

ORIGINAL ARTICLE

The consequences of genetic variation in sex peptide expression levels for egg laying and retention in females

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The accessory gland proteins (Acps) that male *Drosophila melanogaster* produce and transfer to females during copulation are key to male and female fitness. One Acp, the sex peptide (SP), is largely responsible for a dramatic increase in female egg laying and decrease in female receptivity after copulation. While genetic variation in male SP expression levels correlate with refractory period duration in females, it is unknown whether male SP expression influences female egg laying or if any effect of SP is mediated by SP retention in the female reproductive tract. Here we measured the amount of SP retained in the female reproductive tract after mating and female egg laying after copulating with virgin males. We found no correlation between male SP expression levels and egg laying, or the amount of SP in the female reproductive tract after mating. Additionally, the amount of SP retained in the female did not influence egg laying. These findings suggest that additional factors, such as variation in other Acps, are important for the retention of SP in females and its quantitative effects on egg laying. It also shows that egg laying and refractory period response to SP is at least partially uncoupled.

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INTRODUCTION

For female insects, mating often induces large transcriptional, physiological and behavioural changes that are often triggered by components of male seminal fluid (Avila *et al.*, 2011). These changes and the molecules that cause them have been studied in detail in *Drosophila melanogaster* (Findlay *et al.*, 2008). In this species, over 138 seminal proteins and peptides are transferred (along with sperm) to females during mating (Avila *et al.*, 2011). These seminal fluid components are important for male and female fitness and, among other functions, they facilitate successful sperm storage (Adams and Wolfner, 2007), influence the outcome of sperm competition (Begun *et al.*, 2000; Chapman *et al.*, 2001), reduce female receptivity (Chapman *et al.*, 2003; Liu and Kubli, 2003) and stimulate ovulation (Heifetz *et al.*, 2000; Chapman *et al.*, 2001).

The function of one accessory gland protein (Acp), in particular, the sex peptide (SP) has been well-studied in *D. melanogaster*. When injected into (Chen *et al.*, 1988) or ectopically expressed (Aigaki *et al.*, 1991) in females, SP causes several profound changes in female behaviour. Two of the most striking of these are increased egg laying and reduced acceptance of male-mating attempts (Chapman *et al.*, 2003; Liu and Kubli, 2003). These are caused by the sex-peptide binding to a neuronal sex peptide receptor (SPR) in the female (Yapici *et al.*, 2008; Hasemeyer *et al.*, 2009; Yang *et al.*, 2009).

In *D. melanogaster* there is sequence variation within the SP gene and in upstream regions (Cirera and Aguade, 1997), which could potentially influence the regulation of SP. Smith *et al.* (2009) found variation in SP transcript abundance between genetic isolines of *D. melanogaster*, suggesting that transcriptional regulatory elements

may be important for the natural variation of SP expression. There is also evidence that variation in SP expression influences male fitness, with males that produce relatively high and relatively low amounts of SP RNA inducing the longest refractory periods in females (Smith *et al.*, 2009), which suggests that SP transcript levels is subject to disruptive selection. As the genetic variation of SP expression levels in males have been previously attributed to a quadratic relationship with female refractory period, we expect this same relationship to occur with female egg laying, a behaviour also attributed to SP. An alternative prediction is that SP acts on egg laying via a different mechanism to its effect on refractory period. This possibility remains to be tested. It is therefore important to determine if SP expression levels correlate with the egg-laying response by females, and if any potential relationship may be explained by the amount of SP retained by females after mating.

In this study, we tested whether male isolate SP expression levels or SP retention correlated with the number of eggs laid by females. To further investigate the mechanisms involved in this pathway, we also tested if SP expression levels correlated with the amount of SP retained in females after mating.

MATERIALS AND METHODS

Fly lines and quantitative real time polymerase chain reaction

We maintained all flies and measured SP expression levels as in Smith *et al.* (2009) using a subset of the inbred lines used in Ayroles *et al.* (2009). We used isolate numbers 75, 85, 159, 195, 345, 349, 378, 449, 535, 701 and 804, which were kindly donated to us by Frank Jiggins (University of Cambridge). We kept isolines on standard food at 25°C with a 12:12-h light:dark-light cycle. We diagnosed the isolines for the presence of *Wolbachia* infection following

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methods in Snook *et al.* (2000). We reared experimental flies in standardised conditions by placing 40 eggs on ~7 ml of food. As *Wolbachia* infection is known to influence several aspects of reproductive behaviour, we tested for its influence and found no effect (Smith, 2010). Thus, we refrain from any further discussion of effects of *Wolbachia*. We collected virgin males from the standardised egg-density vials and kept them individually with food before all experiments. For *SP* expression analysis, we snap froze 5-day-old males in liquid nitrogen and extracted total RNA (using Tri reagent (Sigma, St Louis, MO, USA)) to measure *SP* RNA levels with FullVelocity SYBR Green one-step Q-RT-PCR Reagents (Stratagene, Santa Clara, CA, USA) with *SP*-specific primers (Smith *et al.*, 2009). We used the housekeeping gene *RP49* to normalise expression levels.

Egg laying

We measured female egg laying after mating to males from the isolines of interest. To minimise unexpected individual differences amongst males within and between isolines, all males were reared individually to 5 days of age before experiments. To minimise individual differences amongst females of the standard isolate, females were collected in the same fashion as the experimental males but housed in groups of 20 before experimentation. We placed females without anaesthesia with individual virgin males from experimental isolines and observed copulation. We used 10 isolines with between 5 and 13 males per isolate. Immediately after the end of mating, we placed individual females in 9×2.5 cm universal tubes with a 2×1 cm laying cap containing standard food with a small amount of red food dye to increase the visibility of eggs laid on the food. Laying caps were replaced and eggs counted daily for 10 days. For analysis, we split the number of eggs laid by each female into two groups. Initially, we assessed the total number of eggs laid after 3 days, as this is when *SP* has its greatest effect on egg laying (Liu and Kubli, 2003). We also assessed the total number of eggs laid after a 10-day period. Previous work on *Drosophila* has shown that short-term measures (7 days) of reproductive output is significantly correlated with lifetime reproductive success (Taylor *et al.*, 2008), therefore 10 days of egg laying should be correlated with long-term reproductive output.

SP retention in females

For the measurement of *SP* retention in the female reproductive tract after mating, we conducted all matings with males from 11 of the isolines quantified for *SP* expression levels (Smith *et al.*, 2009). We reared males as described above and we used individual virgin females from a standard isolate, which was independent from the male isolines. Twenty-four hour before experiments we placed individual females in a single vial with food without live yeast. On the day of mating, we aspirated individual males (without anaesthesia) into a vial containing a single female and observed mating. One hour after each female had finished copulation, we snap froze females in liquid nitrogen and stored them at -80°C .

Following Sirot *et al.* (2009) and Wigby *et al.* (2009) with minor modifications, we used the ELISA technique to determine how much *SP* was present in the female reproductive tract after mating. Briefly, we dissected and homogenised the lower female reproductive tract in 10% Dulbecco's Phosphate Buffering Solution with protease inhibitor (Roche Complete Protease Inhibitor tablets, Indianapolis, IN, USA). We used ELISA plates (BD Biosciences, San Jose, CA, USA) with 1:2000 dilution of primary anti-*SP* antibody (kindly provided by E. Kubli (Peng *et al.*, 2005)) to bind to *SP* present in the samples and 1:2000 diluted horseradish peroxidase-conjugated goat antirabbit secondary antibody (Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) to react with 3,3',5,5'-tetramethylbenzidine substrate (KPL, Gaithersburg, MD, USA) in a colourimetric reaction. We then measured the optical density (OD) at $\lambda = 450$ nm (OD_{450}). For standardisation, we included serial dilutions of protein from the equivalent of the accessory glands from a single Canton-S fly male on each plate of ELISA reactions.

Statistical analysis

We conducted all statistical analysis using R for Macintosh version 2.9.2 (R Development Core Team, 2009). We conducted analysis on male isolate mean and individual values. Isoline means approximate genotypic associations (David *et al.*, 2005) and individual values approximates the phenotypic associations. Females that did not lay any eggs were removed from analysis for the egg-laying assay ($n = 5$). Results using isolate means and medians showed the same relationships, so here we only present results using isolate means.

To examine the impact of variation in isolate mean *SP* transcript levels on egg laying, we regressed isolate mean *SP* expression level values, including the quadratic term, against the number of eggs laid early (over 3 days) or in total (over 10 days) by females after mating to males from our experimental isolines using least-squared regression. We included mean female size as a covariate. We also regressed individual female egg laying against individual female size to test if larger female lay more eggs, a pattern that is expected for this species (Lefranc and Bundgaard, 2000) and hence a test of our egg-laying assay.

We analysed the relationship between *SP* retention and egg laying using least-squared regression with eggs laid either 3 or 10 days after mating as the response variable and *SP* retention and its quadratic term as the explanatory variables and female size as a covariate. We used least-squared regression to investigate the relationship between *SP* expression levels and *SP* retention in a similar manner.

RESULTS

We found no relationship between mean isolate *SP* expression levels and the number of eggs laid by females either 3 ($F_{(1,6)} = 0.08$, $P = 0.72$) or 10 days after mating ($F_{(1,6)} = 0.27$, $P = 0.62$) (Figure 1). Smith *et al.* (2009) previously showed a quadratic relationship between male *SP* expression levels and refractory period duration in females and therefore we considered such a relationship. As seen on Figure 1, there was no quadratic relationship between *SP*

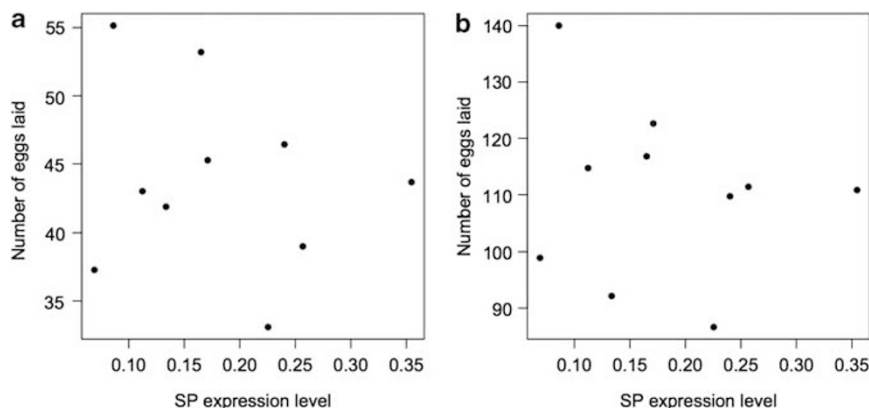


Figure 1 The relationship between the mean cumulative number of eggs laid in (a) 3 or (b) 10 days after mating and the mean *SP* expression levels of males from different isolines ($n = 10$). Results were the same when using line median or rank values (data not shown).

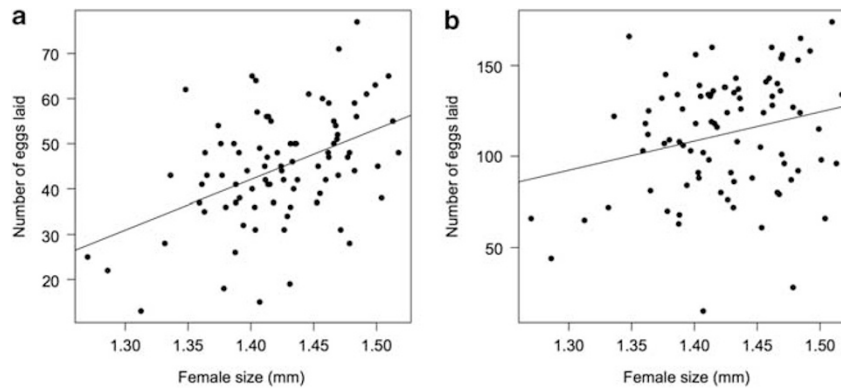


Figure 2 Female size is positively correlated with the cumulative number of eggs laid (a) 3 and (b) 10 days after mating. Five females did not lay any eggs and were not included in this analysis ($n=86$).

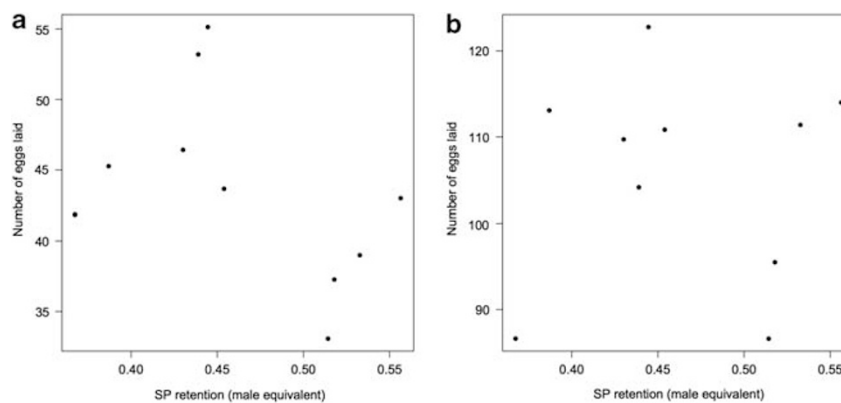


Figure 3 The relationship between the mean cumulative number of eggs laid in (a) 3 or (b) 10 days after mating and the mean SP retention of males from different isolines ($n=10$). Results were similar when using line median or rank values (data not shown).

expression and egg laying after 3 days egg laying ($F_{(1,6)}=0.38$, $P=0.56$) or 10 days egg laying ($F_{(1,6)}=0.14$, $P=0.72$). The mean size of females mated to males from each isolate was not significantly associated with number of eggs laid ($F_{(1,6)}=4.77$, $P=0.072$ and $F_{(1,6)}=1.07$, $P=0.34$ for cumulative eggs laid 3 and 10 days after mating, respectively). There was a significant relationship between the number of eggs laid 3 ($F_{(1,84)}=22.27$, $R^2=0.21$, $t=4.72$, $P<0.0001$) and 10 days ($F_{(1,84)}=5.63$, $R^2=0.06$, $t=2.372$, $P=0.02$) after mating when we used the individual female size and individual female egg laying in the model instead of line means (Figure 2).

We found no correlation between the isolate mean SP retention and isolate mean number of eggs laid after 3 ($F_{(1,6)}=2.48$, $P=0.17$) or 10 days ($F_{(1,6)}=0.03$, $P=0.88$). The quadratic term for isolate mean SP retention did also not significantly influence egg laying after 3 ($F_{(1,6)}=1.50$, $P=0.27$) or 10 days ($F_{(1,6)}=0.43$, $P=0.54$, Figure 3). Isoline mean female size did also not significantly influence egg laying 3 ($F_{(1,6)}=1.87$, $P=0.22$) or 10 days after mating ($F_{(1,6)}=0.01$, $P=0.93$). There was no significant correlation between isolate mean SP expression levels and isolate mean SP retention ($F_{(1,8)}=0.0001$, $P=0.99$ or the its quadratic term ($F_{(1,8)}=0.14$, $P=0.72$, Figure 4).

DISCUSSION

The focus of this study was threefold. First, we investigated whether SP expression levels of different male genotypes correlated with an important fitness correlate influenced by SP, the number of eggs laid by their mates after copulation. We found no correlation between the SP expression levels of male genotypes (as reported in Smith *et al.*

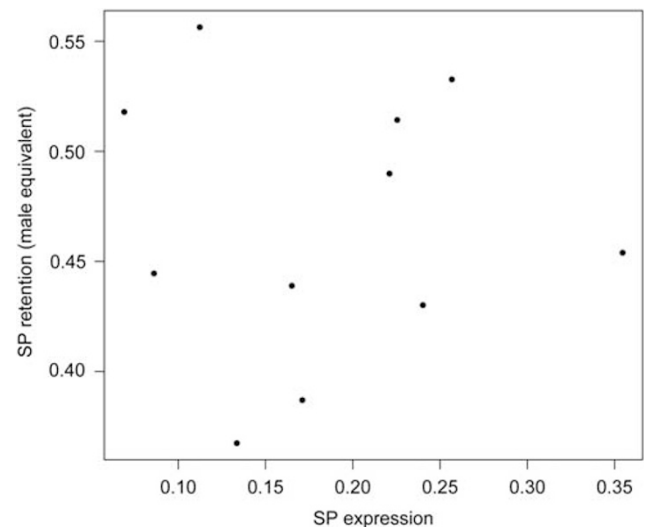


Figure 4 The relationship between isolate mean SP expression levels and isolate mean SP retention in female after mating to a virgin male ($n=11$).

2009) and the number of eggs laid by their mate after copulation. This is surprising given that previously we showed that genetic variation in the expression levels of SP among isolines of *D. melanogaster* males correlated with female propensity to remate in a manner suggestive of disruptive selection (Smith *et al.*, 2009). A possible explanation for the

lack of relationship between SP RNA expression levels and egg laying is that the amount of SP in the female after mating is not reflective by the SP transcript levels. Factors such as translational efficiency of SP RNA, transfer of SP and SP retention could all vary between isolines. It would be of interest to examine the interactions between multiple accessory gland proteins and their fitness effects in the future.

Secondly, to further investigate the potential mechanisms involved in the correlation between SP and egg laying, we tested whether the SP retention of male genotypes were associated with the number of eggs laid by the female that they mated to. We also found no correlation between SP retention and the number of eggs laid by that male's mate after mating. This may be due to effects of other seminal fluid proteins (for example, ovulin, (Heifetz *et al.*, 2000), CG33943 (Ravi Ram and Wolfner, 2007)) that also contribute to the male-induced effects on egg laying. The fact that refractory period duration is correlated with SP expression levels but that egg laying is not suggests that these two female responses to SP are at least partially uncoupled. Perhaps once SP binds to SPR within the female, these two responses are linked to different neuronal or hormonal pathways.

Thirdly, we tested for a correlation between SP expression levels and SP retention in females. Again, we found no correlation in the SP expression levels in males and the amount of SP that is retained in females after mating. Males can differentially allocate the amount of SP that they transfer to females during copulation (Wigby *et al.*, 2009). SP expression levels may then correlate with the amount of SP (protein) produced by males, yet differential male allocation of SP stocks (that is, not transferring all their stock at once) could potentially mean there is no correlation to SP expression levels and the amount of SP delivered and retained in females. It would be of interest to determine if there is genetic variation in the SP allocation patterns of males among sequential matings and to determine the fitness consequences of this. Additionally, a network of other accessory gland proteins is needed for SP retention in females (Ravi Ram and Wolfner, 2009), and variation in any of the members of that network could potentially affect the amount of SP retained by females. Similarly, variation in amounts of stored sperm could affect retention of SP. SP is actively cleaved from sperm within the female (Monsma *et al.*, 1990), and so interactions between female enzymes that cleave SP (Pilpel *et al.*, 2008) or male enzymes that inhibit enzyme activity (Wolfner *et al.*, 1997) could also dramatically alter the measureable amounts of SP in the female reproductive tract relative to male SP expression levels. Here, we tried to minimise the effect of female variation by using females from the same genetic isolate for all our tests. The lack of correlation between SP expression levels and SP retention in female also suggests that perhaps protein measurements will more accurately reflect relationships between peptides and their pathways than expression levels of genes that code for those peptides.

Overall, these results suggest that SP's effects on refractory period duration and egg laying are at least partially uncoupled. This may be female driven as the pathways influencing the two major effects of SP are unlikely to be identical. These results also suggest that expression patterns do not necessarily reflect the amount of protein that is functioning in the downstream pathways. The use of expression patterns in males may not reflect accurately the true interactions between the sexes.

DATA ARCHIVING

Data have been deposited at Dryad: doi:10.5061/dryad.77hn0.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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