

Evolutionary Ecology

The effect of mating on starvation resistance in natural populations of *Drosophila melanogaster*. --Manuscript Draft--

Manuscript Number:	EVEC1124R2
Full Title:	The effect of mating on starvation resistance in natural populations of <i>Drosophila melanogaster</i> .
Article Type:	Research Article
Keywords:	mating effects; starvation resistance; wild flies; <i>Drosophila melanogaster</i> .
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Manuscript Region of Origin:	
Abstract:	<p>In nature, behavioural and physiological process involved in mating may entail different costs and benefits for males and females. However, it has been hypothesized that sexual interactions may have additional costs for <i>Drosophila</i> females like decrease in receptivity to remating and shortening of lifespan. During mating, males transfer seminal fluid proteins to females that exert severe physiological changes that may compromise female's lifespan and reproductive success. However, under specific stressful environmental conditions that organisms usually face in nature, mating may also confer benefits to females. In the present work, we examine the effect of mating on starvation resistance in wild <i>Drosophila melanogaster</i>. We demonstrate that mated females derived from different geographic locations have the benefit of a greater starvation resistance as compared to virgin females. Even though mating status did not affect mean starvation resistance, we detected a strong genotype-specific effect in males. Beyond the obvious advantage of mating, our study reveals that mating might not be perilous for females, as envisaged by sexual conflict theories, but advantageous for flies exposed to shifts in environmental conditions. Thus, our results highlight the importance of studying other ecologically relevant traits that may contribute to the evolution of male-female interactions.</p>
Response to Reviewers:	<p>We detail the actions taken regarding the comments of the Associate Editor:</p> <p>1. In this study I think the main effects are equally as interesting as the interaction term and they should be interpreted in combination. In the current revision, I think some minor qualification is all that is required in order to avoid the concerns that readers, such as the previous reviewers, might have (mistakenly or not). Following the editor's advice we decided to give more emphasis on the interaction between line and mating status all along the revised version of the manuscript.</p>

2. Line 105: Avoiding the mention of "subtle effects of mating status on males" - the effect is strong and line specific - but over all the lines there was no main effect. Please find the correction in lines 106-108 in the revised version of the manuscript.

3. Line 183: perhaps clarify that "the effect of mating status, over all possible iso-female lines, was significant in females but not in males?" Please find the correction in lines 189 in the revised version of the manuscript.

4. Line 196: perhaps include a sentence along the lines of "Therefore, the impact of mating status appears to be line specific and thus accounts for the lack of an overall effect of mating status in males compared to females". Please find the correction in lines 204-206 in the revised version of the manuscript.

5. Line 233: ? and explains the lack of an overall effect of mating status on male starvation resistance. Please find the correction in lines 236-237 in the revised version of the manuscript.

6. Line 238: thus contributing in part to the increase in variance in the mating treatment for female, right? Please find the correction in lines 242-243 in the revised version of the manuscript.

7. Line 240-242: this is simply not true - at the level of the line mating did have effect - it is just that the overall effect is not significant. You can interpret the main effect, but why completely ignore the interactions. What is true is that unlike for females, it is unclear how variation in the number of mating between males would influence the reaction norms. Please find the correction in lines 243-246 in the revised version of the manuscript.

8. I encourage the authors to try to carefully streamline their discussion. The inclusion of the population differences at the start of the interaction seems out of place compared to the general theme of the paper. The a priori expectations (Line 202-210) are not mentioned in the introduction and seem to be not well supported. A discussion of the population differences is required (as per Reviewers 2 suggestion) - I will leave it to the author's whether they tone down the speculation at the start of the discussion and/or move it to elsewhere in the discussion. I also think that the direction of the discussion is overly focused on females - a large portion of the discussion (Line 267 onwards) is on the transfer of peptides. Some discussion is needed on the difference between males and females and the male line specific effects of mating status (without overly lengthening the discussion). This is mentioned briefly Line 262 - 263 and could be made more prominent as an explanation. Following the editor's advice we decided to give less emphasis to the among population variation and reduced the length of the corresponding paragraph. The discussion of the possible implication of seminal and sex peptides in the effect of the mating status on starvation resistance was reduced (see lines 268-277) and we enlarged the sentences dealing with the comparison between sexes (lines 284-289) and possible implications of the different types of genotype by mating status observed in males and females. Please find the discussion of the male line specific effects of mating status in lines 290-296 in the revised version of the manuscript.

Minor comments:

9. Line 67: this should be increased mating rates must be less advantageous for females, mating per se is essential. Please find the correction in the lines 68 in the revised version of the manuscript.

10. Line 103-105: Perhaps it is worth mentioning specifically that for males the impact of mating was iso-line/genotype specific. Please find the correction in the lines 106-108 in the revised version of the manuscript.

11. Line 165: What was used to analyse the data - JMP, R etc etc. Please find the correction in the lines 168-170 in the revised version of the manuscript.

12. Line 183-188: see more detailed comments above. Please see the correction in the point 3 above.

The effect of mating on starvation resistance in natural populations of

Drosophila melanogaster

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Running title: Effects of mating on starvation resistance in wild females flies.

Keywords: mating effects; starvation resistance; wild flies; *Drosophila melanogaster*.

Total word count: 3968

Word counts for:

Abstract: 199

Introduction: 788

Materials and Methods: 840

Results: 480

Discussion: 1477

Acknowledgements: 82

Number of cited references: 66

Number of tables and figures: 2 and 2

Abstract

In nature, behavioural and physiological process involved in mating may entail different costs and benefits for males and females. However, it has been hypothesized that sexual interactions may have additional costs for *Drosophila* females like decrease in receptivity to remating and shortening of lifespan. During mating, males transfer seminal fluid proteins to females that exert severe physiological changes that may compromise female's lifespan and reproductive success. However, under specific stressful environmental conditions that organisms usually face in nature, mating may also confer benefits to females. In the present work, we examine the effect of mating on starvation resistance in wild *Drosophila melanogaster*. We demonstrate that mated females derived from different geographic locations have the benefit of a greater starvation resistance as compared to virgin females. Even though mating status did not affect mean starvation resistance, we detected a strong genotype-specific effect in males. Beyond the obvious advantage of mating, our study reveals that mating might not be perilous for females, as envisaged by sexual conflict theories, but advantageous for flies exposed to shifts in environmental conditions. Thus, our results highlight the importance of studying other ecologically relevant traits that may contribute to the evolution of male-female interactions.

Introduction

In sexually reproducing organisms, a sexual conflict may arise whenever males and females have different reproductive interests (Partridge and Hurst 1998; Chapman et al. 2003a; Pischedda and Chippindale 2006). The importance of such sexual conflict is its potential to drive evolutionary change via sexually antagonistic co-evolution or co-evolutionary arms race between sexes (Arnqvist and Rowe 2002; Pischedda and Chippindale 2006). Such a scenario may produce adaptations in one sex that could be eventually harmful in the other, leading to subsequent counter-adaptations in the latter sex to alleviate the costs imposed by such harmful adaptations. This is the main hypothesis of the sexual conflict theory (Holland and Rice 1998; Chapman et al. 2003a). For instance, male seminal fluid proteins (SFPs) transferred to females with sperm during mating are potential targets of such sexual antagonistic selection. SFPs induce physiological and behavioural changes in mated females (Chapman and Partridge 1996; Chapman et al. 2003b; Wolfner 2002; Kubli 2003; Avila et al. 2011), increasing male reproductive success by delaying female re-mating and increasing egg laying rate (Wigby and Chapman 2005; Fricke et al. 2009). However, some of these proteins may have toxic side-consequences in females by reducing lifespan and reproductive success (Chapman et al. 1995; Civetta and Clark 2000; Wigby and Chapman 2005). In addition, females may suffer additional costs resulting from sexual interactions as the increase of predation, the risk of physical damage and parasite/pathogen infections and the decrease in the number of interactions with other males (Partridge and Hurst 1998; Arnqvist and Nilsson 2000; Johnstone and Keller 2000; Cordero and Eberhardt 2005). In this context, it has been hypothesized that increased mating rate may be less advantageous for females than for males. Nevertheless, there is some uncertainty regarding the importance and ubiquity of such conflict given the observation of female ‘resistance’ behavior, the existence of studies that failed to demonstrate the reproductive costs of mating to females (Cordero and Eberhard 2003) and,

the very few examples documenting the phenomenon. In support to this point, it has been shown that females of the Australian sheep blowfly (*Lucilia cuprina*) gain rather than lose when responding to changes in male genitalia (Cordero and Eberhard 2003, 2005). Moreover, changes in genital structures may lead to an increase in the access of male seminal products into female's reproductive tract (Cordero and Eberhard 2003, 2005).

In nature, organisms often face stressful environmental conditions, which are defined as environmental factors that reduce fitness (Koehn and Bayne 1989). The most important sources of natural selection are the common environmental stressors, desiccation and starvation. Organisms deal with periods of starvation and desiccation by deploying behavioural and physiological mechanisms that allow them to alleviate the consequences of environmental stress (Hoffmann and Parsons 1991; Randall et al. 1997). Studies performed in insect groups as Lepidoptera, Orthoptera and Coleoptera, showed that females mate with multiple males during periods of starvation and desiccation, probably because males transfer nuptial gifts containing large amounts of water and nutrients that may improve female stress resistance (Boucher and Huignard 1987; Butlin et al. 1987; Ivy et al. 1999; Edvardsson 2006).

Also, studies in fruit flies of the genus *Drosophila* inhabiting desertic areas have shown that mated females are more resistant to desiccation than unmated females (Knowles et al. 2004, 2005). These physiological responses suggests that male-female interactions may affect ecologically relevant traits and indicates that, in a natural scenario, mating may not be as harmful for females as previously thought (Chapman et al. 1995; Wolfner 1997; Lung et al. 2002; Chapman and Davies 2004).

Usually, flies living in temperate areas are commonly exposed to seasonal variation in food availability. Consequently, the ability of flies to survive during periods of food shortage is an adult fitness component that acquires special relevance during adverse periods (Izquierdo 1991; Mitrovski and Hoffmann 2001; Boulétreau-Merle and Fouillet 2002). Starvation

resistance is a complex trait influenced by many genes and the external and sexual environments (Harbison et al. 2004; Hoffmann et al. 2005a; Goenaga et al. 2010). Moreover, complex mechanisms are induced that result in an increase of tolerance to food deprivation (Rion and Kawecki 2007).

The goal of the present study is to investigate if mating has an effect on starvation resistance in natural populations of *Drosophila melanogaster* uncovering a wide range of environmental conditions. To address this issue, we measured starvation resistance (SR) in mated and virgin flies derived from three sampling localities in Argentina. We demonstrate that mating has a beneficial effect in females as indicated by a greater SR in mated vs virgins in all natural populations tested. Although differences between mated and virgin males in mean survival under starvation conditions were not significant, the effect of mating status on starvation tolerance was genotype (line) specific in males.

Materials and Methods

Drosophila stocks

This work was performed with flies collected in three sampling localities from Argentina: Güemes (24° 41'S, 65° 03'W and 695 m above sea level), Cachi (25° 07'S, 69° 09'W and 2280 m asl) and Lavalle (32° 50'S, 68° 28'W and 647 m asl). Wild flies were collected using fermented banana baits and sorted by sex upon arrival to the laboratory. Wild inseminated females were isolated and used to establish isofemale lines by rearing the offspring of individual females in vials. All isofemale lines were maintained at a census size of approximately 100 flies per line for 10 generations on a cornmeal-molasses-agar (lab medium) medium at constant temperature (25°C ± 1), humidity (60% - 70%) and 12 h light / 12 h dark cycle before the onset of the experiments described below. The number of lines analyzed in each locality varied from 7 to 11.

Starvation resistance assays

We estimated SR as the time elapsed (in hours) from the moment in which flies were exposed to the starvation diet until death. The starvation diet consisted of 5 ml of 1.7% agar in distilled water, which provided moisture but not food to the flies. For each line, 200 sexually mature flies of both sexes were released in egg-collecting chambers. In each chamber, a Petri dish containing an egg-laying medium (2% agar in distilled water and baker's yeast) was used for egg collection. Petri dishes were removed after 12 h and incubated at 25 °C until egg hatching (approximately 24 h). First-instar larvae were transferred to 10 vials (30 individuals per vial) containing lab medium. Adult flies were recovered from the vials and separated by sex under light CO₂ anesthesia and then transferred to vials containing lab medium before SR assays. Therefore, experimental flies were reared under optimal conditions in order to minimize the influence of environmental variation.

SR was measured in sets of flies that differed in mating status, mated and virgin flies. To obtain mated flies, adults were recovered from the vials every day and separated by sex. Then, groups of 15 males and 15 females were released in new vials and separated by sex after 3 days and maintained in vials (30 individuals per vial) with fresh food before the SR assay. These conditions consistently resulted in females insemination since we detected the presence of larvae in all vials. Although we cannot assure that all flies mated before the SR assay, preliminary experiments revealed that 99% of pairs of virgin females and males tested mated within 10 minutes (results not shown). In order to obtain virgin flies, newly emerged adults were recovered from the vials every 5 hours, separated by sex and maintained in groups of 30 individuals of the same sex in vials with fresh food.

The SR assay was performed with group of five 3 to 6-day-old flies. Seven to ten vials were set up for each line and sex. All vials were incubated at constant temperature (25°C ± 1) and under a 12 h light / 12 h dark cycle. Survival was recorded daily at 8.00 h, 14.00 h and 20.00

h until the death of all flies. Scores of SR for each individual fly were used to estimate mean survival time per replicate which was the variable considered in all statistical analysis.

Statistical analysis

A three-way analysis of variance (ANOVA), considering the line as the experimental unit, was performed to analyze variation in SR, according to the following model:

$$y = \mu + P + M + S + P \times M + P \times S + M \times S + P \times M \times S + \varepsilon_i,$$

where μ is the overall mean, P is the population effect (with three levels: Güemes, Cachi and Lavalle); M is the mating status effect (with two levels: mated and virgin), S is the sex effect and ε_i is the error term which is given by the among line variance. All factors in the ANOVA were considered as fixed. We performed “*a posteriori*” comparisons using Tukey’s method when an interaction term was significant.

We also performed two-way mixed ANOVAs for each combination of population and sex using the following model:

$$y = \mu + L + M + L \times M + \varepsilon_i$$

where μ is the overall mean, M (defined above) and L is the line effect (random, with as many levels as the number of lines analyzed in each population). The error, ε_i , is the among replicate variance term. Vials were considered as experimental units in these ANOVAs. In our experimental design differences among lines may be interpreted as an estimation of the genetic component of phenotypic variance, since lines (genotypes) may be considered as different genotypes (David et al. 2005). Likewise, the line by mating status interaction ($L \times M$) may be construed as an estimate of the genotype by environment interaction (G by E), which may be interpreted as genetic variation in phenotypic plasticity. Statistical analyses were performed using GLM procedure implemented in the STATISTICA 8.0 software package (StatSoft Inc. 2007).

Results

The general ANOVA revealed that populations differed significantly in their tolerance to starvation (Table 1). *A posteriori* pairwise comparisons showed that the significant population effect may be mainly attributed to differences between flies derived from Güemes (52.2 ± 2 h) and Lavalle (46.1 ± 1.1 h) ($p = 0.015$), whereas differences between Cachi (50.8 ± 1.9 h) and the other two populations were not significant. The ANOVA also revealed a significant mating status by sex interaction suggesting that mating differentially affected SR in males and females (Table 1). Further analysis showed that differences between mated and virgin flies were highly significant in females ($p = 0.0001$) but not in males ($p = 0.3$), that differences between males and females were significant in mated ($p = 0.0001$) but not in virgin flies ($p = 0.37$). Actually, mated females outlived virgins by more than 15 h and mated females were significantly more tolerant to starvation than mated males (Figure 1). The non-significant population by mating status by sex interaction suggests that the effect of mating status on SR was homogeneous across populations in both sexes.

Two-way ANOVAs performed for each combination of population and sex showed that differences among lines were significant in all cases (Table 2), indicating that natural populations harbour substantial genetic variation for SR (see also Goenaga et al. 2010). Also, the effect of mating status, over all possible isofemale lines, was significant in females but not in males, pointing to a sex-specific effect of mating status on SR. Though differences in mean SR between mated and virgin females were significant in Güemes and Cachi, and marginally significant in females derived from Lavalle (Table 2), the trends were consistent across all sampling localities. On the other hand, mean SR did not differ significantly between mated and virgin males across all sampling localities. However, it is important to note that the line by mating status interaction was significant in all populations in both sexes (Table 2).

The significance of this interaction may be due either to changes in variance among lines across mated and virgin flies and/or departures of the genetic correlation across mated and virgin flies from unity (see Robertson, 1959, Muir et al. 1992). Further analysis showed that, in females, the line by mating status interaction effect could be mainly accounted for a significantly greater variance among lines in mated than virgins in all sampling localities (Figure 2a, Levene's test $p = 0.0001$). In contrast, differences in variance among lines between mated and virgin males were not significant, suggesting that differences in the ranking order among lines across virgin and mated males may account for the significant line by mating status interaction (Figure 2b, Levene's test $p = 0.87$). Therefore, the impact of mating status appears to be line-specific and, thus accounts for the lack of an overall effect of mating status in males compared to females.

Discussion

The present study demonstrates that the populations analyzed are slightly differentiated and harbor substantial amounts of genetic variation in their ability to survive to food shortage; and, more importantly, that mating status (virgin vs mated flies) has a strong and homogeneous sex-specific effect on starvation resistance in all populations studied.

Concerning among population variation, flies derived from Güemes that inhabit a more benign environment, typical of lowland and tropical areas were, surprisingly, more tolerant to starvation than flies living in harsher environments like in Cachi and Lavalle. These results do not support our hypothesis that flies living in harsher environments, where face longer periods of food shortage, have greater SR (see also Goenaga et al. 2010). These observations are in agreement with the results of reports that propose that the ability to survive periods of food shortage does not correlate with ecologically relevant variables that co-vary with latitude or altitude or climatic variables (Robinson et al. 2000; Hoffmann et al. 2005a; Goenaga et al.

2010). Thus, other authors proposed that variation in SR may likely be a consequence of natural selection acting on genetically correlated traits, among which cold resistance is the main candidate (Hoffmann et al, 2002; Hoffmann et al. 2005b; Kenny et al. 2008; Ayroles et al. 2009).

Our study also shows that mating may confer a benefit to females by affecting survival under starving conditions. Mated females exhibited an increased tolerance to starvation in comparison to virgins, whereas mating status affected male starvation tolerance in a different way. These observations were visualized as a significant line by mating status interaction in all populations, which under our experimental design may be interpreted as a genotype by environment interaction (*G by E*) since each line can be considered as a different genotype and the mating status a particular type of environmental condition. Such particular *G by E* interaction suggests that the effect of mating status varied across genotypes in both sexes. In females, mating had a beneficial effect on most genotypes in terms of SR and the *G by E* interaction may be accounted for by an increased variance in mated than in virgins. In contrast, the significant *G by E* seems to be due to changes in the ranking order of genotypes across mated and virgin males, which explains the lack of an overall effect of mating status on male SR.

However, before further discussing the implications of our results we need to address a critical question of our experimental design, the lack of direct observations of mating.

Additional experiments showed that the great majority of pairs mated within 10 minutes after releasing flies in the vials. On one hand, females in our sample of mated flies probably differed in the number of matings achieved during the confinement with males, contributing in part to the increase in variance in the mating treatment in females. On the other hand, our sample of mated males likely included individuals that mated with 0, 1 or more females due

to differential mating success before the SR assay and this might have some influence in the reaction norms of males.

Similar effects of mating on SR in females have been reported in highly inbred lines with a long history of laboratory rearing (Canton-S and Oregon-R) (Rush et al. 2007) and lines selected for increased lifespan (Service 1989). Interestingly, such positive effect of mating on SR does not seem to be restricted to *Drosophila*. Indeed, mated females of the redback spider (*Latrodectus hasselti*) outlived unmated females when exposed to periods of food deprivation (Stoltz et al. 2010). The authors hypothesized that their observations may be consequence of different investments in mate attraction and reproduction between virgin and mated female spiders. In contrast to these findings, it has been shown that mating is costly for both females and males in the seed bug *Togo hemipterus*, as virgins survived longer than non-virgins when exposed to starvation conditions (Himuro and Fujisaki 2010).

The benefit of mating in females may be related to the transference of nuptial gifts during copulation. In fact, it has been reported that in the desert inhabitant *D. mojavensis* (Pitnick et al. 1997; Markow and O'Grady 2005) and other insects, males transfer nuptial gifts that could be incorporated by females and may help to face periods of food and water shortage (Edvardsson 2006; Gwynne 2008). Nevertheless, there is no evidence of nuptial gifts in *D. melanogaster*. Hence, the benefit that mating confers to females may be related to other factors that *D. melanogaster* males transfer with sperm, as male accessory gland proteins (*Acp*s), which are known to induce changes in female physiology and behavior (Wolfner 2002; Kubli 2003; Chapman and Davies 2004). Among other female post-mating responses (Ravi Ram and Wolfner 2007), it has been shown that food intake increases after mating in *Drosophila* females (Carvalho et al. 2006) and, actually, these results may provide a physiological explanation for the differences in SR between mated and virgin females. A small seminal peptide, the sex peptide (SP), has been identified as a crucial agent to induce

several post-mating responses (Liu and Kubli 2003; Wigby and Chapman 2005; Domanitskaya et al. 2007; Ravi Ram and Wolfner 2007) as sexual maturation and oogenesis in *Drosophila* adult females (Bownes 1989; Riddiford 1993; Soller et al. 1999) which may provoke a rise in food intake (Carvalho et al. 2006) resulting in the accumulation of fat reserves (Rush et al. 2007). Also, it has been shown that the effects of SP on egg laying, receptivity and lifespan are not only dependent on the female's diet (Fricke et al. 2010) but also that diet regulates whether exposure to SP transferring males is beneficial or costly to females (Fricke et al. 2008). Nevertheless, the increase in lipid content seems to be a common mechanism underlying the increase in SR (Chippindale et al. 1996; Harshman and Schmid 1998; Djawdan et al. 1998; Hoffmann et al. 2005b, Goenaga et al. unpublished). Therefore, the ability of mated females to survive longer than virgins to starvation may be explained by previous replenishment of lipid reserves as a consequence of the increase in food intake induced by proteins transferred by males during mating. Anyway, the mechanism through which mated females increase SR remains to be fully established, and is likely to be complex. In contrast, mating did not increase mean SR in mated males as compared to virgins, but affected the trait in a marked genotype-specific way. Unfortunately, studies addressing the putative mechanisms underlying sex differences in the effects of mating on SR (or other stress tolerance traits) are very limited and, most of them focus mainly on females rather than males (see above). Thus, specific experiments testing the hypothesis that the specific responses of males and females to mating are the result of sex-specific feeding responses are necessary. The genotype-specific effect of mating status on male SR is a novel observation in *Drosophila*. Genotypes varied in the magnitude and/or direction of their responses to mating status, implying that mated males of some isofemale lines exhibited greater SR than virgins and *viceversa* in other lines. The genotype-specific effect of mating status suggests some sort of antagonistic effects dependent of the mating status on male SR. Such environment

antagonistic effects are particularly interesting, since this type of interaction may contribute to the maintenance of genetic variation for adaptive traits, like SR, in nature.

Since in nature organisms are likely to face long periods of food shortage, it is reasonable to assume that SR has an adaptive significance (Breitmeyer and Markow 1988; Hoffmann and Parsons 1991). In this respect, our study shows that mating has, apart from the obvious reproductive advantage, an additional significance that may compensate the disadvantages resulting from the sexual conflict. Moreover, our results indicate that female-male interactions also affect other ecologically relevant traits, indicating that in a natural scenario of low food availability, mating may not be as harmful as envisaged by sexual conflict theories. This is an essential point that should be taken into consideration as some studies indicate that costs and benefits of mating could change depending on local environmental conditions (Candolin and Heuschele 2008; Fricke et al. 2008; Fricke et al. 2010). Indeed, studies in other *Drosophila* have shown that newly mated females are more resistant to desiccation, another trait related to stress tolerance, than virgins in the desert inhabitants *D. mojavensis* and *D. arizonae* (Knowles et al. 2004). In this case, the benefit of mating was explained by the effect of the large male ejaculate, which actively provides substantial nutrients to females (Pitnick et al. 1997).

In conclusion, the results reported in this study may have important implications in two directions. On the one hand, cast doubts on the “well” established idea that mating may impair female fitness and, on the other hand, open new avenues of research on the benefits of mating under stressful natural conditions.

Acknowledgements

The authors wish to thank J A Endler (Editor in Chief), D Hosken and M Hall (Associate Editors) and two anonymous reviewers for critical suggestions and advice that helped to improve previous versions of this paper. This research was supported by grants from

321 CONICET, ANPCyT and Universidad de Buenos Aires. J.G. was recipient of a fellowship
322 from University of Buenos Aires, J.M. is recipient of a fellowship from CONICET. J.J.F. and
323 E.H. members of Carrera de Investigador Científico of CONICET (Argentina).

324

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467

468 **Figure 1.** Effect of mating status on SR in females and males. Bars represent mean of SR \pm
469 SEM in virgin (black bars) and mated (white bars) females and males. $p < 0.05^*$; $p < 0.001^{**}$; p
470 $< 0.0001^{***}$.

471 **Figure 2.** Variation in SR in virgin and mated flies in females (a) and males (b) among lines
472 derived from 3 sampling localities.

Table 1. General ANOVA to analyze the effect of populations, mating status and sex on SR.

Source of variation	<i>df.</i>	<i>F</i>	<i>p</i>
Population	2	5.06	<i>0.008</i>
Mating status	1	34.41	<i>< 0.00001</i>
Sex	1	35.52	<i>< 0.00001</i>
Populations x Mating status	2	0.09	0.91
Populations x Sex	2	0.19	0.82
Mating status x Sex	1	13.70	<i>0.0003</i>
Population x Mating status x Sex	2	0.51	0.60
Error	92		

df. degree of freedom

Table 2. Partial ANOVAs performed for each combination of population and sex.

	Cachi		Güemes		Lavalle	
FEMALES	<i>df.</i>	<i>F</i>	<i>df.</i>	<i>F</i>	<i>df.</i>	<i>F</i>
Line	7	44.54***	6	11.32***	10	76.61***
Mating status	1	12.99**	1	17.57**	1	4.71*
Line x Mating status	7	28.30***	6	15.46***	10	60.12***
MALES	<i>df.</i>	<i>F</i>	<i>df.</i>	<i>F</i>	<i>df.</i>	<i>F</i>
Line	7	19.13***	6	7.35***	10	40.13***
Mating status	1	0.78	1	0.56	1	1.74
Line x Mating status	7	9.11***	6	24.01***	10	25.28***

df. degree of freedom * $0.05 < p < 0.10$; ** $p < 0.001$; *** $p < 0.0001$

Figure 1
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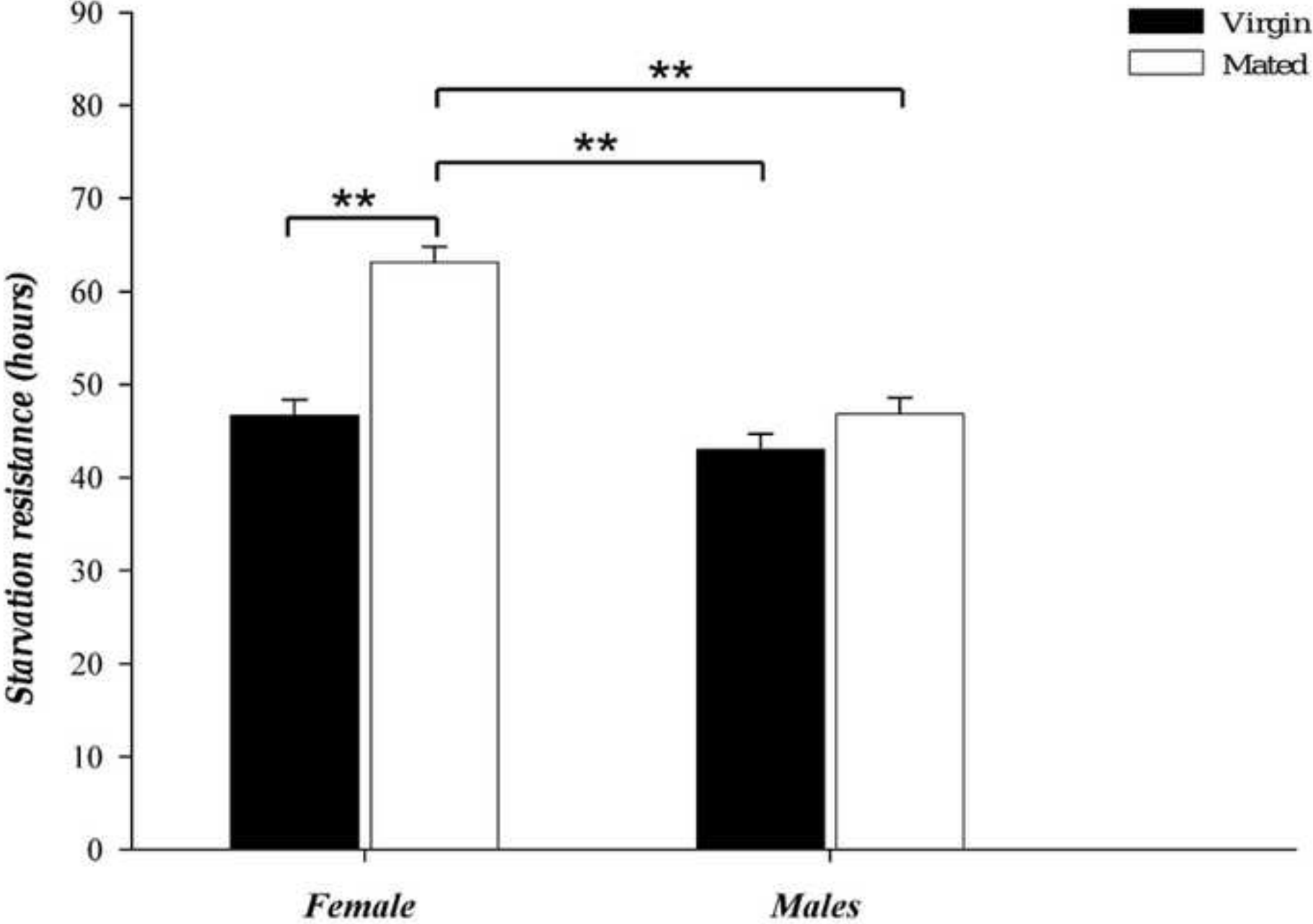


Figure 2
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