A_w$. Before proceeding, we rescale each of our state variables according to the maximum total population size, which by Adekunle et al., 2019^{37} is set by

$$\begin{split} A_u(t) + F_u(t) + A_w(t) + F_w(t) &\leq K + \frac{\tau_u K}{2\mu_u} + \frac{\sigma \tau_w K}{2\mu_u (\mu_w + \sigma)} + \frac{\tau_w K}{2(\mu_w + \sigma)} \\ &\leq K \left(1 + \frac{1}{2} \left(\frac{\tau_u}{\mu_u} + \frac{\tau_w}{(\mu_w + \sigma)} \left(1 + \frac{\sigma}{\mu_u} \right) \right) \right) \\ &\leq \alpha K \end{split}$$

where
$$\alpha = 1 + \frac{1}{2} \left(\frac{\tau_u}{\mu_u} + \frac{\tau_w}{(\mu_w + \sigma)} \left(1 + \frac{\sigma}{\mu_u} \right) \right)$$
.

The closed set

$$\Omega = \{ (A_u, F_u, A_w, F_w) \in \mathbb{R}^4_+ | A_u + F_u + A_w + F_w \le \alpha K \}$$

which is a feasible region for the above system dynamics is positively invariant³⁷. Hence, we let $\bar{A}_u = \frac{A_u}{\alpha K}$, $\bar{A}_w = \frac{A_w}{\alpha K}$, $\bar{F}_u = \frac{\bar{F}_u}{\alpha K}$, $\bar{F}_w = \frac{\bar{F}_w}{\alpha K}$, $\bar{A} = \bar{A}_u + \bar{A}_w$ and $\bar{F} = \bar{F}_u + \bar{F}_w$. Also, letting $\nu = 1$, we assume a perfect maternal transmission for the reproduction outcome of $\bar{F}_w \bar{M}_w$ mating. Therefore, the general Wolbachia model in terms of population proportions is given by Eqs. (12)-(14). Hereafter it is clear that we refer to the scaled values of each state variable and as such drop the overbar from our notation. The scaled model below now evolves in the feasible region $\bar{\Omega}$, where $\bar{\Omega} = \{(\bar{A}_u, \bar{F}_u, \bar{A}_w, \bar{F}_w) \in \mathbb{R}^4_+ | \bar{A}_u + \bar{F}_u + \bar{A}_w + \bar{F}_w \leq 1 \}$.

$$\frac{d\bar{A}_u}{dt} = \left[\frac{\rho_{uu}(\bar{F}_u^2 + (1 - \phi)\bar{F}_u\bar{F}_w) + \rho_{ww}(1 - \delta)\bar{F}_w\bar{F}_u}{\bar{F}}\right](1 - \alpha\bar{A}) - (\tau_u + \mu_{Au})\bar{A}_u,\tag{11}$$

$$\frac{d\bar{F}_u}{dt} = \frac{\tau_u}{2}\bar{A}_u + \sigma\bar{F}_w - \mu_u\bar{F}_u,\tag{12}$$

$$\frac{d\bar{A}_w}{dt} = \left[\frac{\rho_{ww}(\bar{F}_w^2 + \delta\bar{F}_w\bar{F}_u)}{\bar{F}}\right] \left(1 - \alpha\bar{A}\right) - (\tau_w + \mu_{Aw})\bar{A}_w,\tag{13}$$

$$\frac{d\bar{F}_w}{dt} = \frac{\tau_w}{2}\bar{A}_w - \sigma\bar{F}_w - \mu_w\bar{F}_w. \tag{14}$$

The modeling of wAu-Wolbachia transmission dynamics has not been done as this a distinction from other Wolbachia transmission models. Unlike the modeling work in Adekunle et al.³⁷, apart from the non-induction of CI, we considered the loss of Wolbachia infections due to seasonal fluctuation in temperature, a key dynamics that is absent in wAu strain.

Results

Analysis of the model. The above general model (11)–(14) is parametrically adjusted to simultaneously accommodate wAu and wMel Wolbachia strains. For the wAu-Wolbachia model, we set $\phi = \sigma = 0$ and for the wMel-Wolbachia model, we set $\phi=1,\sigma>0$. The wMel-Wolbachia model parameter adjustments correspond to the model studied in Adekunle et al.³⁷.

Here, we want to analyse the general model(11)–(14) with arbitrary values of ϕ and σ to enable comparison with wAu-Wolbachia and Adekunle et al. 2019³⁷ models. Analysing the model for wAu, we have four steady states. The first steady state $e_1 = (0, 0, 0, 0)$ indicates non-existence of mosquitoes. The second $e_2 = (A_u^*, F_u^*, 0, 0)$ signifies the steady state for the uninfected mosquito population only. The third $e_3 = (0, 0, A_w^*, F_w^*)$ describes the equilibrium point for wAu-infected mosquitoes only. Lastly, the $e_4=(A_u^*,F_u^*,A_w^*,F_w^*)$ is the equilibrium point for the co-existence of both uninfected and wAu-Wolbachia-infected mosquito populations.

*Non-existence mosquito population, e*₁. The equilibrium point e_1 is trivial and is not biologically realistic. However, we can gain some insights into the competitive model dynamics by examining the case where there is no interaction between the uninfected and Wolbachia-infected mosquitoes. In other words, we want to investigate how each population would behave in the absence of the other. In particular, we derive the reproduction number of the uninfected R_{0u} and Wolbachia-infected R_{0w} mosquito populations when they do not interact:

$$R_{0u} = \frac{\rho_{uu}\tau_u}{2\mu_u(\mu_{Au} + \tau_u)},\tag{15}$$

$$R_{0w} = \frac{\rho_{ww} \tau_w}{2\mu_w (\mu_{Aw} + \tau_w)},\tag{16}$$

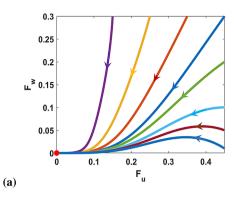
where the factor of $\frac{1}{2}$ in R_{0u} and R_{0w} stems from the choice to set $M = F^{62}$, i.e. $\psi = \frac{1}{2}$.

These reproductive numbers determine if the uninfected and Wolbachia-infected mosquito populations will die out or persist when there is no interaction. Specifically, if $R_{0u} < 1$ and $R_{0w} < 1$, then the two populations will die out (Fig. 2a). We observed in the decoupled case, the expressions for R_{0u} and R_{0w} are independent of the effects of CI (ϕ) and LWI (σ) and are therefore equivalent for both the wAu and wMel-Wolbachia strains (Fig. 2)³⁷.

Uninfected mosquito population, e2. The uninfected-mosquito-only equilibrium point or Wolbachia-free equilibrium is

$$e_2 = \left(\frac{1}{\alpha} \left[1 - \frac{1}{R_{0u}} \right], \frac{\tau_u}{2\mu_u \alpha} \left[1 - \frac{1}{R_{0u}} \right], 0, 0 \right).$$

For e_2 to exist, we require $R_{0u} > 1$. In addition to the uncoupled reproduction numbers (R_{0u} and R_{0w}) we also define the invasive reproduction number $R_{0w|u}$ which describes the average number of secondary offspring that



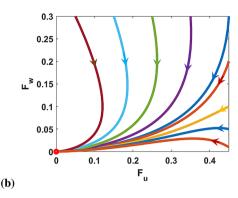


Figure 2. Graphs showing the system trajectories in the (F_u, F_w) plane for (a) wAu ($\phi = \sigma = 0$) and (b) wMel ($\phi = 1, \sigma = 0.04$) Wolbachia models when max[R_{0u}, R_{0w}] < 1. The red ball point indicates the point of stability, that is (F_u, F_w) = (0,0) representing mosquito extinction. We set $\rho_{uu} = 0.01$ and $\rho_{ww} = 0.1$. Other parameters used for these model simulations are provided in Table 2.

will become *Wolbachia*-infected adults after introducing a single adult *Wolbachia*-infected mosquito into an established *Wolbachia* uninfected mosquito population.

To compute $R_{0w|u}$, we use the next generation matrix method⁶³ to obtain

$$R_{0w|u} = \frac{\delta R_{0w}}{R_{0u}},\tag{17}$$

where we have substituted in the definition of R_{0w} from Eq. (16). The invasive reproduction number $R_{0w|u}$ is the same for both wAu and wMel-Wolbachia strains as that derived in Adekunle et al.³⁷. This is because, the expression (17) clearly shows that the invasive reproductive number $R_{0w|u}$ is not dependent on the CI effect, ϕ or LWI, σ .

To check if the equilibrium point e_2 is stable, we compute the Jacobian of the system and evaluate it at e_2 . In particular, letting $z_1 = (\mu_{Au} + \tau_u)$ and $z_2 = (\mu_{Aw} + \tau_w)$, yields

$$J_{e_2} = \begin{pmatrix} -z_1 R_{0u} & \frac{\rho_{uu}}{R_{0u}} & z_1 (1 - R_{0u}) & \frac{(1 - \delta)\rho_{vw}}{R_{0u}} \\ \frac{\tau_u}{2} & -\mu_u & 0 & 0 \\ 0 & 0 & -z_2 & \frac{\delta\rho_{ww}}{R_{0u}} \\ 0 & 0 & \frac{\tau_w}{2} & -\mu_w \end{pmatrix}.$$

To obtain the characteristic equation of J_{e_2} , we have

$$|J_{e_2}-\lambda I|=0,$$

which becomes

$$(\lambda^2 + k_1\lambda + k_2)(\lambda^2 + l_1\lambda + l_2) = 0$$

where

$$k_1 = \mu_u + z_1 R_{0u},$$

$$k_2 = \mu_u z_1 (R_{0u} - 1),$$

$$l_1 = \mu_w + z_2,$$

$$l_2 = \mu_w z_2 (1 - R_{0w|u}).$$

Therefore, e_2 is locally asymptotically stable if and only if $R_{0w|u} < 1$ and $R_{0u} > 1$ (Fig. 4). This is also consistent with the study in Adekunle et al.³⁷ (See Table 3).

Wolbachia-infected mosquito population, e_3 . The wAu-infected-only equilibrium point is $e_3 = \left(0,0,\frac{1}{\alpha}\left[1-\frac{1}{R_{0w}}\right],\frac{\tau_w}{2\mu_w\alpha}\left[1-\frac{1}{R_{0w}}\right]\right)$. This again is consistent with Adekunle et al.³⁷.

For e_3 to exist we require $R_{0w} > 1$. By computation, the invasive reproductive number $R_{0u|w}$ with respect to uninfected mosquitoes is given as,

$$R_{0u|w} = \frac{R_{0u}}{R_{0w}} \left[(1 - \phi) + \frac{\rho_{ww}}{\rho_{uu}} (1 - \delta) \right] = \frac{cR_{0u}}{R_{0w}},\tag{18}$$

where $c = (1 - \phi) + \frac{\rho_{ww}}{\rho_{uu}}(1 - \delta)$. Clearly, $R_{0u|w}$ is dependent on ϕ . For the *w*Mel-*Wolbachia* strain, i.e. $\phi = 1$, $c = \frac{\rho_{ww}}{\rho_{uu}}(1 - \delta)$ which is equivalent to that of Adekunle et al.³⁷. However, for the *w*Au-*Wolbachia* strain, i.e. $\phi = 0$,

we have a modified expression of $c=1+\frac{\rho_{ww}}{\rho_{uu}}(1-\delta)$ in Eq. (18) because we do not assume CI. Therefore, $c\geq 1$ for wAu-Wolbachia strain. Computing the Jacobian at e_3 , we have:

$$J_{e_3} = \begin{pmatrix} -z_1 & \frac{\rho_{uu} + (1-\delta)\rho_{ww}}{R_{0w}} & 0 & 0\\ \frac{\tau_u}{2} & -\mu_u & 0 & 0\\ z_2(1-R_{0w}) & \frac{-(1-\delta)\rho_{ww}}{R_{0w}} & -z_2R_{0w} & \frac{\rho_{ww}}{R_{0w}}\\ 0 & 0 & \frac{\tau_w}{2} & -\mu_w \end{pmatrix}.$$

The characteristic equation of J_{e_3} is then

$$|J_{e_3} - \lambda I| = (\lambda^2 + m_1 \lambda + m_2)(\lambda^2 + n_1 \lambda + n_2) = 0,$$

where

$$m_1 = \mu_u + z_1,$$

$$m_2 = \mu_u z_1 (1 - R_{0u|w}),$$

$$n_1 = \mu_w + z_2 R_{0w},$$

$$n_2 = \mu_w z_2 (R_{0w} - 1).$$

Therefore, e_3 is locally asymptotically stable if and only if $R_{0u|w} < 1$ and $R_{0w} > 1$ (see Fig. 4). The condition is equivalent to that found in³⁷ with generalized expressions for $R_{0u|w}$ used in place of the reduced version presented there (see Table 3).

Coexistent mosquito populations, e4. The equilibrium point for which both the uninfected and Wolbachia-infected populations coexist is

$$e_4 = (\frac{2\mu_u \beta F_w^*}{\tau_u}, \beta F_w^*, \frac{2\mu_w F_w^*}{\tau_w}, F_w^*)$$
 where

$$F_w^* = \frac{1}{2\alpha} \left[\frac{\left(1 - \frac{\xi}{R_{0w}}\right) \tau_u \tau_w}{(\mu_w \tau_u + \beta \mu_u \tau_w)} \right],$$

$$\beta = \frac{R_{0w}(R_{0u|w} - 1)}{R_{0u}(R_{0w|u} - 1)}$$
 and $\xi = \frac{(\beta + 1)}{(\delta \beta + 1)}$. For e_4 to exist, we require $R_{0w} > \xi > 1$ and

- (i) $R_{0w|u}, R_{0u|w} > 1$ or
- (ii) $R_{0w|u}, R_{0u|w} < 1$.

The above conditions (i) and (ii) correspond to the cases for $\delta > \frac{1}{c}$ and $\delta < \frac{1}{c}$ respectively. Comparing these existence conditions with those found above for e_2 and e_3 , we see that condition (ii) for the existence of e_4 matches the combined existence and local asymptotic stability condition for e_2 and e_3 . In other words, e_2 , e_3 and e_4 can coexist, while e_1 always exists (see Fig. 4).

To establish whether e_4 is stable or not, we compute the Jacobian J_{e_4} evaluated at e_4 to obtain the following characteristic equation:

$$|J_{e_4} - \lambda I| := \lambda^4 + s_1 \lambda^3 + s_2 \lambda^2 + s_3 \lambda + s_4 = 0.$$
 (19)

Let

$$z_3 = (\mu_u + \mu_w), z_4 = (\beta \rho_{uu} + \rho_{ww}), z_5 = (\beta + 1)\rho_{uu} + (1 - \delta)\rho_{ww}, z_6 = 1 + \beta(2 + \beta\delta),$$

$$z_7 = (\beta + 1)^2 \rho_{uu} + (1 - \delta)\rho_{ww}, z_8 = (1 + \beta(2 + \beta\delta))\rho_{uu} + (1 - \delta)\rho_{ww},$$
 then we have:

$$\begin{split} s_1 &= z_1 + z_2 + z_3 + \alpha z_4 F_w^*, \\ s_2 &= \mu_u \mu_w + z_3 (z_1 + z_2 + \alpha z_4 F_w^*) - \frac{\xi}{2R_{0w}(\beta+1)^2} (z_6 \rho_{ww} \tau_w + z_7 \tau_u), \\ s_3 &= \mu_u \mu_w (z_1 + z_2 + \alpha z_4 F_w^*) + z_3 \left[z_1 z_2 + \frac{\alpha}{\beta+1} (z_1 (1+\beta\delta) \rho_{ww} + \beta z_2 z_5) F_w^* \right] \\ &- \frac{\xi}{2R_{0w}(\beta+1)^3} \{ \left[(\mu_u + z_1) z_6 + z_8 \alpha \beta F_w^* \right] (\beta+1) \rho_{ww} \tau_w \\ &+ \left[\alpha \beta (1-\delta) z_5 \rho_{ww} F_w^* + z_7 (\alpha (1+\beta\delta) \rho_{ww} F_w^* + (\mu_w + z_2) (\beta+1)) \right] \tau_u \}, \\ s_4 &= \mu_u \mu_w \left[z_1 z_2 + \frac{\alpha}{\beta+1} (z_1 (1+\beta\delta) \rho_{ww} + \beta z_2 z_5) F_w^* \right] - \frac{\xi}{2R_{0w}(\beta+1)^2} \{ \left[z_2 z_6 + z_8 \alpha \beta F_w^* \right] \mu_u \rho_{ww} \tau_w + \left[\alpha \beta (1-\delta) z_5 \rho_{ww} F_w^* + z_7 (\alpha (1+\beta\delta) \rho_{ww} F_w^* + z_2 (\beta+1)) \right] \mu_w \tau_u \\ &- \frac{\xi}{2R_{0w}} \left[z_8 \rho_{ww} \right] \}. \end{split}$$

In order to establish the nature of the equilibrium point e_4 , we performed numerical testing using the Monte Carlo method in ⁵⁰ to verify the conditions (i) and (ii) by computing the real part of the eigenvalues of the Jacobian matrix, evaluated at e_4 . Simulation results are illustrated in Fig. 3.

Although the conditions (i) and (ii) indicated the existence of e_4 , Fig. 3c showed that e_4 is locally stable for condition (i) as all the eigenvalues (real part) are negative (λ_1 , λ_2 , λ_3 , λ_4 < 0). Whilst Fig. 3f showed that e_4 is unstable for condition (ii) as two of the eigenvalues (real part) are positive i.e. λ_3 , λ_4 > 0.

Numerically, we illustrated the existence and stability regions for e_4 in Fig. 4 for the two conditions (i) and (ii) relating to CI and maternal transmission (MT).

Following a modeling study of *Aedes aegypti* mosquitoes and normal *Wolbachia* (in the presence of CI only) interaction analyzed by Ferreira et al. 64 , three equilibrium points: trivial (q_1) ; uninfected only (q_2) ; and coexistence (q_3) , were obtained. However, the *Wolbachia*-only equilibrium point was not computed. The established local stability conditions for q_1 and q_2 correspond to that of the *wMel-like Wolbachia* conditions for e_1 and e_2 respectively. For coexistent populations to persist, the reproductive number for infected mosquitoes only, R_i must be greater than 1 and $R_i > R_u$, where R_u is the reproductive number for wild-type mosquitoes only. The model R_i also described the fitness parameter space between R_u and R_i , showing the change in extinction and persistence of the three equilibria when there is an increase in the initial population proportion of the *Wolbachia*-infected mosquitoes. Our model showed the changes in the no-mosquito, wild-type only, *Wolbachia*-only and coexistence population persistence and extinction in the presence and absence of CI with high and low maternal transmission (MT).

Figure 4 illustrates the existence and local stability regions for the equilibrium points e_1 , e_2 , e_3 and e_4 with respect to the reproduction numbers R_{0u} and R_{0w} as well as the relative magnitude of δ and $\frac{1}{c}$. For $\delta > \frac{1}{c}$ (high MT), Fig. 4a,b describe the dynamics for $\phi = 0$ (CI absent) and $\phi = 1$ (CI present) respectively. Within the subset of the yellow region of these figures bounded by $R_{0u} = 1$, $R_{0w} = 1$, and $R_{0w} = \xi$ we find that only e_1 and e_3 exist. Since e_3 is unstable in this region, we expect the system trajectories to tend to the no-mosquito equilibrium e_1 . This was confirmed through numerical simulations shown in Fig. 5a. For the existence of e_4 we require $R_{0u|w} > 1$, $R_{0w|u} > 1$ and $R_{0w} > \xi$ for stability (within the blue region). But if $R_{0w} < \xi$, e_1 is stable (yellow).

For $\delta < \frac{1}{c}$ (low MT), Fig. 4c,d portrayed the regions of stability for $\phi = 0$ and $\phi = 1$ respectively. The conditions $R_{0u} < 1$, $R_{0w} > 1$, and $R_{0u|w} > 1$ project the trajectiory to tend to e_1 (see Fig. 5b). In the orange region, e_1 and e_3 exist and are simultaneously locally stable as $R_{0u|w} > 1$ and $R_{0w} > \xi$. In addition, we have that e_4 exists where $R_{0u|w} < 1$ and $R_{0w|u} < 1$ (condition (ii)). With these conditions, e_4 exists together with e_2 and e_3 (white region). In this white region, e_2 and e_3 are locally stable even as $R_{0u} > 1$, $R_{0w} > 1$ but e_4 is unstable. Also, e_1 exists when $R_{0w} > 1$ and $R_{0u} < 1$ because the local stability of other equilibrium points is violated with these conditions. When $R_{0u|w} > 1$ but $R_{0u} < 1$ and $R_{0w} > 1$, the only stable outcome is the mosquito-free (no-mosquito) equilibrium e_1 . This occurs when R_{0u} is less than but still close to one. In this region, uninfected mosquitoes are capable of dominating initially when introduced into a *Wolbachia* saturated equilibrium because imperfect maternal transmission achieves $R_{0u|w} > 1$. This competitive advantage drives out the *Wolbachia* infected mosquitoes leaving uninfected mosquitoes only, which then are unable to sustain their population because $R_{0u} < 1$ (Fig. 5).

With the rate of high maternal transmission (MT) in the absence of CI (like-wAu), the reproductive advantage favours the production of uninfected mosquito offspring as it tends to accommodate more coexistent mosquito populations with wild-type than wMel-like strain (presence of CI) due to the presence of CI (Fig. 4a,b). Whilst, with a low MT rate, the CI presence or absence would favour Wolbachia-infected mosquitoes or uninfected mosquitoes respectively. In other words, the coexistent equilibrium point is unstable for the two mosquito populations as these conditions are equivalent to the local stabilities of both Wolbachia-free and Wolbachia-only equilibrium points (Fig. 4c,d). If $R_{0w} < \xi$, the system trajectories tend to the no mosquito equilibrium e_1 .

The conditions for the local stability of all equilibrium points are shown in Table 3 below.

Sensitivity analysis of *Wolbachia* **model**. To carry out the sensitivity analysis we investigate the model robustness due to uncertainties associated with parameter value estimations. In other words, we examine how senitive the invasive reproductive numbers are with respect to these parameters. This in turn, gives insight on influential parameters and their impact in reducing (or increasing) mosquito-type populations. To carry out this, we compute the normalized sensitivity indices of the invasive reproduction numbers with respect to the parameters used in the model.

Definition. The normalized forward sensitivity index of a variable ν with respect to parameter w is defined as:

$$\Lambda_w = \frac{\partial v}{\partial w} \times \frac{w}{v}.\tag{20}$$

Using the above formular (20), we contruct the following plots in Fig. 6.

From Fig. 6 and using the baseline parameter values for the wAu-Wolbachia strain in Table 2, it is clear that the reproductive and mortality rates for both wild-type (ρ_{uu} , μ_u) and wAu-Wolbachia-infected (ρ_{ww} , μ_w) mosquitoes and the proportion of wAu-Wolbachia-infected offspring (δ) have the most sensitivity in the invasive reproductive numbers $R_{0w|u}$. Whilst for $R_{0u|w}$, μ_u and μ_w are the most sensitive parameters. Hence for both invasive reproductive numbers, the most sensitive parameters are μ_u and μ_w . This demonstrates that an increase (or decrease) in the mortality rate of wAu-Wolbachia-infected mosquitoes by 10% will decrease (or increase) $R_{0w|u}$ by 10%.

Does CI (ϕ) **outweigh the LWI** (σ)? For most *Wolbachia* strains except *w*Au, the mating between uninfected female and *Wolbachia*-infected male mosquito crosses generates no viable offspring. However, *Wolbachia*-infected mosquitoes tend to lose their *Wolbachia* infection and lower their maternal transmission rate at high

Parameters	Description	Values (wMel)	Values (wAu)	Dimension	References
ρ_{uu}	Reproduction rate (egg laying rate) from mating between F_u and M_u/M_w mosquitoes	13	13	Eggs/day	32,37,61
ρ_{ww}	Reproduction rate (egg laying rate) from mating between F_w and M_u/M_w mosquitoes	10	10	Eggs/day	30,37,61
δ	The proportion of Wolbachia infected eggs resulting from mating between M_uF_w mosquitoes	0.95	0.95	Dimensionless	30
ν	The proportion of Wolbachia infected eggs resulting from mating between $M_w F_w$ mosquitoes	1	1	Dimensionless	37
ϕ	The CI induction	1	0	Dimensionless	23
ψ	Fraction of eggs that are male	0.5	0.5	Dimensionless	37,62
K	Carrying capacity of the aquatic stage A	10 ⁶	10 ⁶	aquatic mosquitoes	37
σ	Loss of Wolbachia infection (LWI)	0.04	0	day ⁻¹	Assumed
τ_u	Maturation rate of A_u aquatic stage into adulthood (per capita)	0.11	0.11	day ⁻¹	30,61
τ_w	Maturation rate of A_w aquatic stage into adulthood (per capita)	0.11	0.11	day ⁻¹	30,61
μ_{Au}	A_u Aquatic stage mortality rate (per capita)	0.02	0.02	day ⁻¹	57
μ_{Aw}	A_w aquatic stage mortality rate (per capita)	0.02	0.02	day ⁻¹	57
μ_u	F_u adult mortality rate (per capita)	0.061	0.04316	day ⁻¹	23,37
μ_w	F_w adult mortality rate (per capita)	0.068	0.08079	day ⁻¹	23,37

Table 2. Mosquito-Wolbachia model notations.

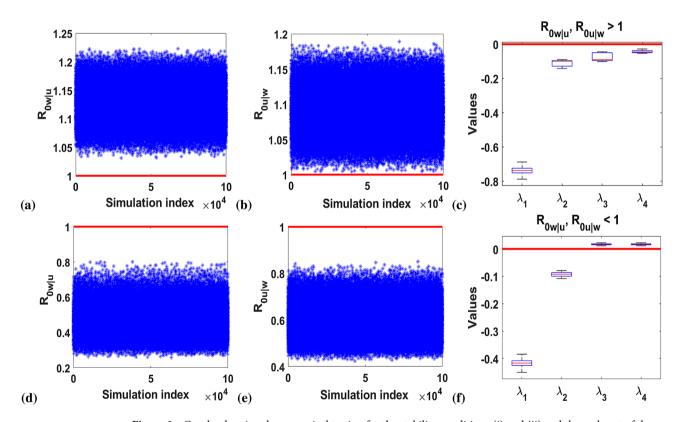


Figure 3. Graphs showing the numerical testing for the stability conditions (i) and (ii) and the real part of the eigenvalues' distribution (λ_1 , λ_2 , λ_3 and λ_4) for e_4 : (**a**,**b**) show that $R_{0w|u}$, $R_{0u|w} > 1$ always hold. (**c**) shows the related distribution of the real part of the eigenvalues for condition (i). (**d**,**e**) show the condition $R_{0w|u}$, $R_{0u|w} < 1$ always hold while (**f**) shows the corresponding distribution of the real part of the eigenvalues for condition (ii).

temperature $(27-37^{\circ}\text{C})^{23}$. With the effect of climate change gradually increasing the temperature by the day, *Wolbachia* strains with moderate or high temperature sensitivity such as *w*Mel may not be able to fully maintain a sufficient frequency level to invade the mosquito population.

In our general *Wolbachia* mathematical model, we describe a modified version of Adekunle et al.³⁷. This modification accommodates parameter adjustments for novel wAu and wMel-Wolbachia strains. For wAu, our mathematical model showed that despite the production of mosquito offspring due to CI absence, the invasive reproduction number due to infected mosquitoes $R_{0w|u}$ remains unchanged compared to the case where CI is present, as with the wMel-like strain³⁷. This further strengthened the fact that CI (inclusion or exclusion) does

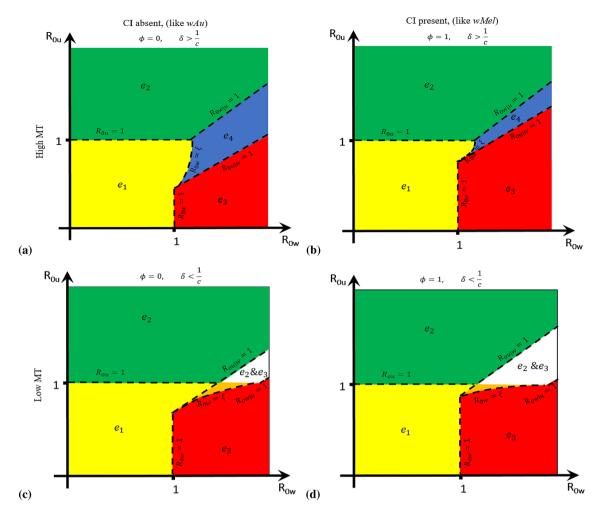


Figure 4. This graph shows the existence and local stability regions for the equilibrium points e_1 – e_4 for the *Wolbachia* model (11)–(14) as a function of the R_{0u} and R_{0w} relating to the cytoplasmic incompatibility (CI), ϕ and maternal transmission (MT), i.e. magnitude of δ and $\frac{1}{c}$. The yellow shaded region indicates the local stability of e_1 equilibrium. The green shaded area illustrates the local stability for the Wolbachia-free equilibrium point (e_2) . e_3 is locally stable at the red shaded part. The blue region indicates the coexistence local stability e_4 . The white region shows the existence of e_2 , e_3 and e_4 and local stability of e_2 and e_3 equilibrium points. And the orange region describes the existence and local stability of e_1 and e_3 . For $\delta > \frac{1}{c}$; (a) describes $\phi = 0$ as the boundary $R_{0w|u}=1$ sits above the boundary $R_{0u|w}=1$ and the arc $R_{0w}=\xi$. The co-existent equilibrium e_4 (blue), always sits in the region between these three boundaries because $R_{0w|u} > 1$, $R_{0u|w} > 1$ and $R_{0w} > \xi$. If $R_{0w} < \xi$, then e_1 becomes stable (yellow). (b) describes similar conditions as in (a) but for $\phi = 1$. We observed that the boundary $R_{0u|w} = 1$ shifts up while $R_{0w|u} = 1$ remained stationary to accommodate more e_3 . For $\delta < \frac{1}{\epsilon}$; (c) describes $\phi = 0$ as the relative position of e_4 boundaries in (a) flips so that boundary $R_{0u|w} = 1$ sits above boundary $R_{0w|u} = 1$ and the arc $R_{0w} = \xi$. Then, $R_{0w|u} < 1$ and $R_{0u|w} < 1$ and $R_{0w} > \xi$ shows the co-existence of e_2 and e_3 (white). However, e_2 and e_3 are locally stable in the white region as $R_{0w} > 1$ and $R_{01} > 1$. For $R_{0u} < 1$, e_2 and e_4 do not exist, only e_1 and e_3 do and if $R_{0w} > \xi$, e_1 and e_3 are locally stable (orange) and if $R_{0w} < \xi$, only e_3 becomes stable (red). (d) describes similar conditions as in (c) but for $\phi = 1$. It was observed that the boundary $R_{0u|w} = 1$ shifts up reducing the region of stability for e_2 .

not guarantee Wolbachia mosquitoes' persistence. Also, the invasive reproduction number due to uninfected mosquitoes expression $R_{0u|w}$ for wAu is similar to wMel, except that the expression depends on CI the effect. This is because, the mosquito gender crosses due to non-induction of CI for wAu, i.e. F_uM_w , generates uninfected offspring with perfect maternal transmission while wMel does not. The chances of establishing Wolbachia infected mosquitoes are lower when CI is ineffective compared to when it is induced. That is, for cytoplasmic inducing wMel-Wolbachia mosquitoes, the effect of LWI outweighs CI effect as mosquitoes still lose their infections (Fig. 7). However, wAu-Wolbachia infection retainment (no LWI) in mosquitoes has shown high level of maintaining the Wolbachia frequency in the absence of CI in mosquitoes (Fig. 7). This suggests that the LWI effect outweighs CI.

The LWI rate $\sigma(t)$ which is dependent on the seasons of the year can be modeled by a sinusoidal equation:

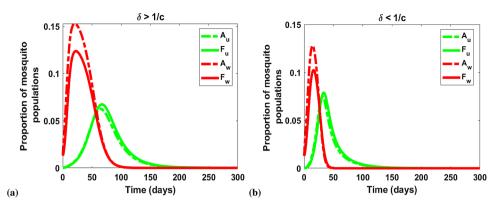


Figure 5. Graphs showing the local stability for e_1 relating to the magnitude of δ and $\frac{1}{c}$. The initial conditions for the state variables are $A_u(0) = 0.00015$, $F_u(0) = 0.00013$, $A_w(0) = 0.013$, $F_w(0) = 0.013$. We set $\rho_{uu} = 1$, $\rho_{ww} = 2.8571$, $\tau_u = \tau_w = 1$, $\mu_{Au} = \mu_{Aw} = 0.2$, $\mu_u = 0.4630$, $\mu_w = 0.6161$. (a) For $\delta > \frac{1}{c}$, where $\delta = 0.4$, c = 2.7143, $R_{0u} = 0.8999$, $R_{0w} = 1.9322$, $R_{0u|w} = 1.2641$, $R_{0w|u} = 0.8588$. (b) For $\delta < \frac{1}{c}$, where $\delta = 0.2$, c = 3.2857, $R_{0u} = 0.8999$, $R_{0w} = 1.9322$, $R_{0u|w} = 1.5303$, $R_{0w|u} = 0.4294$. The equilibrium point e_1 is locally stable if $R_{0u} < 1$, $R_{0w} > 1$, $R_{0w|u} < 1$ and $R_{0u|w} > 1$.

	Conditions for stability		
Equilibrium points	wMel ³⁷	wAu	
(i) No mosquitoes (e ₁)	$R_{0u} < 1 \text{ and } R_{0w} < 1$	$R_{0u} < 1 \text{ and } R_{0w} < 1$	
(ii) Uninfected mosquitoes only (e2)	$R_{0w u} < 1 \text{ and } R_{0u} > 1$	$R_{0w u} < 1$ and $R_{0u} > 1$	
(iii) Wolbachia-infected mosquitoes only (e ₃)	$R_{0u w} < 1 \text{ and } R_{0w} > 1$	$R_{0u w} < 1$ and $R_{0w} > 1$	
(iv) Both mosquitoes (e ₄)	$\delta < 1, \mu_u < \delta \mu_w, R_{0w} > 1 \text{ and } R_{0u} > 1$	$R_{0w u} > 1, R_{0u w} > 1, R_{0w} > 1 \text{ and } R_{0u} > 1$	

Table 3. Expressions for the condition for stability associated with the equilibrium points.

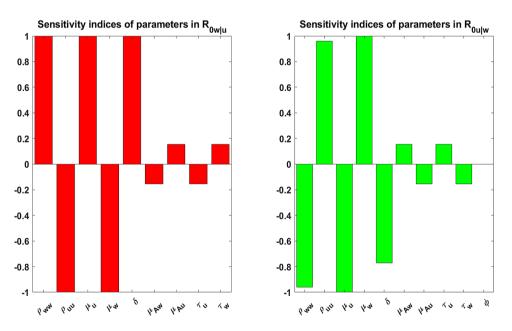


Figure 6. Plots showing the sensitivity indices of $R_{0w|u}$ and $R_{0u|w}$ the model parameters.

$$\sigma(t) = \frac{\sigma_{max}}{2} \left[1 + \cos\left(\frac{2\pi t}{365} - \mathscr{C}\right) \right]$$
 (21)

where σ_{max} is the maximum value of the seasonal variation in LWI, and \mathscr{C} is the phase shift which aligns the model with the seasonal change.

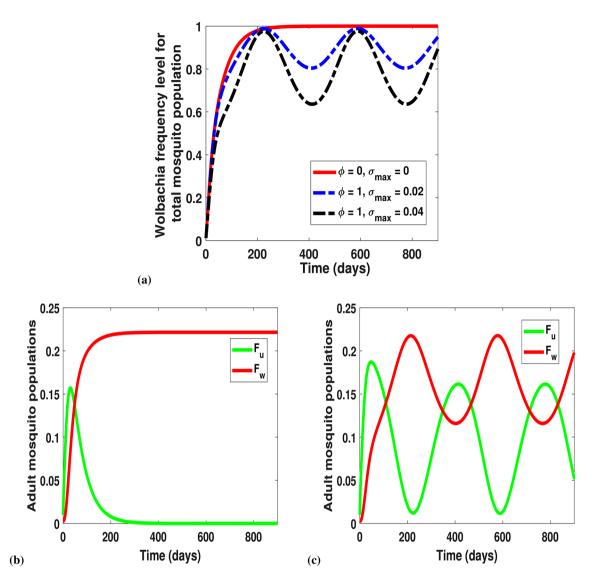


Figure 7. (a) Effect of CI induction ϕ and LWI $\sigma(t)$ on the *Wolbachia* frequency level. The initial conditions for the state variables are $A_u(0)=0.25$, $F_u(0)=0.01$, $A_w(0)=0$, $F_w(0)=0.003$. The red line indicates *Wolbachia* retainment as $\phi=0$ (no CI induction) and $\sigma_{max}=0$ (no LWI) which are features of *w*Au-*Wolbachia* strain. The blue and black dashed lines (for *w*Mel-*Wolbachia* strain) illustrate CI induction and LWI i.e $\phi=1$ for $\sigma_{max}=0.02$ and $\sigma_{max}=0.04$ respectively. Parameters for e_3 were used in these simulations. (b) Shows the dominance of *w*Au-*Wolbachia* infected F_w to uninfected F_u adult mosquitoes due to the retainment of *Wolbachia* infections (not affected by seasonal varying LWI). The *w*Au-*Wolbachia*-infected mosquitoes dominates when there is no CI $\phi=0$ and LWI $\sigma_{max}=0$ (red line). (c) For *w*Mel-*Wolbachia*-infected mosquitoes, the effect of seasonal varying loss of *Wolbachia* infection is shown as infections rise and drop continuously due to LWI $\sigma_{max}=0.04$ and CI induction $\phi=1$.

The effects of CI (ϕ) and LWI ($\sigma(t)$) as features of wAu and wMel Wolbachia strains are shown in Fig. 7. For the total mosquito population, wAu-infected mosquitoes ($\phi=0$, $\sigma_{max}=0$) reach the maximum frequency after approximately 250 days. To see the effect of CI induction and slight LWI i.e. $\phi=1$, $\mathscr{C}=0.25$, for $\sigma_{max}=0.02$ and $\sigma_{max}=0.04$, the Wolbachia frequency level oscillates between (0.8 and 1) and (0.6 and 1) respectively. That is, there is a 20% and 40% drop in the frequency level of Wolbachia when $\sigma(t)$ is at $\sigma_{max}=0.02$ and $\sigma_{max}=0.04$ respectively. This showed that, despite CI induction, LWI reduced the contribution of CI to the Wolbachia invasion (Fig. 7a). Therefore, the LWI gains highly outweigh the CI effect. By this, our analysis suggests that an increase in LWI in the presence of CI results in a drastic decrease in the Wolbachia frequency level (Fig. 7a). On the other hand, Fig. 7b showed the effect of LWI $\sigma(t)$ and CI ϕ with respect to the competitiveness between F_u and F_w . We observed that the F_w population dominates the F_u when there was no CI induction and Wolbachia infection is retained, that is, $\phi=0$, $\sigma_{max}=0$ (Fig. 7b). However, if CI induction occurs with loss of Wolbachia infections, then the seasonal varying effect occurs as seen in Fig. 7c.

Discussion

In this work, we modelled and investigated a general *Wolbachia* model that contained the transmission dynamics of *w*Au and *w*Mel *Wolbachia* strains in *Aedes* mosquitoes as special cases. These transmission dynamics described the competition between the novel *w*Au-*Wolbachia* infected *Aedes* mosquitoes and wild-type mosquitoes and compared the dynamics with the invasive properties of the popular *w*Mel-*Wolbachia* infected mosquitoes. We first derived the *Wolbachia* infection-status reproduction numbers for our *w*Au-*Wolbachia* model and used them to establish the conditions for the local stability of the equilibrium points for the *w*Au-*Wolbachia* invasive model. The reproduction number associated with the uninfected mosquitoes shows the reproductive advantage that the wild type has over the *w*Au strain. The comparison of the *w*Au-*Wolbachia* model (CI and LWI absent) and *w*Mel-*Wolbachia* model (CI and LWI present) showed that the *w*Au strain has the potential of compensating for the undesirable features of the *w*Mel strain.

Additionally, this study has reviewed the main features of different *Wolbachia* strains (Table 1) and shown that the *w*Au *Wolbachia* strain is a promising candidate for efficient *Aedes*-borne arboviral transmission control. Moreover, we analyzed the system dynamics of a general *Wolbachia* invasion model and determined the regions of local stability for each of the identified equilibrium points, highlighting the regions in parameter for which *Wolbachia*-infected mosquito populations persist or go extinct. This work modelled the general *Wolbachia* dynamics which can accommodate various *Wolbachia* characteristics regarding the presence or absence of CI and seasonal changes, unlike Adekunle et al.³⁷, which considers only the presence of CI. We also investigated the advantages gained from CI and LWI. This study has demonstrated that despite the absence of CI, the *Wolbachia* frequency level will drop as much as tenfold of the percentage of *Wolbachia* infection lost. We showed that the advantage of *Wolbachia* retainment in mosquitoes strongly outweighed the negative impact of CI indicating *w*Au *Wolbachia* strains may be suitable for arboviral control. Therefore, this modeling work contributes to the previous studies^{37,54,57,64,65} and helps close the gap between ways of maintaining the *Wolbachia* frequency levels in the absence of LWI and CI.

One implementation question for using the wAu strain as a replacement of the wMel strain is whether the wAu strain is self-sustaining, given that it does not induce CI. In this work, the equilibrium points for the wAu-Wolbachia model are the same as that for the wMel-Wolbachia model except that stricter conditions are required to satisfy the wAu-Wolbachia model equilibrium points. These more stringent conditions translate to additional resources such as the continuous introduction of a larger scale of wAu-infected mosquitoes to ensure replacement⁶⁶. Thus, the wAu strain is a promising alternative strain as it does not suffer from LWI due to high weather temperature and is highly effective in preventing the transmission of the arbovirus^{23,39,67}. Otherwise, combining the two strains may also be a good strategy.

There are limitations associated with any mathematical modeling work, and this study is not exempted. We first assumed the same mosquito gender ratio and expected this proportion to be constant over time. This assumption may be true in a laboratory setting⁶², but not necessarily true in a natural mosquito habitat. However, similar conclusions are expected to be reached as the *Wolbachia* model reduction accurately reproduces the dynamics of the full system⁶⁸. Secondly, we assumed that the absence of CI implies that cross mating resulted in offspring that are uninfected. This may not be true as a small proportion of the offspring may be *Wolbachia* infected²³. If that is the case, then it means that lesser resources will be required to use the *w*Au strain as a *Wolbachia*-based control strategy. Lastly, we assumed the seasonality affects the associated parameters for the *w*Mel dynamics. However, for the *w*Au strain, it is not affected by seasonality as *w*Au-*Wolbachia* infections are retained at high temperature.

Although several studies^{22,38,42,57} have demonstrated that CI drives the persistence of *Wolbachia*-infected *Aedes* mosquitoes, these studies neglected the impact of *Wolbachia* loss in mosquitoes. The CI drive has been shown in four mating lines (see Fig. 1) involving a *Wolbachia*-transinfected *Aedes* mosquitoes mating with wild-type mosquitoes. One of the mating lines for which *Wolbachia*-infected male and uninfected female mosquitoes produced no viable offspring (via CI) truncates the uninfected offspring from being produced as infection is maternally transmitted. With the exception of the mating between the uninfected male and female mosquito line, all other mating lines produce *Wolbachia*-infected offspring leading to persistence. In addition, high temperature affects these *Wolbachia*-infected mosquitoes as they lose their infection due to the unfavourable weather conditions. However, mosquitoes infected with the *w*Au-*Wolbachia* strain have been shown to not only block arboviral transmission efficiently, but also retain the *Wolbachia* infection at typically unfavourable high temperatures. This retainment of infection in mosquitoes strongly outweighed the absence of CI for the *w*Au strain in the establishment and dominance of *w*Au-*Wolbachia* infected mosquitoes.

While vaccine implementation may have been highly effective on dengue seropositive persons in high transmission areas ^{11,12}, the introduction of *Wolbachia*-infected mosquitoes in low and moderate arboviral endemic areas has also effectively shown successful reduction in dengue burden ^{43,51,59,69}. Given that these two strategies could reduce the transmission of *Aedes*-borne diseases, in particular, dengue depending on the transmission level, a modeling study by Ndii⁷⁰ proposed the use of these combined strategies and compared their effectiveness. The author showed that, *Wolbachia* performs better in the presence of low vaccine efficacy, but is outperformed otherwise ⁷⁰. Therefore combining the two strategies may be useful, however understanding both the temperature and seasonality effects on *Wolbachia* intervention programs, and serotypic differences relating to cross-protective immunity to investigate vaccine efficacy is necessary for the reduction and control of *Aedes*-borne arboviral disease transmission.

In conclusion, we have shown that the *w*Au-*Wolbachia* strain could be effective in controlling arbovirus transmission, as its advantages in terms of *Wolbachia* infection retention in mosquitoes may outweigh the absence of CI. This could prove even more promising, especially as the temperature increases due to climate

change. Although wMel and wAlbB-Wolbachia strains only have been rolled out in natural mosquito habitats in replacement programs, combining these strains with wAu is worth exploring.

Data availability

No datasets were generated or analysed during the current study.

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Author contributions

S.T.O and A.I.A. conceived the project concept; S.T.O and A.I.A. performed the model formation and interpretation. S.T.O, A.I.A. and M.T.M. analysed the results. S.T.O., A.I.A., M.T.M., D.P.R. and E.S.M. contributed in drafting the manuscript. All authors have read, reviewed and approved the final manuscript.

Competing interests

The authors declare no competing interests.

Additional information

Correspondence and requests for materials should be addressed to S.T.O.

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Author/s:

Ogunlade, ST; Adekunle, AI; Meehan, MT; Rojas, DP; McBryde, ES

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