- 1 Symbiont-mediated fly survival is independent of defensive symbiont
- 2 genotype in the *Drosophila melanogaster-Spiroplasma*-wasp interaction
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13 Abstract

When a parasite attacks an insect, the outcome is commonly modulated by the presence of 14 defensive heritable symbionts residing within the insect host. Previous studies noted markedly 15 different strengths of Spiroplasma-mediated fly survival following attack by the same strain of 16 wasp. One difference between the two studies was the strain of *Spiroplasma* used. We therefore 17 performed a common garden laboratory experiment to assess whether *Spiroplasma*-mediated 18 19 protection depends upon the strain of *Spiroplasma*. We perform this analysis using the two 20 strains of male-killing *Spiroplasma* used previously, and examined response to challenge by two 21 strains of *Leptopilina boulardi* and two strains of *Leptopilina heterotoma* wasp. We found no 22 evidence Spiroplasma strain affected fly survival following wasp attack. In contrast, analysis of the overall level of protection, including the fecundity of survivors of wasp attack, did indicate 23 24 the two *Spiroplasma* strains tested varied in protective efficiency against three of the four wasp strains tested. These data highlight the sensitivity of symbiont-mediated protection phenotypes 25 to laboratory conditions, and the importance of common garden comparison. Our results also 26 indicate that Spiroplasma strains can vary in protective capacity in Drosophila, but these 27 differences may exist in the relative performance of survivors of wasp attack, rather than in 28 29 survival of attack per se.

31 Introduction

The outcome of natural enemy attack has traditionally been considered a function of factors 32 encoded within the genome of the host and infecting parasite. Within this interaction may exist 33 34 a degree of specificity whereby a subset of parasite genotypes are able to infect a subset of host genotypes and, reciprocally, a subset of host genotypes are able to resist a subset of parasite 35 genotypes (Woolhouse et al. 2002; Lambrechts et al. 2006). Specificity between host and 36 parasite genotypes can lead to negative-frequency dependent selection between players and 37 can contribute to the maintenance of heritable variation for defence and attack factors within a 38 39 population (Woolhouse *et al.* 2002; Schmid-Hempel & Ebert 2003).

40 More recently it has been observed that the outcome of natural enemy attack is not solely 41 determined by host and parasite genotypes, but also by the presence and genotype of defensive heritable microbial symbionts residing within the host (Brownlie & Johnson 2009; Oliver et al. 42 2009; Ballinger & Perlman 2019). Defensive microbial symbionts have been identified in a wide 43 range of organisms. For example, microbial symbionts are known to provide protection against 44 45 ssRNA viruses (Hedges et al. 2008; Teixeira et al. 2008), nematodes (Jaenike et al. 2010), fungal 46 pathogens (Scarborough et al. 2005; Lukasik et al. 2013) and parasitic wasps (Oliver et al. 2003; Xie *et al.* 2010). 47

Recently, studies have described how microbial strain identity can complement host and 48 49 parasite genotype as an additional driver of the outcome of a host – parasite interaction. In 50 aphid systems, this is commonly manifested in symbiont strain x host strain x enemy strain 51 interaction terms (Sandrock et al. 2010; Schmid et al. 2012; Cayetano & Vorburger 2013, 2015; Parker *et al.* 2017). Beyond the aphid systems, it is known that the strain of infecting *Wolbachia* 52 is an important source of variation in Wolbachia-mediated protection against viruses in 53 Drosophila, associated with different titre achieved by the strains (Osborne et al. 2009; Bian et 54 al. 2013; Chrostek et al. 2013, 2014; Martinez et al. 2017). Similarly, in the bumblebee, Bombus 55 terrestris, the defensive gut microbiota type is predominantly responsible for resistant 56

phenotypes against the virulent gut trypanosomatid, *Crithidia bombi* (Koch & Schmid-Hempel
2012).

59 The heritable endosymbiont *Spiroplasma*, has been shown to protect *Drosophila* from attack by 60 nematodes and parasitoid wasps (Jaenike et al. 2010; Xie et al. 2010, 2014; Mateos et al. 2016). 61 The ability of Spiroplasma to protect Drosophila is thought to be orchestrated through a combination of RIP toxin activity (secreted by *Spiroplasma*) and exploitative competition 62 between *Spiroplasma* and the infecting parasite for lipid stores (Paredes *et al.* 2016; Ballinger & 63 64 Perlman 2017). Despite being regarded as an important model system, little is known about the role of host, symbiont and parasite identity in determining the outcome of the interaction. 65 Recent work has revealed that the genotype of attacking parasitoid wasp is important for the 66 67 degree of protection conferred by Spiroplasma (Jones & Hurst 2020). It was observed that Spiroplasma (MSRO-Br strain) conferred protection of 40% against the Lh-Fr and Lh-Mad L. 68 *heterotoma* wasp strains, contrasting with 5% protection against the Lh14 strain. The reasons 69 70 underpinning the variation observed is unknown, but intraspecific differences in the toxicity of wasp venom transferred along with the wasp egg during parasitization may be a factor. 71

72 A more general understanding of how symbiont and parasite genotypes are likely to interact is 73 essential for predicting the dynamics of symbionts in natural populations. In this study, we 74 determine whether parasite genotype x symbiont genotype interactions exist within the Spiroplasma-Drosophila melanogaster system. Most studies concerning Spiroplasma-mediated 75 76 protection have reported the outcome of experiment in which a single symbiont strain defends 77 against a single enemy strain. Analysis across these studies indicates that the strain of 78 *Spiroplasma* may be an important component of *Spiroplasma*-mediated protection. For instance, survival of flies exposed to the Lb17 strain of the specialist parasitoid wasp L. boulardi was 79 80 recorded at 5% in *D. melanogaster* infected with the MSRO-Br strain (Xie *et al.* 2014), and 50% in D. melanogaster infected with the MSRO-Ug Spiroplasma strain (Paredes et al. 2016). One 81 interpretation of these results is that the *Spiroplasma* strains differ in protective capacity in *D*. 82

melanogaster. However, analysis of these two strains within a common experimental design
(controlling for potential lab practice, wasp strain and fly strain differences) is required to
determine the precise importance of symbiont strain in determining the outcome of the
parasite-host interaction.

We here present an analysis of the capacity of MSRO-Br and MSRO-Ug to defend *D. melanogaster*against wasp attack. This analysis is performed for two strains of the specialist parasitoid *L. boulardi*, and two strains of the generalist *L. heterotoma*. We compare survival following wasp
attack, mirroring previous studies, and additionally estimate overall protection combining fly
survival data with data on the fertility of flies that survived wasp attack to establish a protective
index for each wasp strain by *Spiroplasma* strain combination.

94 Materials and methods

95 Strains and maintenance

96 Two strains of *Spiroplasma* were used in this study. The first, Red 42, was originally collected in Campinas, São Paulo State, Brazil in 1997 (Montenegro et al. 2000) and later transinfected and 97 98 maintained in the laboratory on a Canton-S background. The second *Spiroplasma* strain was collected from Namulonge, Uganda in 2005 (Pool et al. 2006) which was later transferred and 99 maintained in the laboratory on an Oregon-R background. It should be noted that all larvae from 100 the Spiroplasma infected treatments are female due to the high efficiency of male-killing. 101 However, there does not appear to be any differences in survival between the sexes against 102 parasitoid wasp attack (Xie et al. 2014). All flies were maintained on ASG corn meal agar vials 103 (10 g agarose, 85 g sugar, 60 g maize meal, 40 g autolysed yeast in a total volume of 1 L, to 104 which 25 mL 10% Nipagin was added) at 25 °C on a 12:12 light:dark cycle. 105

The L. boulardi strains used were the NSRef strain, established from an initial female collected in 106 107 Gotheron, near Valence, France (Varaldi et al. 2003), and the Lb17 strain, initially collected in 108 Winters, California in 2002 (Schlenke et al. 2007). The L. heterotoma strains used were the inbred Lh14 strain also collected in Winters, California in 2002 (Schlenke et al. 2007) and the 109 Lh-Mad strain established from a single female collected in Madeira, Portugal in March 2017 110 (Jones & Hurst 2020). The wasp stocks were all maintained on second instar Oregon-R larvae at 111 25°C on a 12:12 light:dark cycle. After emergence, wasps were maintained on grape agar vials 112 supplemented with a flug moistened with honey water and allowed to mature and mate for 7 113 days prior to exposure to *D. melanogaster* L2 larvae. 114

115 Artificial infection of *Spiroplasma*

The *Spiroplasma* strains (MSRO-Br and MSRO-Ug) were artificially transferred into a common host background (Canton-S) to remove any effect of host nuclear background on the level of protection conferred. Canton-S stocks carry the naturally occurring *Wolbachia* strain wMel. *Wolbachia* has been shown to provide a weak positive effect on fly larva-to-adult survival and a negative effect on wasp success in flies attacked against *L. heterotoma* (Lh14 strain) (Xie *et al.*2014). Artificial infections were carried out as described by Nakayama *et al.* (2015).
Hemolymph was extracted from the thorax of *Spiroplasma*-infected *D. melanogaster* and mixed
with sterile PBS. Virgin female Canton-S were artificially injected in the abdomen with 0.1-0.2 μl
of PBS-hemolymph, using a hydraulic positive-pressure microinjection apparatus (Model IM-6,
Narushige Ltd, Tokyo, Japan).

126 Confirmation of *Spiroplasma* infection status

127 Three weeks post injection, the infection status of the artificially infected flies was confirmed via Spiroplasma-specific PCR. DNA extraction was carried out using the Wizard® Genomic DNA 128 129 Purification Kit (Promega). To this end, each injected mother was taken and macerated in 150 μl of Nuclei Lysis Solution and incubated at 65 °C for 30 min. After incubation, 50 μl of Protein 130 131 Precipitation Solution was added to each sample and then placed on ice for 5 min. Samples were then centrifuged for a further 4 min at 16,000 x g and the supernatant was transferred into a 132 new tube containing 150 µl of isopropanol. Samples were centrifuged for 2 min at 16,000 x g 133 134 and the supernatant discarded. 150 µl of 70% ethanol was added to each sample and 135 centrifuged for 1 min at 16,000 x g. The supernatant was discarded. Pellets were dried before re-suspending in 25 µl of molecular grade water at 4 °C overnight before use in subsequent PCR 136 assays. 137

138 PCR amplifications were conducted using *Spiroplasma* specific primers, SpoulF (5'-GCT TAA CTC CAG TTC GCC-3') and SpoulR (5'-CCT GTC TCA ATG TTA ACC TC-3') (Montenegro et al. 2005). 139 Each reaction was carried out in 15 µl volume containing 7.5 µl of GoTaq® Hot Start Green 140 Master Mix (Promega), 0.5 µl each of the forward and reverse primer, 5.5 µl of Molecular Grade 141 Water and 1 µl of DNA. All reactions were conducted alongside the positive and negative 142 143 controls. This included a PCR negative control containing the PCR reaction mixture only 144 (excluding DNA template). The PCR thermal program consisted of an initial denature of 5 min at 95 °C, followed by 35 cycles of 15 s at 94 °C, 1 min at 55 °C and 40 s at 72 °C. The PCR products 145

were electrophoresed in a 1.5% agarose gel at 155 V for 15 min and the products were
visualised. Offspring sex ratio of infected mothers were also checked to determine *Spiroplasma*efficiency. Only mothers which were infected with *Spiroplasma* and produced all female broods
were used to create new lines.

150 To confirm the Spiroplasma strain status of each artificially injected line of Drosophila *melanogaster*, sequencing was performed on 5 individual flies from each strain. To this end, the 151 DNA of 5 flies from each Spiroplasma strain line were extracted using the Wizard® Genomic 152 DNA Purification Kit following the methodology from above. PCR amplifications were conducted 153 154 using Spiroplasma specific primers, Spiro_MSRO_diff_F (5'-TAC GAC CAA TGG CTT GTC CC-3' and Spiro MSRO diff R (5'- CTG GCA TTG CTT TTT CCC CA-3'). The PCR thermal program consisted 155 156 of an initial denature of 2 min at 94 °C, followed by 35 cycles of 15 s at 94 °C, 1 min at 56 °C and 157 40 s at 72 °C. To prepare the PCR reaction for sequencing, PCR products underwent an ExoSAP digest clean up to remove excess primers. To this end, 5 μ l of PCR product was added to a 158 159 mixture containing 0.2 µl Shrimp alkaline phosphate, 0.05 µl of Exonuclease I, 0.7 µl 10X RX 160 Buffer and 1.05 µl of molecular grade water. Samples were then incubated for 45 min at 37 °C followed by 15 min at 80 °C and sent for Sanger sequencing. The *Spiroplasma* strain status of the 161 MSRO-Br and MSRO-Ug line were confirmed by the presence of a Guanine and Thymine 162 respectively in position 414193, coding for a type III pantothenate kinase. The expected 163 164 amplicon size is 509bp. Transinfected fly lines were passaged for >10 generations before 165 experiments were conducted.

166 Wasp attack assay

To ensure efficient vertical transmission of *Spiroplasma*, infected females were aged to at least ten days prior to egg laying. Flies were allowed to mate in cages and lay eggs on a grape Petri dish painted with live yeast for 24 h. Grape Petri dishes were incubated for a further 24 h to allow larvae to hatch. First instar larvae were picked from the grape plate into the experimental vials at 30 larvae per vial. A fully factorial design was created for each of the

four wasp strains described which included *Spiroplasma* strain (MSRO-Br, MSRO-Ug and uninfected control) and wasp (presence or absence). Five experienced, mated female wasps were transferred into the wasp treatment vials. Adult wasps were allowed to parasitise for 2 days before being removed. All vials were maintained at 25 °C on a 12:12 light:dark cycle. For each vial, the number of puparia, emerging flies and emerging wasps were recorded. Experiments using *L. boulardi* and *L. heterotoma* were conducted in separate blocks, one week apart.

179 Measuring female fecundity

Spiroplasma infected flies that survive wasp attack generally have a lower fecundity than 180 Spiroplasma infected flies which were not exposed to wasps (Xie et al. 2011; Jones & Hurst, 181 2020). To determine whether the wasp attacked survivors were differentially impacted by 182 Spiroplasma strain the average daily emerged offspring of Spiroplasma infected survivors 183 ("Exposed") and *Spiroplasma* infected flies which did not undergo wasp attack ("Unexposed") 184 was measured for the MSRO-Br and MSRO-Ug strains. The Spiroplasma uninfected wasp 185 186 attacked group was not included due to the extremely low number of flies which emerged, 187 which were also likely to have avoided wasp attack all together. After emergence, flies from the 188 wasp attack assay were stored in vials containing sugar yeast medium (20 g agarose, 100 g 189 sugar, 100 g autolysed yeast in a total volume of 1 L, to which 30 mL 10% Nipagin w/v propionic acid was added) at mixed ages. A week after emergence commenced, a subset of flies 190 191 from each of the *Spiroplasma* treatments were placed into an ASG vial with two Canton-S males with a single yeast ball and allowed to mate. Approximately 25 replicates per treatment were 192 created. Flies were transferred onto fresh ASG vials each day for five days. Flies were given two 193 194 weeks to emerge to ensure every fly had emerged before counting. Female fecundity was 195 measured as the average number of offspring produced over four days (day 2-5).

196 Statistical analysis

All statistical analyses were performed using the statistical software R, version 3.5.0 (R CoreTeam 2018). Fly and wasp survival data were analysed by fitting a generalized linear model

with binomial distributions. In all cases, a fully saturated model including all factors and their
interaction was reduced to a minimum adequate model through step-wise simplification. Nonsignificant factors are reported as the output of the model comparisons. The effect of significant
independent variables are reported from the analysis of the minimum adequate model using the
'car' package.

204 To produce a composite measure of protection, a Protective Index (PI) was calculated by comparing the survival and fecundity of *Spiroplasma*-infected flies in the presence/absence of a 205 206 given strain of wasp. The PI was calculated as the ratio of p(survival) x p(fertile) x fecundity of 207 fertile individuals for attacked vs unattacked Spiroplasma-infected flies and reflects the benefit 208 of Spiroplasma in the face of wasp attack. Credible intervals for PI were calculated through 209 simulation. By assuming prior probability distributions for each parameter (Survival probability = beta distribution; Fertility probability = beta distribution; Fecundity = normal distribution). 210 the 'rbeta' and 'rnorm' functions were used to calculate 95% credible intervals for PI. The 211 212 simulation data was also used to establish the posterior probability of PI differing between attacking wasp strains. 213

215 Results

- 216 Fly survival and wasp success
- 217 Leptopilina boulardi experiment
- 218 In the absence of *L. boulardi* wasps, *Spiroplasma* strain had a significant effect on fly larva-to-

adult *D. melanogaster* survival ($\chi^2 = 7.74$, d.f. = 1, *P* = 0.005). The mean survival of MSRO-Br

infected and MSRO-Ug infected *D. melanogaster* was 72.2% and 83%, respectively (Fig. 1A). In

221 the presence of *L. boulardi* wasps, there was no significant effect of wasp strain ($\chi^2 = 0.281$, d.f. =

1, P = 0.596), *Spiroplasma* strain ($\chi^2 = 0.0008$, d.f. = 1, P = 0.977), nor a significant interaction

between wasp strain and *Spiroplasma* strain on larva-to-adult survival of *D. melanogaster* (χ^2 =

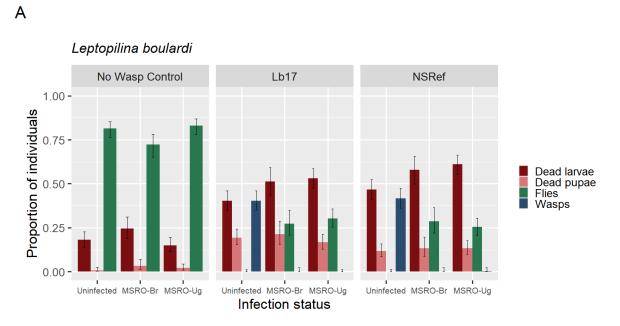
224 0.284, d.f. = 1, P = 0.594) (Fig. 1A). There was no significant effect of wasp strain on wasp

success (χ^2 = 0.121, d.f. = 1, *P* = 0.728) (Fig. 1A), and wasps were observed only in the absence of

226 Spiroplasma.

227 Leptopilina heterotoma experiment

228 In the absence of *L. heterotoma* wasps, there was no significant effect of *Spiroplasma* strain on fly larva-to-adult survival (χ^2 = 0.345, d.f. = 1, *P* = 0.557). The mean survival of uninfected, 229 MSRO-Br infected and MSRO-Ug infected D. melanogaster was 81.1% and 83%, respectively 230 (Fig. 1B). In the presence of *L. heterotoma* wasps, there was a significant effect of wasp strain on 231 fly larva-to-adult survival of *D. melanogaster* (χ^2 = 34.21, d.f. = 1, *P* < 0.001). Fly larva-to-adult 232 survival of *Spiroplasma*-infected *D. melanogaster* attacked by the Lh-Mad strain of *L. heterotoma* 233 was approximately double that observed for flies attacked by the Lh14 strain of L. heterotoma 234 (Fig. 1B). Similar to the *L. boulardi* experiment, there was no significant effect of *Spiroplasma* 235 strain ($\chi^2 = 0.740$, d.f. = 1, P = 0.390), nor a significant interaction between wasp strain and 236 237 *Spiroplasma* strain (χ^2 = 0.674, d.f. = 1, *P* = 0.412) on larva-to-adult survival of *D. melanogaster* (Fig. 1B). There was a significant effect of wasp strain on wasp success ($\chi^2 = 4.805$, d.f. = 1, P =238 239 0.028) (Fig. 1B). The average wasp success of the Lh14 and Lh-Mad wasp strains were 27.7% 240 and 37.3% respectively. Wasps only emerged in the absence of *Spiroplasma*, with both symbiont 241 strains preventing development of both wasp strains.



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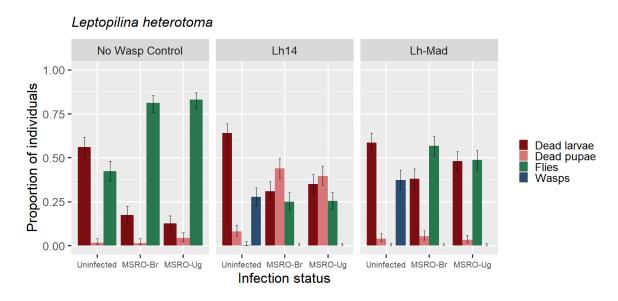


Figure 1: Proportion of dead larvae (red), dead pupae (pink), emerging flies (green) and
emerging wasps (blue) for *Spiroplasma*-infected (MSRO-Br and MSRO-Ug strains) and
uninfected *Drosophila melanogaster* attacked by A) *L. boulardi* (Lb17 and NSRef strains) and B) *L. heterotoma* (Lh14 strain and Lh-Mad strains). Error bars represent 95% binomial confidence
intervals.

248 Overall protection index

Despite finding no difference between the survival of flies infected with MSRO-Br and MSRO-Ug 249 250 against each of the four wasp strains tested, previous work has shown that it is also important 251 to consider, in combination with survival, the fertility of wasp-attacked flies compared to non-252 attacked controls to produce a complete model of protection (Xie et al., 2011; Jones & Hurst 253 2020). Taking into account the survival, proportion of adults fertile, and the fecundity of wasp attack survivors, compared to unexposed *Spiroplasma*-infected controls, a protection index (PI) 254 was calculated as the product of fly survival x p(fertile) x fecundity of exposed vs unexposed 255 256 *Spiroplasma*-infected flies (Table 1). This metric assumes complete mortality from wasps in the 257 absence of *Spiroplasma*, which is approximately true as <1% of individuals tested survived wasp 258 attack. Against the Lb17, NSRef and Lh-Mad strains of wasp, the posterior probability that the protection index for MSRO-Br is greater than the protection index for MSRO-Ug is >0.97 (Table 259 260 2). However, against the Lh14 strain of wasp, the posterior probability that the protection index 261 for MSRO-Br is greater than the protection index for MSRO-Ug is 0.44 (Table 2).

Table 1: The overall protection conferred by MSRO-Br and MSRO-Ug *Spiroplasma* strains against a) *Leptopilina boulardi* (Lb17 and NSRef strains)

and b) Leptopilina heterotoma (Lh14 and Lh-Mad strains) in Drosophila melanogaster. Exposed S- = wasp attacked Spiroplasma-uninfected flies;

264 Exposed S+ wasp attacked Spiroplasma-infected flies; Unexposed S+ Spiroplasma-infected flies not attacked. Protective Index is calculated as

265 [p(survival) x p(fertile) x fecundity of fertile individuals] of exposed vs unexposed individuals with credible intervals calculated as given in methods.

a) Leptopilina boulardi

Spiroplasma		Fly Survival (binomial 95% CI intervals	Proportion fertile (binomial 95%	Fecundity measure ±	Estimated protective index (95%			
strain	Treatment	(lower, upper))	CI intervals (lower, upper))	SE	Credible interval (lower, upper))			
MSRO-Br	Lb17 exposed S+	0.27 (0.20, 0.35)	0.96 (0.75, 0.99)	15.8 ± 1.31	0.37 (0.25, 0.55)			
	NSRef exposed S+	0.29 (0.22, 0.36)	1.00 (0.86, 1.00)	15.6 ± 1.45	0.40 (0.27, 0.59)			
	Unexposed control S+	0.72 (0.65, 0.78)	0.92 (0.72, 0.98)	16.9 ± 1.47				
MSRO-Ug	Lb17 exposed S+	0.30 (0.25, 0.36)	1.0 (0.85, 1.0)	11.0 ± 1.77	0.30 (0.14, 0.32)			
	NSRef exposed S+	0.25 (0.21, 0.30)	0.88 (0.68, 0.96)	14.3 ± 1.58	0.20 (0.14, 0.30)			
	Unexposed control S+	0.83 (0.78, 0.87)	0.96 (0.776, 0.99)	19.4 ± 1.49				

b) Leptopilina heterotoma

<i>Spiroplasma</i> strain	Treatment	Fly Survival (binomial 95% CI intervals (lower, upper))	Proportion fertile (binomial 95% CI intervals (lower, upper))	Fecundity measure ± SE	Estimated protective index (95% Credible interval (lower, upper))
MSRO-Br	Lh14 exposed S+ Lh-Mad exposed S+ Unexposed control S+	0.25 (0.20, 0.30) 0.57 (0.51, 0.62) 0.81 (0.76, 0.85)	0.79 (0.59, 0.91) 0.91 (0.71, 0.98) 0.92 (0.71, 0.98)	$13.2 \pm 1.75 \\ 14.0 \pm 1.58 \\ 14.3 \pm 1.72$	0.24 (0.15, 0.39) 0.68 (0.54, 1.16)
MSRO-Ug	Lh14 exposed S+ Lh-Mad exposed S+ Unexposed control S+	0.25 (0.20, 0.31) 0.49 (0.43, 0.54) 0.83 (0.78, 0.87)	1.0 (0.82, 1) 0.91 (0.70, 0.98) 1.0 (0.82, 1)	$12.8 \pm 1.14 \\ 11.0 \pm 1.34 \\ 15.3 \pm 1.58$	0.25 (0.18, 0.36) 0.39 (0.26, 0.56)

Table 2: The posterior probability that the estimated protective index for MSRO-Br is greater than the MSRO-Ug for each wasp strain tested.

Wasp strain	Estimated protective ind (lower)	Posterior probability (EPI MSRO-Br > EPI MSRO-Ug)				
	MSRO-Br	MSRO-Ug	_			
Leptopilina boulardi						
Lb17	0.37 (0.25, 0.55)	0.30 (0.14, 0.32)	0.97			
NSRef	0.40 (0.27, 0.59)	0.20 (0.14, 0.30)	0.99			
Leptopilina heterotoma						
Lh14	0.24 (0.15, 0.39)	0.25 (0.18, 0.36)	0.44			
Lh-Mad	0.68 (0.54, 1.16)	0.39 (0.26, 0.56)	0.99			

268 Discussion

Defensive symbionts can contribute to the outcome of a host-parasite interaction. Previous 269 studies in aphids have shown that the strain of symbiont is an important determinant of 270 271 symbiont-mediated protection across multiple model systems (Schmid et al. 2012; Cayetano & Vorburger 2013, 2015; Parker et al. 2017). However, whether strains of the Drosophila 272 273 defensive symbiont, Spiroplasma poulsonii, vary in their capacity for protection is unknown. 274 The contrasting levels of fly survival observed between two previous studies on the 275 Drosophila-Spiroplasma-L. boulardi interaction suggested that the strain of Spiroplasma may be an important determinant of protection capacity in Drosophila (Xie et al. 2014; Paredes et al. 276 2016). We therefore performed an experiment to determine whether the strength of 277 278 Spiroplasma-mediated protection depended on the strain of infecting Spiroplasma using two 279 known strains of MSRO Spiroplasma (MSRO-Br and MSRO-Ug). We found no evidence that the 280 strength of Spiroplasma-mediated fly survival differed between the MSRO-Br and MSRO-Ug 281 strains against any of the four *Leptopilina* wasp strains tested. However, the overall protective index, including the fecundity of survivors of wasp attack, did vary between the two 282 283 *Spiroplasma* strains for three of the attacking wasp strains.

284 The strain of *Spiroplasma* did not alter the strength of *Spiroplasma*-mediated fly survival in *D*. *melanogaster* in our experiment. This result raises the question as to why fly survival following 285 286 attack differed between the two previous independent studies. Fly survival against the 287 parasitoid wasp, L. boulardi (strain Lb17) was observed to vary from 5% with MSRO-Br (Xie et al. 2014), to 50% with MSRO-Ug (Paredes et al. 2016). Comparisons across studies indicate 288 289 that the strength of symbiont-mediated fly survival appear to be highly variable across 290 laboratory studies. In this study, we found survival of 30% against the L. boulardi (Lb17 291 strain), yet Paredes et al. (2016) found survival of 50% against the same wasp strain despite using the same fly strain. Similarly, we found survival of 25% against the Lh14 strain of L. 292 293 heterotoma, despite survival of <8% observed in previous studies (Xie et al. 2014, Jones & 294 Hurst 2020).

295 The variability in *Spiroplasma*-mediated survival observed across studies may be the result of 296 variability in wasp success. Whilst wasp attack rate was very high in all cases (with very low fly 297 survival in uninfected controls), wasp success was highly variable across the studies and correlated to some extent with fly survival. Specifically, against the Lb17 wasp strain, Xie et al. 298 (2014) found high wasp success of \sim 70% and low fly survival of \sim 5%. In contrast, this study 299 observed reduced wasp success of $\sim 40\%$ and increased fly survival of $\sim 30\%$. Thus, the 300 variability in Spiroplasma-mediated fly survival across studies could be associated with 301 302 condition of the attacking wasps. Associated with this, it is notable that larval-to-pupa survival following attack is lower in our studies than previously observed, and this may potentially 303 304 explain differences in wasp survival. These studies may highlight the sensitivity of symbiont-305 mediated protection to husbandry conditions of both fly and wasp.

306 From several studies, it has been demonstrated that symbiont-mediated survival against 307 natural enemies can be highly sensitive to particular environmental conditions. Temperature 308 is one environmental factor known to impact the strength of symbiont-mediated protection 309 (Corbin et al. 2017). For example, in the pea aphid, higher temperatures can negatively impact H. defensa-mediated survival against Aphidius ervi (Doremus et al. 2018). Similarly, heat shock 310 311 also negatively impacts X-type-mediated survival against A. ervi wasps in the pea aphid (Heyworth & Ferrari 2016). Another possibility, raised by studies of the strength of CI and 312 313 male-killing exhibited by Wolbachia, is that protection strength is influenced by parental 314 Spiroplasma titre (Dyer et al. 2005; Layton et al. 2019). It is notable that both thermal 315 environment and age at reproduction are known to affect *S. poulsonii* titre and male-killing 316 strength in D. melanogaster (Anbutsu & Fukatsu 2003; Montenegro & Klaczko 2004; Anbutsu et al. 2008). Finally, wasp husbandry and attack protocols may vary. Wasp attack success is 317 318 thought to be higher when wasps are previously conditioning before assays and may also be 319 impacted by the arena in which attack occurs. Wasps attack fly larvae at the surface of the food, 320 and the surface area available for attack, and indeed the medium in which the larvae are 321 feeding, may impact success. The variable strength of protection afforded by symbionts across

laboratories may be due to unmeasured differences in stock maintenance/ambient
environmental conditions and reinforce the need for common-laboratory experiments when
comparing outcomes.

Our experiment nevertheless did indicate differences in protection associated with Spiroplasma 325 326 strain, but these were reflected in the overall phenotype, including the survival and fecundity of wasp-attack survivors. Surviving flies infected with the MSRO-Br strain of Spiroplasma had an 327 328 overall higher protective index against the NSRef, Lb17 and Lh-Mad strains of wasp compared 329 to flies infected with the MSRO-Ug strain. The reasons as to why fly survivors infected with MSRO-Ug had a lower protective index compared to MSRO-Br survivors remains unclear. One 330 possible factor which cannot be ruled out from this study is the effect of Wolbachia. Although 331 332 from the results it does not appear that *Wolbachia* is having an effect on fly survival, it may be 333 possible that the presence of *Wolbachia* is differentially impacting the fertility of wasp-attacked survivors among the MSRO-Br and MSRO-Ug strains tested. Another factor which is difficult to 334 determine is the possibility that a proportion of flies in the *Spiroplasma* treatments were not 335 attacked. Although fly emergence from the Spiroplasma negative controls suggests that all 336 larvae were successfully parasitized, this does not exclude the possibility that not all larvae 337 338 were parasitized in the *Spiroplasma* positive treatments, although past work found no evidence for discrimination by wasps (Xie et al. 2010, Jones & Hurst, 2020). However, the result that 339 340 there was no difference in the overall protection between wasp-attacked survivors infected 341 with MSRO-Br and MSRO-Ug against the Lh14 strain of wasp indicates that the reasons for this 342 difference may be a consequence of wasp strain.

This study clearly demonstrates two important features of protection. First, there is a need for common-laboratory experiments to compare levels of protection, as this phenotype has both genetic and environmental drivers. Second, there is a clear distinction between symbiontmediated survival and symbiont-mediated protection within defensive symbiont studies. Symbiont-mediated protection is often measured as the relative survival of an infected-

348 individual compared to an uninfected individual when faced with natural enemy attack. However, symbiont-mediated protection is not only the ability of an infected-host to survive, 349 but also the relative capacity it has to reproduce compared to un-attacked comparators. Despite 350 finding no evidence that fly survival differed between the two strains of *Spiroplasma* against all 351 four wasp strains tested, differences between *Spiroplasma* strains were observed on the overall 352 353 strength of symbiont-mediated protection. Assessment of the relative survival and reproductive ability of un-attacked vs. attacked survivors is essential for revealing the true protective 354 capacity of a defensive symbiont. 355

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- 365 Conflict of interest
- 366 The authors declare no conflicts of interest.

367 Data accessibility

368 Data generated and analysed during this study are available at figshare 369 (https://doi.org/10.6084/m9.figshare.c.4856790.v1).

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488 Supplementary material

Table S1: Replicate identity and number.

Experiment	Figure	Replicate identity	Treatment	Number of replicates					
					Wasp strain				
				Lb17	NSRef	Control	Lh14	Lh-Mad	Control
Survival	1	Vial of 30 larvae	Uninfected	10	10	10	10	10	10
			MSRO-Br	5	5	6	10	10	9
			MSRO-Ug	10	10	10	10	10	10
Proportion	2	Single female fly	MSRO-Br	23	24	24	24	23	24
fertile			MSRO-Ug	23	24	24	19	22	19
	_								
Number of	3	Single female fly	MSRO-Br	22	24	22	19	21	22
daughters			MSRO-Ug	23	21	23	19	20	19
produced									