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Wolbachia improves the performance of an invasive fly after a diet shift

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ABSTRACT

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Polyphagy is an adaptative strategy allowing species to develop and survive on multiple diets. Phytophagous insects can be exposed to sub-optimal resources, as host plants vary in nutritional quality. Some insects may rely on symbiotic bacteria to better utilize less favorable substrates. The invasive fruit fly *Drosophila suzukii* is a highly successful pest worldwide. The use of ripening and ripe fruits, a niche neglected by other drosophilids, and its polyphagy may have contributed to its global spread. The role of D. suzukii's Wolbachia strain (wSuz) remains unconfirmed, although a mutualistic role has been hypothesized via virus protection or dealing with abiotic stress. In some insect species, Wolbachia acts as a nutritional mutualist. This study explored the role of wSuz in D. suzukii adaptation to fluctuating diet regimes. To simulate a diet shift, we used Wolbachia-infected and uninfected flies, maintained on a corn-based (low-starch) or a grape-based (highstarch) diet for at least 10 generations. Then, we placed individuals from each line on one or the other diet for 30 days. The effects of Wolbachia and the diet treatments were assessed by quantifying female fecundity, eggto-adult development time and survival, and offspring mass. The presence of Wolbachia positively affected female fecundity and offspring mass after a diet shift. Wolbachia also increased survival during larval development regardless of the diet, supporting its mutualistic role. Our results underline the role of wSuz on D. suzukii diet tolerance and performance. A better understanding of the role of symbionts in invasive species could help to improve management strategies.

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Keywords: Drosophila suzukii, endosymbiont, fecundity, mutualism, nutrition, polyphagy

KEY MESSAGE

- Generalist phytophagous insects may face diets varying in nutritional quality.
- Endosymbionts may contribute to better tolerance of diet shifts.
- The role of *Drosophila suzukii*'s *Wolbachia* strain (wSuz) is currently unconfirmed.
- We demonstrate a mutualistic role of wSuz, increasing survival and offspring mass after a diet shift.
- Understanding the role of symbionts in invasive species could help to improve management strategies.

4647 INTRODUCTION

The ability to develop and reproduce on several diets – known as polyphagy – is an adaptive strategy that is often successful in heterogeneous environments (Singer 2008). However, polyphagy among phytophagous insects is the exception rather than the rule as most hypotheses on niche breadth relate on host specialization (reviewed by Jousselin and Elias 2019). While specialists can only develop and survive on a very limited number of plant species, polyphagous insects can complete their cycle on a large spectrum of plants with varying quality. Encountering both low- and high-quality resources, the generalist strategy can be costly for individuals as offspring performance (e.g., developmental rate, survival) may vary between host plants (Janz 2003; Poyet et al. 2015). This cost could be offset by a female laying eggs on numerous host plants, ensuring that at least some offspring survive or achieve optimal development (Jaenike 1990; Singer 2008) and leading to a trade-off between host quality and availability (Singer 2008). Many successful pest insects are polyphagous (Singer 2008). This is the case for the fruit flies (Diptera: Tephritidae), which include some species with extreme polyphagy, for which selection pressure acts towards the ability of individuals to use any fruit that are readily available (Wilson et al. 2012).

Extreme polyphagy may act in synergy with other strategies to help improve performance when encountering a poor-quality host plant. A particularly important factor affecting insect host range is their associated microbiome (Frago et al. 2012; Sugio et al. 2015; Giron et al. 2018). For example, facultative secondary endosymbionts broaden the range of suitable host plants for the whitefly *Bemisia tabaci* (Benhamou et al. 2021). Associations between microbes and insect hosts can be beneficial or deleterious, facultative or obligatory (Sugio et al. 2015; Gurung et al. 2019). Amongst those with beneficial interactions, bacteria are the most studied organisms. They have a role in insect immunity, protection against parasitoids and parasites, helping insects coping with abiotic stress, as well as with nutritional stress (Dillon and Dillon 2004; Oliver et al. 2010; Eleftherianos et al. 2013; Sannino et al. 2018; Gurung et al. 2019). Bacteria can play multiple nutritional roles in insects, including improving their ability to live on suboptimal diets, increasing the efficiency of digestion, providing digestive enzymes, detoxifying defense compounds and provisioning vitamins (Dillon and Dillon 2004; Ashra and Nair 2022). For example, *Candidatus Erwinia dacicola*, an obligate symbiont of the tephritid olive fruit fly, *Bactrocera oleo* (Rossi), provides adult flies with essential amino acids that are absent in unripe olive fruits (Ben-Yosef et al. 2014). In *Drosophila melanogaster* larvae,

Acetobacter pomorum and Lactobacillus plantarum allow them to better cope with changes in nutrient availability, optimizing host fitness (Consuegra et al. 2020).

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Among bacterial endosymbionts, Wolbachia is widely acknowledged to be the most abundant and prevalent in insect species (Zug and Hammerstein 2015). Initially known for their reproductive parasitism that allows them to spread, research has increasingly shed light on Wolbachia's complex effects on hosts, ranging from parasitism to mutualism (Brownlie et al. 2009; Zug and Hammerstein 2015; Newton and Rice 2020; Pimentel et al. 2021). As mutualists, Wolbachia can protect the host against viruses (Chrostek et al. 2013; Cattel et al. 2016b; Pimentel et al. 2021), increase thermal tolerance (Saeed et al. 2018) and provide nutritional benefits (Brownlie et al. 2009; Hosokawa et al. 2010; Zug and Hammerstein 2015; Newton and Rice 2020; Lindsey et al. 2023). Nucleotides and vitamins have been suggested to be supplemented or modulated by Wolbachia (Newton and Rice 2020; Lindsey et al. 2023). In the bed bug Cimex lectularius, the Wolbachia strain wCle provides B vitamins to its host that are essential for growth and reproduction (Hosokawa et al. 2010; Nikoh et al. 2014). In D. melanogaster, Wolbachia plays a role in iron metabolism (Brownlie et al. 2009) and was recently found to be highly beneficial under nutrient limited conditions (Lindsey et al. 2023). The nutritional role of Wolbachia could explain its prevalence in some populations even when no reproductive manipulation is found (Newton et al. 2020). Overall, however, the role of Wolbachia as a potential promotor of invasion and/or pest success has received little attention (but see Lu et al. 2016). In this study, we explored its role in an important invasive fruit fly.

Originating from Asia, Drosophila suzukii Matsumara spread in the late 2000s into Europe, America and, more recently, Africa (Asplen et al. 2015; Kwadha et al. 2021). Unlike other *Drosophila* species that oviposit on rotting fruit, D. suzukii lays eggs on still ripening and ripe fruits using a serrated ovipositor (Atallah et al. 2014; Clemente et al. 2018). This ability, in addition with multiple generations per year, makes D. suzukii an important pest of soft and thin-skinned crops, resulting in significant economic losses for fruit growers (Walsh et al. 2011). Being highly polyphagous, the fly exploits a wide variety of cultivated (e.g., red fruits, cherry, plum, peach, grape; Walsh et al. 2011) and wild fruits (e.g., elderberry, mistletoe, wild black cherry; Poyet et al. 2015; Kenis et al. 2016; Ulmer et al. 2022), ensuring its persistence year-round (Poyet et al. 2015). However, this polyphagy exposes the fly to diets of differing quality, which results in various effects on its development (Poyet et al. 2015). The Wolbachia strain wSuz has been reported in D. suzukii (Cattel et al. 2016b). In nature, this strain has an imperfect maternal transmission (Hamm et al. 2014), while in laboratory the maternal transmission is perfect (Cattel et al. 2016a). It does not induce reproductive manipulation in American populations (Hamm et al. 2014; Cattel et al. 2016b), yet it induces moderate cytoplasmic incompatibility in European populations (Cattel et al. 2016b, 2018). Its prevalence in natural populations is higher than expected by chance, indicating that wSuz must provide benefits to its host (Cattel et al. 2016b), buts costs (see below) and drift (Oliver et al. 2014) certainly can play against the maintenance of Wolbachia within populations. Studies have attempted to identify these beneficial effects, but the available information remains inconclusive. Some studies have reported an increase in fecundity (Mazzetto et al. 2015) and others a decrease (Hamm et al. 2014) associated with Wolbachia presence. Hamm et al. (2014) found no effect of wSuz on starvation tolerance or desiccation resistance while Saeed et al. (2018) reported an increase in thermal tolerance. These results must be taken with caution as different host genetic backgrounds or the presence/absence of antimicrobial agents in the diet may have led to confounding effects (Cattel et al. 2016b). Cattel et al. (2016b) revealed protection against two viruses, Drosophila C virus and Flock House virus. However, no studies have yet addressed the potential effects of wSuz on nutritional provisioning or diet tolerance. Taking into account the high level of polyphagy displayed by *D. suzukii*, the variable resource availability across time and space, and the difference in wSuz prevalence across geographical ranges, we hypothesized that wSuz benefits *D. suzukii* by improving the fly's ability to deal with the diet shifts it experiences.

This study therefore investigated the role of the *D. suzukii Wolbachia* strain *w*Suz in dealing with nutritional shifts. We used two fly lines sharing the same genetic background but differing in the presence or absence of *Wolbachia*. Two diets were tested: a corn diet and a grape diet, which differed mainly in their starch content. The numbers of eggs produced, developmental time, offspring survival to adulthood and offspring mass were assessed following different combinations of nutritional diet shifts. We predicted differences in fecundity and developmental parameters between the two diets, regardless of the *Wolbachia* infection status of the flies. Also, we hypothesized that the presence of *Wolbachia* would increase fly survival after a diet shift, especially if *Wolbachia* plays a nutritional mutualistic role in the context of polyphagy, as described above.

MATERIALS AND METHODS

Rearing of D. suzukii stocks

Drosophila suzukii flies originated from a field collection made on infested raspberries in Rennes, France (48°7'2.158"N, 1°40'40.053"W), in October 2020, by the ECOBIO Laboratory (Rennes). Five hundred fruits from multiple plants were individually placed in vials for D. suzukii emergence. One hundred and ninety fly pairs were obtained and maintained separately on artificial diet (banana-based). In November 2020, nine lines infected by Wolbachia (wSuz) were detected using specific PCR assays with Wolbachia-specific primers targeting the wsp gene (81F/691R), following Braig et al. (1998). These infected lines were mixed in cages, and the population amplified. To obtain separate lines differing by the presence (W+) or absence (W-) of Wolbachia, half of the population was reared on a diet supplemented with tetracycline for four generations, as previously described by Hague et al. (2020). Then, the two lines were reared in 100 ml plastic bottles containing ~30 ml of standard cornmeal diet and incubated at 20°C, LD 12:12 h and 75% RH (STRADER, EV1300, Angers, France). At least 30 bottles of 50–200 flies per line were used for continuous maintenance. Before and after experiments, Wolbachia presence in the two lines was assessed by diagnostic PCR following the abovementioned method. To evaluate the impact of the tetracycline treatment on the other microbiota of the flies, we restored the microbiota of W- flies that have been raised for about 20 generation in the laboratory using W+ flies' faeces for three generations, as described by Strunov et al. (2022). Then we compared Wflies with or without microbiota restoration by performing a metabarcoding analysis with primers targeting the

variable region V3-V4 of the 16S rRNA gene to characterize Bacteria diversity and ITS2 to assess the diversity of fungi (see Supplementary Material).

Experimental diets

We used two diets to analyze the influence of different nutritional substrates on *D. suzukii* development, a standard cornmeal diet (Corn) and an industrial diet made from grape (Grape). These two media were contrasted in their composition, both were artificial but the Grape diet included grape powder, a fruit that *D. suzukii* can use for its development (e.g., Shu et al. 2022). The flies were divided into two populations reared on each diet and maintained for several generations (~10, six months) before starting the experiment. A detailed analysis of macronutrients (carbohydrates, lipids and proteins) in each diet was performed as described by

 $\begin{table}{l} \textbf{Table 1} Treatment codes corresponding to diet combinations for $Wolbachia$-infected and uninfected flies. N_i = initial sample size; N_f = final sample size after removing the data from females that died before the end of the experiment. } \label{eq:table_size}$

Foray et al. (2012). For analyses of amino acid, B vitamins and iron content, samples were sent to Eurofins[©].

Code	Treatment	N_i	$N_{\rm f}$
W+ C/C	Wolbachia-infected flies from corn culture placed on Corn	25	21
W+G/C	Wolbachia-infected flies from grape culture placed on Corn	25	21
W+G/G	Wolbachia-infected flies from grape culture placed on Grape	25	22
W+ C/G	Wolbachia-infected flies from corn culture placed on Grape	25	23
W-C/C	Wolbachia-uninfected flies from corn culture placed on Corn	25	19
W-G/C	Wolbachia-uninfected flies from grape culture placed on Corn	25	14
W-G/G	Wolbachia-uninfected flies from grape culture placed on Grape	25	22
W-C/G	Wolbachia-uninfected flies from corn culture placed on Grape	25	22

Effect of diet on D. suzukii development

Egg laying boxes were designed for the experiment: a hole was drilled in 60×45×50 mm crystal polystyrene boxes to receive a 6×30 mm 3D print egg laying cylinder containing 5 ml of diet. One female and three males aged 24 h and collected randomly from the culture vials were placed in each laying box. There were four diet combinations for *Wolbachia*-infected and uninfected flies, resulting in eight treatments (Table 1). Each treatment had 25 replicates (i.e., 25 females). C/C and G/G treatments were used as control as they correspond to the routine rearing. A pinch of active yeast was dusted on each laying cylinder to stimulate egg laying (Plantamp et al. 2017). The laying boxes were maintained under standard culture conditions (see above). Every day over 30 days, eggs were counted on each laying cylinder under a stereomicroscope. The laying cylinder was replaced every 48 h and the one removed was placed on a Petri dish containing ~30 ml of diet to continue larval development. Developmental conditions were the same as for the rearing (see above). For each Petri dish, the numbers of developed pupae and the numbers of emerged flies were recorded daily. Egg-to-pupa, pupa-to-adult and egg-to-adult survival and development time were assessed. The emerging adults were

counted and frozen at -20°C for later weighing. From each female and treatment, 120 emerging individuals were weighed on a laboratory precision balance (0.01 mg; QUINTIX125D-1S; Sartorius, Göttingen, Germany).

Statistical analyses

All data were analyzed in R version 4.1.2 (R Core Team 2022). Differences in macronutrient composition between diets (i.e., glucose, starch, proteins and lipids) were assessed using the Wilcoxon-Mann-Whitney test. Data from females that died before the end of the experiment were removed from the analyses (Table 1). The total numbers of eggs laid and numbers of offspring were compared among treatments with generalized linear models using a Poisson distribution. The chosen explanatory variables were: the *Wolbachia* infectious status, the laying diet and the presence of a diet shift between the diet from which the female originated and the one it was made to oviposit on. Egg-to-pupa, pupa-to-adult and egg-to-adult survival, mean development time and offspring mass were compared using analysis of variance (ANOVA) after square root transformation to achieve normality and homoscedasticity of error variance. Offspring sex-ratio was also analysed with ANOVA. Where ANOVA was significant, significant differences between means were then identified using Tukey's HSD tests or the emmeans() function from the emmeans package.

RESULTS

Diet analyses

Macronutrient analysis indicated that the Grape diet was enriched in carbohydrates (more starch) compared to the Corn diet, but that both diets provided a similar amount of proteins and lipids (Table S1). Therefore, the Grape diet provided more energy than the Corn. Analyses of amino acid, B vitamins and iron contents revealed that both diets were comparable in these three compound groups (Table S2). Therefore we concluded that any differences found in fly performance between the diet treatments must be interpreted as a consequence of varying starch levels, as the protein:total carbohydrate ratios were similar between the two diets (P:C Corn = 0.011 ± 0.006 ; P:C Grape = 0.010 ± 0.005 ; p = 0.84).

Microbiota analysis

The microbiota consisted almost exclusively of *Acetobacter* for bacteria and *Saccharomyces* and *Wallemia* for fungi (Fig. 1, see Supplementary Material for more details). No significant difference was found between W– flies with or without microbiota restoration. Because this assessment is indirect, we still cannot totally exclude the confounding effect of microbiota. However, the absence of difference in the microbiota of W– flies (many generations after exposure to antibiotic) before and after exposure to W+ faeces strongly suggests

that our W+ and W- lines shared the same microbiota. Moreover, the protocol to restore the microbiota of W- flies was used in previous studies and we are therefore confident that it provides adequate outputs. Therefore, we considered that the two lines W+ and W- share similar microbiota and that the results reported hereafter reflect the presence/absence of *Wolbachia*.



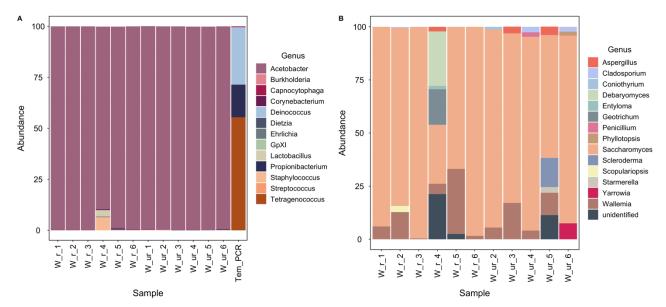


Fig.1 Metabarcoding analysis to characterize Bacteria (A) and fungi (B) diversity in *Drosophila suzukii*. W_r samples correspond to W- flies with microbiota restoration using W+ flies' faeces for three generations as described by Strunov et al. (2022) and W_ur to W- flies without microbiota restoration (see Supplementary Material for more details). Each column corresponds to one individual.

Effect of diet treatment and Wolbachia status on female fecundity

Overall, there was a large variation between females in the total numbers of eggs laid, from 18 to 568 eggs during the 30 days of the experiment. The total number of eggs laid by a female was significantly influenced by the laying diet, the diet shift, Wolbachia presence and the interaction terms (Table 2, Fig. 2A). Females laid significantly more eggs on Grape (mean \pm sd: 214 ± 134 eggs) than on Corn (157 ± 96 eggs). When confronted to a diet shift, females from the Corn line laid more eggs on Grape than they did on Corn (263 ± 143 vs. 201 ± 111 eggs; z = -32.4, p < 0.0001) and females from the Grape line laid fewer eggs on Corn than they did on Grape (107 ± 38 vs. 163 ± 105 eggs; z = 31.3, p < 0.0001). In addition, W+ females laid fewer eggs than W-, although the difference was not significant for females from C/C treatment.

Effect of diet treatment and Wolbachia status on offspring survival

Wolbachia presence was the only variable to impact significantly the egg-to-adult survival (Table 2, Fig. 2B). Mean egg-to-adult survival was about $66 \pm 14\%$ and did not differ significantly between diet treatments. Wolbachia presence increased survival across diet conditions (F = 39.2, p < 0.0001), although this was not

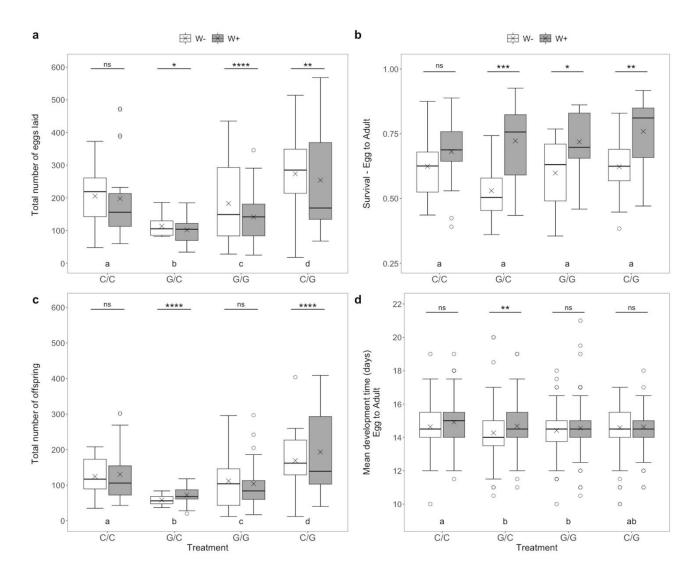


Fig. 2 Influence of diet treatment and *Wolbachia* infectious status (W+ or W-) on *Drosophila suzukii* reproductive parameters: (A) Total number of eggs laid in 30 d, (B) Egg to adult survival, (C) Total number of offspring per female (emerged adult flies), and (D) Mean development time from egg to adult emergence in days. The diet treatments were: flies from corn culture placed on corn diet (C/C), flies from grape culture placed on corn diet (G/G), and flies from corn culture placed on grape diet (G/G). Significance of differences between diet treatments and *Wolbachia* infection status are represented by different letters (a, b, c, d; bottom of each panel) or by asterisks, respectively (*: p < 0.05, **: p < 0.01, ***: p < 0.001, ***: p < 0.001, ns: non-significant; top of each panel)

significant for the C/C treatment, with a mean increase of 17%. *Wolbachia* presence primarily acted on larval survival (egg-to-pupa) and did not have a significant effect on pupal survival (Table S3, Fig. S1).

Overall, individual females produced between 12 and 409 surviving adult offspring over the experiment, with means of 145 ± 98 and 99 ± 60 offspring when laying on Grape and Corn respectively. The effect of diet treatment on the total number of offspring per female was the same as on the total number of eggs laid (Table 2, Fig. 2C). A mean of 67 ± 23 offspring was produced in the G/C treatment, almost a third of those produced in the C/G treatment (182 ± 106 offspring). The numbers of offspring produced in treatments not involving a diet shift were 108 ± 75 and 127 ± 67 for G/G and C/C, respectively. *Wolbachia* had a significant effect only

in the treatments involving a diet shift (C/G and G/C), where the numbers of offspring were increased by 14.1% and 23.1%, respectively.

Effect of diet treatment and Wolbachia status on developmental time

Across all treatments, the development time from egg to adult was 14.6 ± 1.1 days, with a minimum of 10 days and a maximum of 21 days. The laying diet, *Wolbachia* status and occurrence of a diet shift had an interactive effect on the development time (Table 2). The longest development time was observed on Corn when no diet shift occurred while the shortest times were observed on the G/C and G/G treatments (Fig. 2D). *Wolbachia* presence had a significant effect only on the G/C treatment, increasing the development time by almost half a day (W-: 14.3 ± 1.4 days, W+: 14.7 ± 1.2 days; F = 14.7674, p = 0.008). *Wolbachia* presence had no effect on egg-to-pupa development time (Table S4). The larval phase lasted about 8.2 ± 1.0 days and was significantly shorter in the G/C and G/G treatments, where larvae pupated from 3 to 7 hours earlier than in the other treatments (Fig. S2A). The pupal phase lasted 6.4 ± 0.8 days and even if *Wolbachia* presence had an overall effect on pupa-to-adult development time, the effect was not significant anymore within the diet treatment (Table S4, Fig. S2B). Diet treatment did not have a significant influence on pupa-to-adult development time (Table S4, Fig. S2B).

Effect of diet treatment and Wolbachia status on offspring mass and sex-ratio

Overall, female offspring were almost twice as heavy as males, with a mean mass of 1.28 ± 0.40 mg vs. 0.75 ± 0.24 mg. The diet treatments had comparable effects on female and male mass, with the heaviest

Table 2 Summary of results for generalized linear models for total number of eggs laid, egg-to-adult survival, total number of offspring, and egg-to-adult development time. Df = degree of freedom; Dev. = deviance; $P > |\chi^2| = P$ -value

	Total number of eggs laid			Egg-to-adult survival		Total number of offspring			Egg-to-adult development time			
Effect	Df	Dev.	$P > \chi^2 $	Df	Dev.	$P > \chi^2 $	Df	Dev.	$P > \chi^2 $	Df	Dev.	$P > \chi^2 $
Wolbachia (Wol)	1	136.71	<0.0001	1	0.605	<0.0001	1	9.05	0.0026	1	19.1	0.0001
Laying diet (LD)	1	670.46	<0.0001	1	0.041	0.105	1	723.91	<0.0001	1	5.5	0.039
Diet shift (S)	1	32.33	<0.0001	1	0.003	0.650	1	43.89	<0.0001	1	2.1	0.201
$Wol \times LD$	1	0.20	0.651	1	0.001	0.763	1	1.70	0.192	1	6.2	0.029
$\text{Wol} \times S$	1	0.07	0.791	1	0.045	0.090	1	13.92	0.0002	1	0.0	0.864
$LD \times S$	1	2112.90	<0.0001	1	0.029	0.174	1	1526.86	<0.0001	1	21.5	<0.0001
$Wol \times LD \times S$	1	24.75	<0.0001	1	0.035	0.134	1	0.46	0.498	1	1.5	0.282

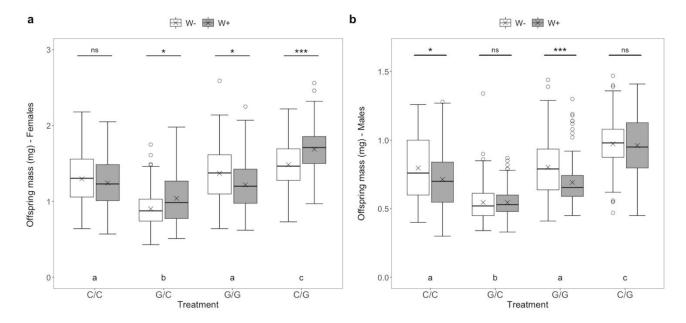


Fig. 3 Influence of diet treatment and *Wolbachia* presence (W+ or W-) on *Drosophila suzukii* offspring mass in mg with (A) females and (B) males. The treatments are the same as in Fig. 2. N = 120. Analysis of variance (ANOVA) was performed after square root transformation. Statistically significant differences are represented either by different letters (a, b, c, d; bottom of each panel) or by asterisks (*: p < 0.05, ***: p < 0.001; ns: non-significant; top of each panel)

individuals obtained in the C/G treatment and the lightest ones from the G/C treatment. When there was no diet shift (C/C and G/G treatments), the adult masses were intermediate (Table 3, Fig. 3). *Wolbachia* presence had an effect on the mass of both sexes produced, but only when a diet shift occurred (Table 4). When there was no diet shift, the mass of W+ offspring was significantly lower than those of W- for both males and females, although this was not significant in females from the C/C treatment (Fig. 3). In contrast, under diet shifts, the offspring mass did not differ in relation to infection status for males (Fig. 3B), and the W+ offspring were significantly heavier than W- offspring for females (Fig. 3A).

The mean offspring sex-ratio (females/males) was 0.51 ± 0.06 . Wolbachia had no effect on sex ratio (ANOVA, F = 1.099, p = 0.2961). The sex-ratio of offspring from G/C treatment was significantly higher (more females) than the sex-ratio of offspring from the G/G treatment (G/C: 0.53 ± 0.07 , G/G: 0.49 ± 0.06 , p = 0.0258). The sex-ratio did not differ between other treatments.

DISCUSSION

Across their lifetime, polyphagous insects are exposed to various diets that may change across generations and phenology. Polyphagy at the population level is generally interpreted as an individual developing on a single plant species, yet being able to develop on several hosts (Singer 2008), so that the female has a range of host plants on which to oviposit. The larvae produced can thereby be exposed to diets varying in quality. Endosymbiotic partners may contribute to improve the performance of insects faced with such diet shifts, and

Table 3 Summary of results from generalized linear models for female and male offspring mass. For details, see legend of Table 3

	Offs Fem	pring mass ale		Offs Male			
Effect	Df	Dev.	$P > \chi^2 $	Df	Dev.	$P > \chi^2 $	
Wolbachia (Wol)	1	0.048	0.132	1	0.93	0.0001	
Laying diet (LD)	1	5.006	<0.0001	1	19.61	<0.0001	
Diet shift (S)	1	0.015	0.409	1	0.09	0.238	
$Wol \times LD$	1	0.007	0.555	1	0.06	0.342	
$Wol \times S$	1	0.840	<0.0001	1	0.77	0.0004	
$LD\times S$	1	4.310	<0.0001	1	19.91	<0.0001	
$Wol \times LD \times S$	1	0.046	0.142	1	0.00	0.896	

could be especially critical for invasive species. Here, we show that the invasive fly *D. suzukii* performs better in terms of survival when it hosts the *Wolbachia* strain *w*Suz, in particular if it is faced with diet shifts. Hosting this endosymbiont may, however, also result in costs, since a negative effect was observed on other life history traits, such as fecundity or offspring mass. As the fly is an invasive species, and because it is present year-round in temperate regions, the influence of *Wolbachia* presence on its survival likely provides an overall advantage to *D. suzukii*. This effect may provide an important element explaining the maintenance of *w*Suz in natural populations and the success of this pest fly species worldwide. Caution is needed however when interpreting our findings, as they are specific to a single *Wolbachia* genotype and one *D. suzukii* population. To ensure broader applicability, it is essential to evaluate our results with various *D. suzukii* genotypes and *Wolbachia* strains, given that numerous studies has revealed that the influence of *Wolbachia* is contingent on both its own genotype and that of its host (Fry et al. 2004; Capobianco et al. 2018; Serga et al. 2021; Strunov et al. 2022).

Wolbachia presence and the tradeoff between fecundity and offspring survival

In all treatments, *Wolbachia*-infected females laid fewer eggs than non-infected ones, although this tendency was not significant for flies on Corn without diet shift. This apparent fitness cost was counterbalanced by the large increase seen in offspring survival, specifically larval survival, across all treatments, which resulted in almost perfect compensation in the number of offspring when the diet was stable, and even overcompensation (i.e., more surviving offspring in total) under diet shift treatments. This effect of *Wolbachia* on overall offspring survival contrasts with some previous studies. Hamm et al. (2014) reported a negative effect of *Wolbachia* presence on progeny production while Mazzetto et al. (2015) reported an increase of 30-50% in progeny production by *Wolbachia*-infected flies. However, the diet history of the flies was not taken into account in these studies. Mazzetto et al. (2015) reared flies on various organic fruits, then transferring

them to a standard diet, while Hamm et al. (2014) collected flies from an organic raspberry field and then reared them on blueberry. For both, no antibiotic was used on the fruits they used as diet and they only had a few populations, which may have led to genetic confounding effects. In our study, larval survival included both hatch rate and direct larval survival, which means *Wolbachia* could have acted either on one parameter or the other or both. *Wolbachia* is already known to act on hatch rate but its effects are inconclusive. Saeed et al. (2018) showed that *Wolbachia* maintained a high egg hatch rate in *D. suzukii* even at increased temperature while in Cattel et al. (2018) infected flies had a lower hatch rate than uninfected flies.

Saeed et al. (2018) concluded that *Wolbachia* presence does not offer any fitness benefit linked to nutritional tolerance. However, their study mainly focused on survival of eggs and subsequent development time on a yeast-deprived diet, which may have been insufficient to detect any *Wolbachia*-associated benefit. Our results suggest that the influence of *Wolbachia* presence may differ depending on the diet history (i.e., the maintenance of a stable diet or occurrence of a diet shift). The increases we observed in offspring survival also provide support for the hypothesis that *Wolbachia* plays a mutualistic role for *D. suzukii*. Increasing host survival certainly is beneficial for the bacteria since this effect likely contributes to its higher prevalence in the population (Newton and Rice 2020).

The mechanisms at play in this nutritional mutualism were not addressed directly in this study. As in other insect-Wolbachia nutritional mutualisms (Nikoh et al. 2014), Wolbachia may produce and supply nutrients that contribute to D. suzukii fitness (Newton and Rice 2020). Indeed, studies of other insect species have confirmed that Wolbachia genomes can possess pathways for B vitamins, either complete (e.g., riboflavin B2) or partial (e.g., thiamine B1, pyridoxine B6, biotin B7, and folate B9) (Nikoh et al. 2014), as well as pathways for nucleotide and heme syntheses (Newton and Rice 2020). For example, Wolbachia supplements biotin and riboflavin to bedbugs (Moriyama et al. 2015; Balvín et al. 2018) and planthoppers (Ju et al. 2020). The bacteria also influence iron metabolism in the parasitoid wasp Asobara tabida, the mosquito Aedes aegypti and the flies Drosophila simulans (Kremer et al. 2009) and D. melanogaster (Brownlie et al. 2009). Supplementation seems of particular importance for insect species that feed on single diets (e.g., blood, plant-sap, and grain), as some important nutrients (mostly amino acids and vitamins) are missing from these diets (Ju et al. 2020). Similar effects may also be useful for polyphagous insects developing using diets of varying quality. In our diets, riboflavin and tyrosine differed quantitatively, even if we cannot assess statistical significance. If wSuz have a role in riboflavin and tyrosine metabolism, we would have expected W- flies to struggle in Grape diet, resulting in a lower survival or mass. It was not the case (Fig. 2, 3), either because wSuz does not play a role in riboflavin or tyrosine metabolism, or because our diet was not lacking these nutrients and just had a reduced amount, limiting the potential beneficial impact of Wolbachia. The diets were suboptimal in their P:C ratio for D. suzukii (Silva-Soares et al. 2017; Young et al. 2018) and differed mostly in starch content (Table S1). The role of starch reserves, which likely determine the energetic status of flies, in this nutritional mutualism requires further investigation to identify the mechanisms involved.

The possible contributions of other bacteria, such as *Acetobacter*, to the observed differences also cannot at present be completely excluded. Acetic acid bacteria are important components of the microbiota associated with *Drosophila* (Winans et al. 2017; McMullen et al. 2021), and a role in *D. suzukii* gut response to diet

modification has been suggested (Vacchini et al. 2016). Also, *Wolbachia* presence can itself modulate the relative titers of these bacteria, although the mechanisms involved are not known (Simhadri et al. 2017). We compared the microbiota of W– flies and W– flies with restored microbiota after being in contact with faeces of W+ flies, but some bacteria species may not be transferred by this mechanism and future research should explore these aspects. It is also known that cultivating insects for numerous generations on the same diet stabilizes the gut-microbiome (Sugio et al. 2015; Bing et al. 2018). Overall, we assume that the differences observed in this study are most likely directly linked with *Wolbachia* presence rather than other bacteria such as *Acetobacter*.

Finally, our data indicate that experiencing stressful conditions can increase the beneficial effects of *Wolbachia* presence. In the current study, the stress was induced by the diet shift, through exposure to a new nutritional environment. In other studies on the effect of *Wolbachia* in a nutritional context, benefits are seen in nutrient-deprived situations. For example, *Wolbachia* presence increased *D. melanogaster* fecundity under iron-restricted diets (Brownlie et al. 2009). Similarly, *w*Cle conferred fitness advantages to the host *Cimex lectularius* when fed on B-vitamin-deprived food (Nikoh et al. 2014). Considering other microbes, Bing et al. (2018) showed that the general microbiota of *D. suzukii* is advantageous in the context of a nutrient-poor diet, while having no influence or even deleterious effects when the fly was cultivated on a nutrient-rich diet. Thermal stress is also known to have differential effects on *Wolbachia* and the host insect (Strunov et al. 2013; Saeed et al. 2018). As *Wolbachia* can be both mutualistic and parasitic with the same host (Zug and Hammerstein 2015), studies of its effects should include stressful conditions that can impact the entire holobiont.

Diet affects reproductive traits

As expected, fecundity was affected by the laying diet treatment applied. However, the direction of the effect observed was unexpected. When the diet was stable, the maximum number of laid eggs and surviving offspring were observed on Corn rather than Grape, with the latter containing grape powder, a fruit that *D. suzukii* can use for its development (e.g., Shu et al. 2022). *Drosophila suzukii* is known to favor a low protein diet on which to oviposit (Silva-Soares et al. 2017; Young et al. 2018). This is related to the particular behavior of *D. suzukii* of ovipositing in ripening and ripe fruit, stages at which the fruit reach their maximum sugar content (Burrack et al. 2013) and lowest protein content (Silva-Soares et al. 2017). Sugars subsequently decrease and proteins increase as the fruit starts decaying (Silva-Soares et al. 2017). The overall protein content of our two diets did not differ. Both diets had anti-microbial agent (Table S1), which could have led to different microbial proliferation and divergence in microbiota between Corn and Grape, but it did not seem to influence our results. However, as with proteins, the sugar content influences the number of eggs laid (Burrack et al. 2013). Therefore, females may have relied on sugar concentrations to drive their reproductive behavior in our experiments, thereby preferring to oviposit on the Corn diet, which had lower sugars content even if it had the same P:C ratio of 1:10.

Another factor which may have influenced oviposition is the penetration force required by the diet. Burrack et al. (2013) found that the optimal force required by *D. suzukii* is around 30 cN, which corresponds to the Grape diet. Our direct observations of oviposition suggest that the penetration force required on the Corn diet may have been inferior and therefore suboptimal, even though the females laid more eggs on Corn. Conversely, Silva-Soaeres et al. (2017) reported no difference in the proportions of eggs laid by females on diet patches differing in agar concentration. These observations might explain why, in our experiment, the difference observed between Corn and Grape diets was small, about 40 eggs and 19 offspring. In their study of *D. suzukii*, Olazcuaga et al. (2021) described positive fitness changes across generations when a population was exposed to a new diet after 12 generations and more. As our populations were reared on Grape for about ten generations, it is therefore possible that the differences we observed between Grape and Corn could disappear completely over a longer period.

Notably, diet shift had a strong directional effect on fecundity. Females originating from Grape that were forced to lay eggs on Corn laid the minimum amount of eggs while, in the inverse sequence, females originating from Corn and forced to lay eggs on Grape laid the maximum amount of eggs. In the short term, the Grape diet possible allowed better performance of *D. suzukii* than the Corn diet. In the current study we only tested two artificial diets, but Olazcuaga et al. (2021) used eight fruit and obtained similar results: depending on the diet on which the population were reared, fitness would increase or decrease depending on the new diet, but this effect would disappear after >12 generations on the same diet. In the natural environment, this feature has clear relevance to the multitrophic context in which *D. suzukii* develops. Depending on the context (i.e., availability of different food sources), the species' high polyphagy could either be an advantage or a disadvantage. For instance, after several generations laying eggs on the same fruit (e.g., during a fruit harvest season), females would have to find new fruit to oviposit on, either increasing or decreasing their abundance. This is a general pattern in polyphagous insect pests that depend on dynamic cropping systems (Kennedy and Storer 2000). It would also be of importance in temperate climates when fruit resources are scarce during winter and alternative resources provide nutritional support (Stockton et al. 2019).

Offspring survival was not affected by any diet treatment. The slight differences in diet composition may not have been sufficient to induce a difference in larval mortality (see Young et al. 2018). However, the diet treatment did affect surviving offspring mass and development time. When the diet was stable, offspring mass did not differ between Corn and Grape, though the development time in Corn was longer than in Grape. However, the differences in development time were small and may not have been sufficient to induce differences in offspring mass. These findings are in agreement with Silva-Soares et al. (2017), who found a faster development and larger offspring when *D. suzukii* was raised in intermediate protein concentrations, and also with Jaramillo et al. (2015), who reported faster development on a low sugar diet (blueberry) than on a sugar-rich diet (corn-molasses based) and no difference on phenotypic traits (e.g., wing length). Our results differ however from those of Young et al. (2018) who observed longer egg-to-adult development time and/or smaller adult mass when flies developed in high sugar environments. One possible explanation relies on egg density, which can lead to overcrowding. Overcrowding generally induces longer development time and smaller adult size (Da Silva et al. 2019) and can promote cannibalism in some insects (Richardson et al. 2010),

including *D. suzukii* (Da Silva et al. 2019). We did not control for egg density, but the replacement of the diet every two days contributed to avoid reaching densities that can be detrimental. As seen in real fruit (Da Silva et al. 2019), when larval density is higher, thanks to the large quantity of resource in the experimental Petri dishes larvae pupated around the Petri dish edges in order to avoid competition.

When a diet shift was enforced the development time was shorter on Corn and offspring were lighter, while on Grape the development time was longer and offspring were heavier. Fellous and Xuéreb (2017) found the same pattern when comparing the larval performance of D. suzukii on diets differing in their protein / sugar concentrations, with the shortest larval development time on high protein, low sugar diets. On the other hand, Shu et al. (2022) found larval development time to be shorter and adult mass to be greater on high protein, low sugar diets. However, the latter study took place on actual fruits, which might have impacted larval development differently to artificial diets, especially because of the absence of antimicrobial agents (Young et al. 2018). Generally, it might be expected that, the longer the development time, the heavier the offspring will be as they can benefit from the diet over a longer period. However, this is not always the case as insects are also able to increase their foraging rate, becoming heavier in a shorter time (Da Silva et al. 2019). Faster development without effects on body size can also indicate a higher quality diet (Edgar 2006). In this study, the diet shift induced modifications in both development time and in offspring mass and it is plausible that the diet shift affected both the foraging rate of the larvae and their nutrient intake. Even if both Corn and Grape diets had the same P:C ratio and mainly differed in starch content, the slight differences in nutrients between them may have induced the same consequences observed in studies with diet differing in their P:C ratio. The role and even importance of starch in the performance of *Drosophila* flies remains undetermined and future studies should investigate its role in modulating fitness during diet changes.

CONCLUSIONS

In our study, *Wolbachia* presence improved the survival of *D. suzukii* in the earlier stages of development. The beneficial effects on offspring performance were particularly clear when females experienced a diet shift. Our results unveil an unappreciated role of *Wolbachia* in polyphagous insects: the bacteria support survival of the fly and this effect is especially important during a diet shift. These findings suggest that *Wolbachia* could supplement the diet of its host when it is developing on a low-quality resource, but this hypothesis requires further investigation. Such a role could also explain the maintenance of the *w*Suz strain in *D. suzukii* globally. Depending on specific host fruit availability and quality, *Wolbachia* presence could be more or less advantageous to *D. suzukii*. *Wolbachia* presence is also associated with cost that can offset the benefits, such as costs related to the energy needed to host the endosymbiont (Clavé et al. 2022). Further studies are required to investigate how the cost-benefice balance is modulated in a multitrophic context, depending on the host plants. Abiotic variables (e.g., temperature, humidity) also need to be taken into account as it has been shown that the effect of *Wolbachia* can vary with temperature (Saeed et al. 2018). An integrative approach is needed to properly understand the role *Wolbachia* plays in polyphagous insects. If *Wolbachia* has a mutualistic role, its presence would benefit the fly during invasion, helping with establishment and maintenance of populations

472	in heterogeneous environments. A better understanding of these dynamics is crucial to identify and apply
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493	
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