Title: Warm temperature inhibits cytoplasmic incompatibility induced by endosymbiotic
Rickettsiella in a spider host

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ABSTRACT

Bacterial endosymbionts manipulate reproduction in arthropods to increase prevalence in the host

population. One such manipulation is cytoplasmic incompatibility (CI), wherein the bacteria sabotage

sperm in infected males to reduce hatch rate when mated with uninfected females, but zygotes are

"rescued" when that male mates with an infected female. In the spider *Mermessus fradeorum*

(Linyphiidae), *Rickettsiella* symbionts cause variable levels of CI. We hypothesized that temperature

affects CI strength and rescue in *M. fradeorum,* potentially mediated by bacterial titer. We reared

Rickettsiella-infected spiders in two temperature conditions (26°C vs 20°C) and tested CI induction in

males and CI rescue in females. In incompatible crosses between infected males and uninfected females,

hatch rate from warm males was doubled (Mean±S.E. = 0.687±0.052) relative to cool males

(0.348±0.046), indicating that CI induction is weaker in warm males. In rescue crosses between infected

females and infected males, female rearing temperature had a marginal effect on CI rescue, but hatch

rate remained high for both warm (0.960±0.023) and cool females (0.994±0.004). Bacterial titer as

measured by qPCR was lower in warm than cool spiders, particularly in females, suggesting that

bacterial titer may play a role in causing the temperature-mediated changes in CI.

Keywords

INTRODUCTION

Maternally-inherited bacterial symbionts are common in arthropods, some of which can

manipulate host reproduction to increase bacterial prevalence within the host population.

Cytoplasmic incompatibility (CI) is the most common and well-studied manipulation, sabotaging

matings between infected males and uninfected females, preventing the production of

uninfected offspring, and increasing the relative proportion of infected offspring in the

population over time (Shropshire et al. 2020). CI is typically viewed as a modification/rescue

system: The symbiont induces CI by modifying male gametes, which cannot successfully fertilize

and develop into progeny unless the egg is also infected and contains an appropriate rescue

factor (Shropshire et al. 2020). Numerous bacterial clades have been shown to cause CI,

including *Wolbachia* (Alphaproteobacteria; reviewed in Shropshire et al. 2020), *Cardinium*

(Bacteroidetes; Hunter et al. 2003), *Mesenet* (Alphaproteobacteria; Takano et al. 2021),

Spiroplasma (Mollicutes; Pollman et al. 2022), and *Rickettsiella* (Gammaproteobacteria;

Rosenwald et al. 2020). The mechanistic underpinnings of CI are only well characterized for

Wolbachia (Shropshire et al. 2020), but genomic and phenotypic studies with other symbionts

suggest novel modes of action may be present (Poinsot et al. 2003, Penz et al. 2012, Pollmann

et al. 2022). Consequently, caution should be used when extrapolating inferences from

Wolbachia CI studies to other bacterial symbionts.

There is growing evidence that symbiont-induced host phenotypes, such as CI, can be thermally

sensitive but potentially idiosyncratic (Corbin et al. 2017, Doremus et al. 2019, Shropshire et al.

2020, Chrostek et al. 2021, Corbin et al. 2021, Hague et al. 2022, Jones et al. 2023, Martins et al.

2023). *Wolbachia*-induced phenotypes generally appear to weaken with increased temperature

(e.g., Clancy and Hoffmann 1998, Ross et al. 2017), which likely has consequences for the global

deployment of this symbiont in vector management programs (Ross et al. 2023). Other

symbionts, such as *Spiroplasma*, show a reversed pattern, with weaker effects on their hosts at

decreased temperature (Anbutsu et al. 2008, Corbin et al. 2021). The effect of temperature on

Rickettsiella-induced CI has not previously been investigated, but in a non-CI-inducing

Rickettsiella strain in aphids, temperature affects *Rickettsiella* titer, host phenotype, and spread

in the host population (Gu et al. 2023). More generally, several studies have suggested that

thermal conditions might limit the geographic distribution of symbionts, their phenotypes, and

consequently even their host species (Zhang et al. 2019, Ross et al. 2020, Hague et al. 2022).

Thus, under global climate change, it becomes critical to understand thermal effects on

arthropod symbionts, because symbiont-induced phenotypes are likely to change and affect the

spread and performance of both pest and beneficial arthropod hosts.

Here, we test whether temperature affects the strength of *Rickettsiella*-induced CI and bacterial

titer in the linyphiid spider *Mermessus fradeorum* (Berland, 1932). Few studies of manipulative

symbionts have been conducted in non-insect arthropods, which can be infected with well-

known symbiotic taxa such as *Wolbachia*, and lesser known manipulative clades such as

Rickettsiella. Populations of *Mermessus fradeorum* are composed of individuals that are

naturally co-infected with different combinations of several such maternally inherited

symbionts, of which only *Rickettsiella* has been shown to induce CI (Curry et al. 2015,

Rosenwald et al. 2020). The mechanism by which *Rickettsiella* induces CI is unknown; however,

we have observed the strength of CI to be quite variable. We hypothesized that temperature

mediates the penetrance of the phenotype, potentially mediated by bacterial titer. Using

laboratory-reared colonies of spiders that were either infected only with *Rickettsiella* or were

uninfected because they came from lineages that had been experimentally cured via antibiotics

(Rosenwald et al. 2020), we reared spiders under two temperature regimes (26°C or 20°C) and

evaluated the ability of infected males to induce CI and infected females to rescue CI as a

function of temperature. Because the strength of symbiont-induced phenotypes is often

mediated by bacterial quantity (Lopez-Madrigal and Duarte 2019), we also evaluated bacterial

titer as a function of temperature.

EXPERIMENTAL PROCEDURES

To test the effect of temperature on *Rickettsiella*-induced CI in *M. fradeorum*, we conducted

separate experiments on CI-induction in males versus CI-rescue in females. We assessed

induction versus rescue separately to maximize experimental power on informative crosses. We

selected two temperature treatments, warm (26°C) and cool (20°C) that fall within the range of

temperatures the spiders would experience in the field, and at which we knew the spiders

could be successfully reared and propagated. *Rickettsiella*-infected (R-infected) spiders were

reared under both temperature regimes, while uninfected spiders were only reared under cool

conditions because the experiments were designed to test for effects of temperature on CI, not

on the spiders *per se*.

A cohort of R-infected (n=17) and uninfected (n=13) mother *M. fradeorum* were mated and

allowed to lay eggmasses over the span of 2-3 weeks. As each clutch of spiderlings hatched,

they were split into individual 4 cm diameter rearing cups with moistened plaster at the bottom

for humidity control (Rosenwald et al. 2020). Half of the R-infected spiderlings were randomly

assigned to be reared in a 20°C environmental chamber and half in a 26°C chamber; all

uninfected spiderlings were reared at 20˚C. All were maintained on a diet of *Sinella curviseta*

collembola when young and *Drosophila melanogaster* fruit flies when older (Rosenwald et al.

2020). After spiders reached maturity, they were randomly assigned to mating crosses in either

male CI induction or female CI rescue experiments. All experimental spiders were 7-10 weeks of

age at mating; these spiders typically live 6-10 months under laboratory conditions, hence age

differences of a couple of weeks are trivial.

For the male CI induction experiment, the experimental test cross was to compare CI induced

by R-infected males reared under warm versus cool conditions. These males (n=30 per

treatment) were mated to uninfected females to evaluate CI induction. To ensure that hatch

rate reductions observed in the CI crosses were not due to innate male infertility, these same

males were mated one week later to cool-reared R-infected females (compatible cross). Male

M. fradeorum are capable of mating multiple times and we have not detected decreases in

fertility or CI strength with subsequent matings (J. White, unpublished data). To control for

potential differences in innate female fertility, additional sets of cool-reared R-infected (n=15)

and uninfected (n=15) females were mated with uninfected males.

For the female CI rescue experiment, the experimental test cross of interest was to compare CI

rescue by R-infected females reared under warm versus cool conditions. These females (n=30

per treatment) were mated to cool-reared R-infected males to evaluate CI rescue. Because we

could not use the same re-mating technique with females as males, we mated separate sets of

warm and cool R-infected females (n=15 per treatment) to uninfected males, as controls for

female fertility. We included an additional control cross between cool R-infected males and

uninfected females (n=15), to ensure that males used in this experiment were capable of

causing CI. Finally, an additional set of warm-reared R-infected males (n=15) was also crossed

with uninfected females to aid in comparison across the two experiments.

After matings for each experiment were complete, male spiders were placed in 1.5mL

microcentrifuge tubes and stored at their rearing temperature without food for five to seven

days to ensure that gut contents had been emptied. Specimens were then preserved in 95%

ethanol and stored at -20˚C for further analysis. Female spiders were maintained under

standard rearing conditions at their assigned temperature until they deposited three

eggmasses. Then, females were moved to microcentrifuge tubes and underwent the same food

deprivation and preservation sequence as the males. To validate expected infection status, we

extracted DNA from a subset of 20 spiders using DNeasy Blood & Tissue Kit (Qiagen) following

the manufacturer's protocol. We performed diagnostic PCR for *Rickettsiella*, and two other

symbiont genera (*Tisiphia* and *Wolbachia*) known to be present in some *M. fradeorum* using

previously published protocols (wsp primers for *Wolbachia*, Baldo et al. 2006; Ricklong primers

for *Tisiphia*, Curry et al. 2015; RLA primers for *Rickettsiella*, Rosenwald et al. 2020). All tested

spiders were positive or negative for *Rickettsiella* as expected, and all spiders were negative for

the other symbionts.

To evaluate the hatch rate, we dissected each eggmass after spiderlings emerged, and counted

the remaining unhatched eggs as in Rosenwald et al. (2020). Total hatched and unhatched eggs

per female were used for a series of logistical regression contrasts, using Williams' correction

for overdispersion (Arc v 1.06; Williams 1982). In the CI induction experiment, we conducted

three statistical contrasts. First, to ensure CI was in evidence in the experiment, we compared

hatch rates in expected CI treatments (warm and cool R-infected males mated to uninfected

females) to control matings between uninfected males and uninfected females. Second, to

ensure that neither temperature nor *Rickettsiella* infection negatively affected male fertility, we

then contrasted hatch rates when these same males were mated to cool-reared *R*-infected

females, relative to uninfected males mated to R-infected females. Finally, we directly

contrasted the hatch rates of the warm and cool CI treatments to one another, to evaluate the

effect of rearing temperature on CI induction. In the CI rescue experiment we likewise

conducted three statistical contrasts. First, to ensure rearing temperature didn't directly affect

hatch rate, we contrasted hatch rates from warm versus cool R-infected females when mated

to uninfected males. Second, to ensure CI was in evidence in the experiment, we contrasted

cool R-infected females mated to cool R-infected males (cool rescue treatment), relative to cool

uninfected females mated to cool R-infected males (CI control). Finally, we directly contrasted

hatch rates of the warm and cool CI rescue treatments to one another, to evaluate the effect of

rearing temperature on CI rescue.

We evaluated symbiont titer via qPCR for a subset of 10-15 R-infected spiders from each sex

and temperature treatment. For the sake of consistency, all evaluated spiders originated from

the same experiment, the CI rescue experiment. We placed specimens into 50% PBS (EtOH:

PBS) for 30 min, followed by 75% PBS (EtOH: PBS) for 30 min, and finally kept in 100% PBS

overnight. We crushed these specimens in cryogenic vials containing solid-glass beads using

FastPrep® Tissue Homogenizer (MP Biomedicals), then extracted DNA using DNeasy Blood &

Tissue Kit (Qiagen) following the manufacturer's protocol. We designed short primers specific

to the *Rickettsiella recA* gene and *M. fradeorum 18S rRNA* gene using Geneious Prime Version

2021.2 (Table S1). Primer specificity was tested *in-silico* and using diagnostic PCR (Table S2). We

cloned these short specific sequences using a vector from CloneJET™ PCR Cloning Kit and

JM109 Mix & Go Competent Cells. Serial dilutions with serial factor 1:10 for the symbiont gene

and 1:3 for the spider gene were used for a standard calibration curve. qPCR was performed

using the Step One Plus real-time PCR system (Applied Biosystems, CA, USA) and Fast SYBR

green master mix (Thermo Fisher Scientific, MA, USA). For each 20 μL reaction mixture, we used

30 ng of gDNA, and each sample was measured in three technical triplicates. Cycling conditions

are listed in Table S2. We used output Ct, Intercept, and Slope values to calculate the amount

(ng) of the symbiont's gene and the spider's gene from each sample ($ng = 10^{\left(\frac{Ct-Intercept}{Slope}\right)}$). We

then used these weights to calculate the gene copy number (gcn) ($gcn =$

$\frac{ng \times gcn/mole}{bp \times ng/g \times g/mole\ of\ bp}$), followed by data normalization ($normalized\ symbiont\ titer =$

$\frac{symbiont\ recA\ gcn}{spider\ 18S\ gcn}$). We log-transformed the normalized titer values to reduce heteroscedasticity,

and compared means using a general linear model using Type III sums of squares in IBM SPSS v

28.0., with sex, temperature, and their interaction included as fixed factors.

RESULTS

When we reared R-infected spiders at either 26°C or 20°C and then evaluated the strength of CI

induction (Figure 1A), we found that both warm and cool R-infected males caused reduced

hatch rates when mated to uninfected females, relative to control matings between uninfected

Figure 1. Proportion hatch of eggs from

crosses between *Rickettsiella*-infected

Mermessus fradeorum spiders that were

individually reared at 26°C or 20°C, and

subsequently mated to either R-infected

or uninfected spiders reared at 20°C as

indicated on the x-axes. Realized sample

sizes per cross (excluding failed matings)

are indicated below each column.

Crosses to evaluate male CI induction

(Panel A) showed a significantly higher

hatch rate (weaker CI) for offspring of

warm-reared than cool-reared infected

males when mated to uninfected

females. All control crosses had high

hatch rates. Crosses to evaluate female

CI rescue (Panel B) showed marginally

lower hatch rate for warm-reared than

cool-reared infected females mated to

infected males, but high hatch rates

overall indicate temperature had

relatively little effect on CI rescue.

Incompatible crosses between infected

males and uninfected females were

included as controls to aid comparison

across the two experiments and validate

that infected males used within this

experiment were capable of inducing CI.

males and females (ΔDeviance = 58.95, d.f. = 2, P<0.001). When these same R-infected males

were mated with R-infected females, hatch rates were nearly perfect and did not differ from

control crosses between uninfected males and R-infected females (ΔDeviance = 2.87, d.f. = 2,

P=0.238), indicating that neither rearing temperature nor *Rickettsiella* infection intrinsically

decreased fertility of males. When compared directly, the hatch rate of incompatible crosses

between uninfected females and warm males was twice as high (0.687±0.052) as with cool

males (0.348±0.046; ΔDeviance = 21.30, d.f. = 1, P<0.001), indicating that warmer rearing

temperatures weakened CI induction.

In the CI rescue experiment (Figure 1B), hatch rates were high and equivalent for warm and

cool R-infected females mated to uninfected males (ΔDeviance=0.20, d.f.=1, P=0.654),

indicating that rearing temperatures did not directly affect hatch rate. Cool R-infected females

also exhibited nearly perfect hatch rates when mated to cool R-infected males, much higher

than cool uninfected females mated to cool R-infected males (ΔDeviance=84.49, d.f.=1,

P<0.001). This control contrast establishes both that the R-infected males in this experiment

induced CI, and that cool R-infected females were capable of rescue. When we contrasted

rescue between the two female rearing temperatures, we found that maternal rearing

temperature had a marginally significant effect on hatch rate (ΔDeviance = 3.825, d.f. =1,

P=0.0505), suggesting that CI rescue ability can be weakened in at least some R-infected

females. Nevertheless, most females in these rescue crosses exhibited perfect hatch rates,

indicating that the CI rescue ability of *Rickettsiella* usually remains intact regardless of rearing

temperature.

When we evaluated bacterial titer in a subset of warm-reared versus cool-reared spiders

(Figure 2), we found an interactive effect between sex and temperature (Sex $F_{1,44}$ =0.329,

P=0.529, Temperature $F_{1,44}$ =6.026, P=0.018, Sex × Temp interaction $F_{1,44}$ =5.22, P=0.027). Titer

was lower in the warm treatments, but this lower titer was associated with females more than

males, even though temperature effects on symbiont phenotype were more evident in males

(CI induction) than females (CI rescue).

DISCUSSION

Warmer rearing temperatures

strongly decreased CI induction by

Rickettsiella in male *M. fradeorum*

spiders, but only mildly decreased

CI rescue in female spiders.

Thermal sensitivity for *Rickettsiella*-

induced phenotypes is consistent

with results for a different strain of

Rickettsiella in an aphid host, which

exhibits a different set of host

phenotypic effects that don't

include CI (Gu et al. 2023). In the

present study, CI strength overall

was notably weaker than in prior

Figure 2. *Rickettsiella* titer in a subset of warm (26°C) and cool (20°C) reared spiders from the CI rescue experiment. Log transformed titer (normalized against spider 18S quantity) was significantly affected by the interaction between temperature and sex, primarily driven by lower titer in warm reared females.

studies by our group (Rosenwald et al. 2020), which were conducted at room temperature, but

in a laboratory that was cool throughout the duration of the experiments (J. White, pers. obs.).

Highly variable hatch rates across all CI crosses (Fig 1A), suggest that other factors may

additionally be driving CI strength in this system (e.g., Shropshire et al. 2021). More generally,

bacterial symbionts often show reduced phenotypic penetrance at warmer temperatures

(Martins et al. 2023), although the phenomenon is not universal, and instances where

symbiont-induced phenotypes increase with warmer temperatures also exist (e.g., Corbin et al.

2021).

Warmer rearing temperature also decreased *Rickettsiella* titer, particularly in female spiders,

which also tended to have lower titer than males. The latter was surprising, because gravid

females are usually expected to carry higher loads of maternally transmitted symbionts than

males (e.g., Noda et al., 2001). However, it is also possible that relatively high loads of bacteria

would be allocated to eggs, and these post-ovipositional females, which had laid 3 eggmasses

prior to preservation and extraction, may have exhibited substantially reduced bacterial titer

relative to their pre-ovipositional state. It is also worth noting that our standard laboratory

protocol calls for imposing several days of food deprivation to clear gut contents (and

associated prey symbionts) before spider specimen preservation. This protocol likely

diminished the utility of the qPCR assay, by temporally disassociating the titer of the preserved

specimen from its titer at biologically relevant timepoints (fertilization, egg deposition). Despite

these caveats, we nevertheless did detect a signal of temperature treatment on bacterial titer,

suggesting that stronger CI penetrance may be associated with higher bacterial titer. We will be

following up with assays more specifically designed to probe the mechanism and timing of CI

modification and rescue in this novel host/symbiont system. Associations between

temperature and symbiont titers have previously been demonstrated with other

endosymbionts (Breeuwer and Warren 1993, Lopez-Madrigal and Duarte 2019, Ross et al.

2020), and thus our results further support the potential links between temperature, titer, and

phenotypic outcome.

Our study demonstrates the thermal sensitivity of CI in a novel arthropod/symbiont system,

implying that temperature may be modulating host/symbiont interactions quite broadly in

nature (e.g., Kreisner et al. 2016, Hague et al. 2022). However, much additional study will be

needed before we can predict how temperature affects CI dynamics in the field. For *M.*

fradeorum, we selected only two relatively moderate temperatures to test, which the spider

likely experiences routinely in the field. We also kept each temperature constant, because we

do not yet know the mechanism or timing of *Rickettsiella*-induced CI modification and rescue.

Future studies will explore variations of timing, duration, and extent of thermal exposures on CI

in this system, which will inform our understanding of spider/symbiont biogeography for this

widespread introduced species within the context of global climate change.

More broadly, environmental effects on symbiont phenotypic penetrance may have important

implications in both managed and natural ecosystems. *Wolbachia* is already in use in vector

management in several areas of the globe, with mixed results (Hoffmann et al. 2011, Dos

Santos et al. 2022, Hoffmann et al. 2024). Symbiont thermal tolerance is one factor that

appears to limit the success of this technique in some regions (Ross et al. 2020, 2023). As the

use of symbionts to control pests and vectors comes under consideration in a range of other

systems (e.g., Gong et al. 2020, Mateos et al. 2020), it becomes critical to characterize the

environmental constraints that might influence efficacy. Even in less-managed systems, the

thermal optima of symbionts might play a critical role in determining the distribution of pests

and the ability of natural enemies to control them (Zhang et al. 2019, Hague et al. 2022). Our

study contributes to a growing body of literature that suggests environmental contingency may

be the rule for interactions between hosts and reproductive manipulators, not the exception.

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REFERENCES
Anbutsu, H., Goto, S. and Fukatsu, T. (2008). High and low temperatures differently affect
infection density and vertical transmission of male-killing *Spiroplasma* symbionts in *Drosophila*
hosts. Applied and Environmental Microbiology, 74, 6053-6059.

Breeuwer, J.A. and Werren, J.H. (1993). Cytoplasmic incompatibility and bacterial density in
Nasonia vitripennis. Genetics, 135, 565-574.

Chrostek, E., Martins, N., Marialva, M.S. and Teixeira, L. (2021). *Wolbachia*-conferred antiviral
protection is determined by developmental temperature. mBio, 12, 10-1128.

Clancy, D.J. and Hoffmann, A.A. (1998). Environmental effects on cytoplasmic incompatibility
and bacterial load in *Wolbachia*-infected *Drosophila simulans.* Entomologia Experimentalis et
Applicata, 86, 13-24.

Corbin, C., Heyworth, E.R., Ferrari, J., and Hurst, G.D.D. (2017). Heritable symbionts in a world
of varying temperature. Heredity, 118, 10-20.

Corbin C., J. E. J., Chrostek E., Fenton A., and Hurst G.D.D. (2021). Thermal sensitivity of the
Spiroplasma-Drosophila hydei protective symbiosis: The best of climes, the worst of climes.
Molecular Ecology, 30, 1336-1344.

Curry, M. M., Paliulis, L. V., Welch, K. D., Harwood, J. D., and White, J. A. (2015). Multiple
endosymbiont infections and reproductive manipulations in a linyphiid spider population.
Heredity, 115, 146-152.

Doremus M.R., K. S. E., and Hunter M.S. (2019). Exposure to opposing temperature extremes
causes comparable effects on *Cardinium* density but contrasting effects on *Cardinium*-induced
cytoplasmic incompatibility. PLoS Pathogens, 15, 1008022.

Dos Santos, G.R., Durovni, B., Saraceni, V., Riback, T.I.S., Pinto, S.B., Anders, K.L., Moreira, L.A.
and Salje, H. (2022). Estimating the effect of the wMel release programme on the incidence of
dengue and chikungunya in Rio de Janeiro, Brazil: a spatiotemporal modelling study. The Lancet
Infectious Diseases, 22, 1587-1595.

Gong, J.T., Li, Y., Li, T.P., Liang, Y., Hu, L., Zhang, D., Zhou, C.Y., Yang, C., Zhang, X., Zha, S.S. and
Duan, X.Z. (2020). Stable introduction of plant-virus-inhibiting *Wolbachia* into planthoppers for
rice protection. Current Biology, 30, 4837-4845.

Gu, S., Ross, P.A., Gill, A., Yang, Q., Ansermin, E., Sharma, S., Soleimannjad, S., Sharma, K.,
Callahan, A., Brown, C., Umina, P.A., Kristensen, T.N., and Hoffmann. A.A. (2023). A rapidly
spreading deleterious aphid endosymbiont that uses horizontal as well as vertical transmission.
PNAS, 120, 2217278120.

Hague, M.T., Shropshire, J.D., Caldwell, C.N., Statz, J.P., Stanek, K.A., Conner, W.R. and Cooper,
B.S. (2022). Temperature effects on cellular host-microbe interactions explain continent-wide
endosymbiont prevalence. Current Biology, 32, 878-888.

Hoffmann, A. A., Montgomery, B.L., Popovici, J., Iturbe-Ormaetxe, I., Johnson, P.H., Muzzi, F.,
Greenfield, M., Durkan, M., Leong, Y.S., Dong, Y., Cook, H., Axford, J., Callahan, A.G., Kenny, N.,
Omodei, C., McGraw, E.A., Ryan, P.A., Ritchie, S.A., Turelli, M., and O'Neill, S.L. (2011).
Successful establishment of *Wolbachia* in *Aedes* populations to suppress dengue transmission.
Nature, 476, 454-457.

Hoffmann, A.A., Ahmad, N.W., Keong, W.M., Ling, C.Y., Ahmad, N.A., Golding, N., Tierney, N.,
Jelip, J., Putit, P.W., Mokhtar, N. and Sandhu, S.S. (2024). Introduction of *Aedes aegypti*
mosquitoes carrying *wAlbB Wolbachia* sharply decreases dengue incidence in disease
hotspots. iScience, 27.

Hunter, M. S., Perlman S.J., and Kelly S.E. (2003). A bacterial symbiont in the Bacteroidetes
induces cytoplasmic incompatibility in the parasitoid wasp *Encarsia pergandiella*. The Royal
Society, Proceedings B, 270, 2185-2190.

Jones, J. E., Hurst, G.D.D. (2023). History matters: thermal environment before, but not during
wasp attack determines the efficiency of symbiont-mediated protection. Molecular Ecology, 32,
3340-3351.

Kriesner, P., Conner, W.R., Weeks, A.R., Turelli, M. and Hoffmann, A.A. (2016). Persistence of a
Wolbachia infection frequency cline in *Drosophila melanogaster* and the possible role of
reproductive dormancy. Evolution, 70, 979-997.

López-Madrigal, S. and Duarte, E.H. (2019). Titer regulation in arthropod-*Wolbachia*
symbioses. FEMS Microbiology Letters, 366, 232.

Martins, M., César, C.S. and Cogni, R. (2023). The effects of temperature on prevalence of
facultative insect heritable symbionts across spatial and seasonal scales. Frontiers in
Microbiology, 14, 1321341.Mateos, M., Martinez Montoya, H., Lanzavecchia, S.B., Conte, C.,
Guillén, K., Morán-Aceves, B.M., Toledo, J., Liedo, P., Asimakis, E.D., Doudoumis, V. and Kyritsis,
G.A. (2020). *Wolbachia pipientis* associated with tephritid fruit fly pests: from basic research to
applications. Frontiers in Microbiology, 11, 521402.

Noda, H., Koizumi, Y., Zhang, Q. and Deng, K. (2001). Infection density of *Wolbachia* and
incompatibility level in two planthopper species, *Laodelphax striatellus* and *Sogatella*
furcifera. Insect Biochemistry and Molecular Biology, 31, 727-737.

Penz, T., Schmitz-Esser, S., Kelly, S.E., Cass, B.N., Müller, A., Woyke, T., Malfatti, S.A., Hunter,
M.S., and Horn, M. (2012). Comparative genomics Suggests an independent origin of
cytoplasmic incompatibility in *Cardinium hertigii*. PLoS Genetics, 8, 1003012.

Poinsot, D., Charlat, S., and Merçot, H. (2003). On the mechanism of *Wolbachia*-induced
cytoplasmic incompatibility: confronting the models with facts. BioEssays 25, 259-265.

Pollmann M., M., L.D., Krimmer, P.D., Hasselmann, M., Perlman, S.J., Ballinger, M.J., Steidle,
J.L.M., and Gottlieb, Y. (2022). Highly transmissible cytoplasmic incompatibility by the
extracellular insect symbiont *Spiroplasma*. iScience, 25,104335.

Rosenwald L.C., Sitvarin, M. I. and White, J. A. (2020). Endosymbiotic *Rickettsiella* causes
cytoplasmic incompatibility in a spider host. The Royal Society, Proceedings B, 278, 20201107

Ross, P.A., Wiwatanaratanabutr, I., Axford, J.K., White, V.L., Endersby-Harshman, N.M. and
Hoffmann, A.A. (2017). *Wolbachia* infections in *Aedes aegypti* differ markedly in their response
to cyclical heat stress. PLoS Pathogens, 13, e1006006.

Ross, P.A., Axford, J.K., Yang, Q., Staunton, K.M., Ritchie, S.A., Richardson, K.M. and Hoffmann,
A.A. (2020). Heatwaves cause fluctuations in *wMel Wolbachia* densities and frequencies in
Aedes aegypti. PLoS neglected tropical diseases, 14, e0007958.

Ross, P.A., Elfekih, S., Collier, S., Klein, M.J., Lee, S.S., Dunn, M., Jackson, S., Zhang, Y., Axford,
J.K., Gu, X. and Home, J.L. (2023). Developing *Wolbachia*-based disease interventions for an
extreme environment. PLoS Pathogens, 19, e1011117.

Shropshire J.D., L. B., and Bordenstein S.R. (2020). Symbiont-mediated cytoplasmic
incompatibility: What have we learned in 50 years? eLife, 9, e61989.

Shropshire, J.D., Hamant, E. and Cooper, B.S., 2021. Male age and *Wolbachia* dynamics:
investigating how fast and why bacterial densities and cytoplasmic incompatibility strengths
vary. *MBio*, *12*, e02998-21.

Takano, S.I., Gotoh, Y. and Hayashi, T. (2021). "Candidatus *Mesenet longicola*": novel
endosymbionts of *Brontispa longissima* that induce cytoplasmic incompatibility. Microbial
Ecology, 82, 512-522.Williams, D.A. (1982). Extra-binomial variation in logistic linear
models. Journal of the Royal Statistical Society: Series C (Applied Statistics), 31, 144-148.

Zhang, B., Leonard, S.P., Li, Y. and Moran, N.A. (2019). Obligate bacterial endosymbionts limit
thermal tolerance of insect host species. Proceedings of the National Academy of Sciences, 116,
24712-24718