

# Effects of maternal age and stress on offspring quality in a viviparous fly

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## Statement of authorship

SE and LH designed the study with inputs from ST, JSL and MB. RL and LH carried out the laboratory experiments. JSL carried out the analyses with input from SE, MB and ST and led the writing of the manuscript. RL, LH, AB, MB, ST and SE contributed to manuscript content and structure.

## Data accessibility statement

The data and R scripts to reproduce the analyses will be available at

[https://github.com/jenniesuz/tsetse\\_senescence.git](https://github.com/jenniesuz/tsetse_senescence.git).

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## Abstract

Many organisms show signs of deterioration with age, both in terms of survival and reproduction. Theory suggests that variation in such senescence patterns can be driven by resource availability or reproductive history. Here, we test this theory by manipulating nutritional stress and age at first reproduction and measuring age-dependent reproductive output in tsetse (*Glossina morsitans morsitans*), a viviparous fly with high maternal allocation. Across all treatments, offspring wet weight followed a bell-shaped curve with maternal age. Nutritionally stressed females had higher probability of abortion, produced smaller offspring with lower starvation tolerance. Despite this, there was no strong evidence of differences between treatments in the pattern of abortion probability, offspring wet weight or offspring starvation tolerance with age. Therefore, although we found strong evidence of general reproductive senescence in tsetse, variation in the onset and rate of senescence was not explained by resource allocation trade-offs or the costs of reproduction.

## 64 Introduction

65 Across the natural world, it is commonly observed that as individuals get older, they are more  
66 likely to die and have lower reproductive output (Nussey *et al.* 2013; Hoekstra *et al.* 2019;  
67 Zajitschek *et al.* 2019). Broad patterns of actuarial and reproductive senescence have been  
68 well described for a wide range of taxa (Nussey *et al.* 2013; Hoekstra *et al.* 2019; Zajitschek  
69 *et al.* 2019), particularly birds and mammals (Gaillard & Lemaître 2020). There is also  
70 extensive variation in the onset and rate of ageing, both within and across populations  
71 (Holand *et al.* 2016; Rodríguez-Muñoz *et al.* 2019; Cayuela *et al.* 2020). The role of  
72 individual heterogeneity, and subsequent variation in the onset and rate of ageing among  
73 individuals, however, has been neglected (Gaillard & Lemaître 2020).

74  
75 A central tenet of the evolution of senescence is that the strength of selection declines with  
76 age (Hamilton 1966), with consequences in terms of antagonistic pleiotropy (Williams 1957)  
77 or the accumulation of somatic damage (Kirkwood 1977). Life-history theory suggests both  
78 antagonistic pleiotropy and the accumulation of somatic damage can be explained by trade-  
79 offs between reproduction and other physiological processes (Kirkwood 1977; Boggs 2009;  
80 Baudisch & Vaupel 2012; Davison *et al.* 2014). A key assumption underlying this theory is  
81 that resources are limited and must be allocated either to reproduction or somatic  
82 maintenance (Partridge 1987; Boggs 2009). Reproductive senescence could also occur  
83 through physiological damage incurred directly from reproduction, or can be an adaptive  
84 strategy to prolong survival (McNamara *et al.* 2009).

85  
86 Experimental manipulation of access to mates or food can yield insights on the factors  
87 driving variation in senescence, including costs of reproduction and allocation of resources.  
88 For such experiments, insects are useful organisms as they have relatively short generation  
89 times, can be reared in large numbers, and access to mates and resources can be easily  
90 manipulated. Experiments with Lepidoptera, for example, have shown that delayed mating  
91 reduces fecundity but extends longevity (Unnithan & Paye 1991; Jiménez-Pérez & Wang  
92 2009). A large body of studies have also shown that females with access to fewer resources  
93 have lower overall reproductive output but longer lifespan (De Sousa Santos & Begon 1987;  
94 Ernsting & Isaaks 1991; Kaitala 1991; Chippindale *et al.* 1993; Tatar & Carey 1995; Curtis  
95 Creighton *et al.* 2009). These studies support the theory that senescence is, at least in part,  
96 caused by direct costs of reproduction through physiological damage, or arises as an indirect

consequence of resource availability. However, experimental manipulation of both age at reproduction and nutrition is required to tease apart the contributions of reproductive history and resource availability to senescence.

Here, we present such an experimental study, focussing on reproductive senescence in tsetse (*Glossina morsitans morsitans*). Tsetse are vectors of human and animal trypanosomiasis in Africa. They give birth to a single live larva weighing the same as the mother (Hargrove & Muzari 2015; Haines *et al.* 2020), approximately every nine days. Immature stages receive energy and nutrients from the mother only. Adults do not increase in size after emergence and have a relatively long lifespan for their small size, living for weeks rather than days (Hargrove *et al.* 2011).

Evidence for age-related changes in reproductive output in field and laboratory tsetse is mixed (Jordan *et al.* 1969; Langley & Clutton-Brock 1998; McIntyre & Gooding 1998). Key limitations to these studies are that flies were kept only under optimal laboratory conditions, not tracked individually and frequently grouped across ages. Preliminary data from the tsetse colony at the Liverpool School of Tropical Medicine (LSTM) provided evidence of reproductive senescence (Supporting Information, S1 File), but these data did not allow for teasing apart of within-individual from among-individual patterns (Monaghan *et al.* 2020). In this study, we therefore used a novel method of housing tsetse females to track individual mothers and their offspring from this colony.

Our first objective was to confirm reproductive senescence in tsetse, as shown by a decline in the number and quality of offspring with maternal age. Our second objective was to determine the effect of maternal nutrition and the physiological costs of reproduction on age-dependent patterns of maternal allocation. To achieve this aim, we manipulated nutritional stress by feeding adult females on a high- or low-quality diet and we varied potential costs of reproduction by delaying the age at which females were mated. Our third objective was to assess variation in reproductive senescence patterns among individuals, due to unmeasured heterogeneity in factors including size, condition and mate quality.

We hypothesised that: i) females mated later would experience a delayed and/or slower increase in the probability of abortion and decline in offspring quality, due to physiological costs of reproduction, resource allocation trade-offs, or both; and either: ii) nutritionally

stressed mothers would have an earlier, and potentially steeper increase in probability of abortion and decline in offspring quality, due to resource allocation-trade-offs; or iii) nutritionally stressed mothers may senesce more slowly, due to lower physiological costs of reproduction, if stressed females have lower reproductive output.

## Methods

Tsetse from a colony of *G. m. morsitans* maintained at LSTM were used for the experiments. The colony was established using pupae from Zimbabwe and has been maintained at LSTM since 2004. The colony was kept at 26°C ( $\pm 2$  °C), 68–76% relative humidity and a 12 h photoperiod. Flies were fed on a Monday, Wednesday and Friday on defibrinated horse blood (TSC Bioscience) which comprises 40–50% red blood cells.

### Measures of offspring quality

In addition to spontaneous abortions, we recorded the wet weight of viable pupae and, for a subset, we measured either fat, or whether they emerged and their starvation tolerance upon emergence (Fig. 1). Wet weight of pupae correlates with pupal volume in field flies and does not require destructive sampling (Hargrove 1999). We recorded emergence as we hypothesised that offspring with more reserves would be more likely to emerge. We measured starvation tolerance because, in the wild, newly emerged tsetse have a high probability of mortality (Hargrove *et al.* 2011), likely due to failure to find a blood meal before running out of energy reserves.

From preliminary data on the tsetse colony at LSTM (S1 File), we hypothesised that there would be an increase in the probability of abortion with age. If resource-allocation trade-offs exist we would also expect that the increase would be steeper for the nutritionally stressed treatment and potentially shallower for the mating delay treatment, relative to the control. However, for offspring wet weight and starvation tolerance we predicted that there would be an initial increase, reflecting the requirement for newly emerged flies to also divert energy and nutrients to the development of thoracic muscles (Anderson & Finlayson 1973), followed by a decline due to senescence. Lastly, we hypothesised that the effect of maternal age on starvation tolerance would be determined largely by energy reserves, as indirectly measured by offspring wet weight. Heavier offspring would have more reserves and therefore would be more likely to emerge and have higher starvation tolerance. Maternal age could also influence

offspring starvation tolerance, independent of effects on wet weight, through the quality of the resources provided.

### **Selection and mating of female tsetse for the experiment**

The *G. m. morsitans* colony at LSTM is maintained in 12 trays corresponding to each week of life. Females in the colony older than this are killed due to a decline in reproductive output. To select females for the experiment, we collected equal numbers of pupae from each tray over a one-week period for emergence (S2 File). At the time of mating (upon emergence or after a delay), females were mated at a ratio of 2:1 (female:male) with one-week old males for 48 hours. Males were then removed, and females were selected for the experimental treatments.

### **Treatments**

We set up control, nutritional stress and mating delay treatments, each consisting of 96 adult female *G. m. morsitans* (Fig. 1, S2 File). Because reducing the amount of haemoglobin in a bloodmeal results in lower pupal wet weights (Kabayo & Langley 1985), we chose to dilute red blood cells with serum to produce a low-quality diet for the nutritional stress group. Trials testing different ratios of red blood cells to serum showed that flies fed on c. 10% red blood cells produced lighter pupae but had similar survival over a 50-day period, compared to flies fed on c. 45% red blood cells (S3 File). We thus ascribed the 10% red blood cell treatment as the ‘nutritional stress’ treatment and 45% red blood cells as ‘control’. For the mating delay treatment, virgin females were kept in communal cages for three weeks post-emergence. Virgin females continue to ovulate, but mature eggs eventually disintegrate (Ejezie & Davey 1977). Once mated, they were separated into individual cages. The experiment was then run until mothers were 100 days old. (Hargrove *et al.* 2011). In addition, unpublished data from the LSTM colony indicated that by 100 days the probability of abortion had increased from c. 0 to 0.25 and the size of pupae produced had declined to similar sizes seen from first-time mothers (S1 File).

### **Housing of individual adult females and feeding regime**

Flies were housed in individual cages, which were placed on pupal collection trays, made from acrylic extrude (described in S2 File), to allow individual tracking of female reproductive output and survival. Cameras were fixed below the cages to record weekend activity and to identify exact larviposition dates. Females were monitored daily between

Monday and Friday and if death occurred, we recorded the date of death. To determine the exact date of death for females found dead on a Monday morning, we reviewed video recordings from Friday evening to Monday morning (S2 File).

Experimental flies were fed following the colony schedule (see above). To feed individual females, we placed cages on a shallow grid (S2 File). The grid ensured each female was provided with the same amount of blood and had equal opportunity for feeding, with c. 100  $\mu$ l of blood provided per female. We heated the tray to 37°C and covered the blood with a silicon membrane, then left flies to feed for 45 minutes, which was sufficient time for most flies to feed.

### **Offspring measurements**

Time of larviposition was recorded, including whether or not larvae produced were viable (Baldry *et al.* 1992). Production of non-viable offspring was considered a spontaneous abortion ('abortion' hereafter). It was not possible to record egg abortions as they were not visible due to their small size. For pupae collected on a Monday morning, we consulted the video recordings to determine the exact day of larviposition (S2 File). For aborted larvae, however, we could not determine the date for those aborted over the weekend, because early larval stages were also not visible on the camera and were therefore recorded as occurring on the Friday. For viable pupae, we measured the wet weight, to 0.1 mg.aining 30% were destructively sampled for fat analysis (S2 File) in order to quantify how wet weight correlates with fat reserves.

Pupae assigned for emergence studies were placed singly into 50 ml Falcon tubes with a 3 mm hole drilled in the centre of the screw cap to allow air flow. Pupae were observed daily for date of emergence on working days and again video recordings were used to determine the date of emergence on weekends. Pupae were observed for a maximum of 50 days. Any pupae that had not emerged by 50 days, were recorded as a failed emergence (Hargrove 2004). Each emerging fly remained in the tube until it died of starvation. Sex of emerging flies and date of death were recorded.

### **Statistical analyses**

For each analysis described below, models were compared using Akaike's information criterion (AIC). If the ratio of the sample size to the number of model parameters was <40,

we used AIC corrected for small sample size (AICc) (Burnham & Anderson 2002). Figures were produced using the model with the lowest AIC. For model comparison, the difference in AIC between each model and the lowest AIC model ( $\Delta_i$ ) and Akaike weights ( $\omega_i$ ) were calculated. Akaike weights sum to one and provide a relative indication for the weight of evidence for any one model as the best approximating model (Burnham & Anderson 2002). We analysed the data using R version 3.6 (R Core Team 2014). The data and R scripts to produce the results can be accessed at ([https://github.com/jenniesuz/tsetse\\_senescence.git](https://github.com/jenniesuz/tsetse_senescence.git)).

For the probability of abortion, offspring wet weight, and starvation tolerance, we carried out statistical analyses for each treatment separately using linear and generalised linear mixed effects models implemented with the ‘lme4’ and ‘nlme’ R packages (Bates *et al.* 2015; Pinheiro *et al.* 2018). For each analysis described below maternal age in days was incorporated as a continuous variable. All models with and without random effects were fitted and simultaneously compared using maximum likelihood estimation, assuming that the bias in the variance components would be relatively small ( $n/(n-p)$ : 1.009 for the probability of abortion; 1.049 for offspring wet weight; and 1.095 for offspring survival, largest values across treatments reported). Models with and without a random intercept were compared to assess evidence for variation among individual mothers in the probability of abortion, offspring wet weight and offspring starvation tolerance. We also compared models with and without a random slope for maternal age, to assess evidence for variation in senescence patterns among individual mothers.

The effect of maternal age on probability of abortion, offspring wet weight and offspring starvation tolerance were compared between treatments using fitted coefficients and 95% confidence intervals from the model with the lowest AIC. We took this approach to analysis rather than including treatment as a factor in models as it is simpler and does not require the assumption of equal variance between treatments.

Logistic regression was used to quantify the effect of maternal age on the probability of abortion for each treatment, assuming a linear relationship between maternal age and the log odds of abortion. For the effect of maternal age on offspring wet weight and starvation tolerance, models including maternal age as a cubic, quadratic, logistic or linear effect were compared. For wet weight, we also compared the most parsimonious model with a model fit using generalised additive modelling (GAM). Cubic regression splines were fitted for each

treatment, with maternal age as the explanatory variable and accounting for multiple offspring from individual mothers, using the mgcv R package (Wood 2017). Generalised additive model fits with knots – locations where the slope changes – ranging from 3 to 10 were compared using AICc. The correlation between offspring wet weight and fat was summarised using Pearson's correlation coefficient.

For offspring starvation tolerance, there was no censoring and the data were approximately normally distributed (S4 Fig.). The relationship between offspring wet weight, sex, maternal age and number of days to starvation was therefore modelled using linear mixed effects models for each treatment. Maternal age was included in models as described for offspring wet weight. In addition to maternal age and wet weight, we also included offspring sex as females are larger than males on emergence (Hargrove *et al.* 2019).

We repeated the above analyses for the nutritional stress treatment, excluding females that had died, to ensure our results were not affected by the possibility that females who died allocated more to their offspring. For the other two treatments this was not done as >90% of mothers were still alive by the end of the experiment. Lastly, an analysis of maternal survival is provided in S5 File.

## Results

### Abortions

The probability of abortion increased with age across all treatments (models without age: control  $\omega = 0.000$ ; mating delay  $\omega = 0.001$ ; and nutritional stress  $\omega = 0.00$ ) (Tables S6 1-3). The increase in the odds of abortion with age were similar across treatments (from the model with lowest AIC: control 1.072, C.I. 1.054 – 1.094; mating delay 1.053, CI 1.025 – 1.086; and nutritional stress 1.047, CI 1.034 – 1.061) (Fig. 2, S6 4-6). Females in the nutritional stress treatment had higher probability of abortion at any age compared with the other two treatments (Fig. 2).

For all treatments, including a random intercept and slope in models resulted in a singular fit indicating that models were overfitted. We did not therefore assess evidence for individual variation in the effect of maternal age on probability of abortion. There was, however, strong

evidence for variation among individual mothers in the probability of abortion for the nutritional stress treatment (model with random intercept  $\omega = 0.985$ ; model without  $\omega = 0.015$ ). This was not the case for the control (model with random intercept  $\omega = 0.289$ ; model without  $\omega = 0.711$ ) or mating delay treatments (model with random intercept  $\omega = 0.368$ ; model without  $\omega = 0.631$ ) (Tables S6 1–3). Plots of the raw data for this analysis, and analyses described below, can be viewed in S7 File.

### **Offspring wet weight and fat**

Across all treatments, wet weight of pupae increased and then declined with maternal age until the end of the experiment (Fig. 3). Model selection results provided strong evidence for a quadratic effect of maternal age on offspring wet weight (control – quadratic effect including random intercept and slope  $\omega = 0.719$ ; mating delay – quadratic effect with random intercept and slope  $\omega = 0.562$ , without random slope  $\omega = 0.272$ ; and nutritional stress – quadratic effect with random intercept and slope  $\omega = 0.866$ ) (Tables S6 7-9). There was no support for a log, or a linear effect of maternal age for any of the treatments ( $\omega = 0.000$ ). In addition, across treatments, the predicted effect of mother age at the population level, assuming a quadratic fit, was within the standard errors of predictions from GAM fits to the data (S8 File).

The fitted quadratic curves and coefficients were similar between treatments (from the model with lowest AIC, which included a random intercept and slope, coefficient for maternal age: control 0.520 (0.466 – 0.574); mating delay 0.564 (0.406 – 0.723); and nutritional stress 0.502 (0.420 – 0.584), coefficient for maternal age squared: control -0.0040 (-0.0045 – -0.0035); mating delay -0.0042 (-0.0054 – -0.0030); and nutritional stress -0.0040 (-0.0048 – -0.0032)) (Tables S6 10 - 12).

For all three treatments, there was strong evidence for individual variation among mothers in the wet weight of their pupae (for all treatments, all models without random effects  $\omega = 0.000$ ). This variation was highest in the mating delay treatment with a random intercept variance from the most parsimonious model of 11.820, compared with 7.307 for the control and 2.379 for the nutritional stress treatment (Tables S6 13 - 15).

There was evidence for variation among mothers in the effect of maternal age on wet weight for the control and nutritional stress treatments (quadratic fit with random intercept and slope:

control  $\omega = 0.719$ ; and nutritional stress  $\omega = 0.866$ ). There was less support for this in the mating delay treatment (quadratic fit with random intercept and slope  $\omega = 0.562$ , next lowest AIC model - quadratic fit with random intercept only  $\omega = 0.272$ ).

Offspring fat increased linearly with offspring wet weight (Pearson's correlation coefficient 0.554) (File S9), thus for brevity we focus on wet weight as our trait of interest, but the patterns are qualitatively similar if offspring fat is considered instead.

### **Offspring emergence and starvation tolerance**

During the 50-day pupal observation period, 96% of 355 pupae from the control, 93% of 187 pupae from the mating delay, and 91% of 232 pupae from the nutritional stress group emerged successfully. All emerged offspring subjected to starvation were dead by 15 days post emergence. Although the wet weight of pupae was similar between females and males (Fig. 4a), across all three treatments there was strong evidence that female offspring survived on average longer than males, but not by more than a day (Tables S6 19 – 21), with  $\omega \leq 0.01$  for models not including sex (Tables S6 16 - 18, Fig. 4b).

There was strong evidence for an additional effect of maternal age on offspring starvation tolerance, in addition to offspring wet weight, for the nutritional stress treatment (models without age  $\omega = 0.000$ ). Offspring from young mothers in the nutritional stress treatment were particularly vulnerable to starvation (Fig. 4b), with evidence for a cubic effect of maternal age in this treatment (cubic effect including random intercept  $\omega = 0.705$ ). While there was also evidence for maternal age effects on offspring starvation tolerance in the other two treatments, there was not sufficient information in the data to quantify the relationship between maternal age and offspring starvation tolerance for the control or mating delay treatments (lowest AIC model: control – quadratic effect and no random effects  $\omega = 0.298$ ; mating delay – quadratic effect with no random effects  $\omega = 0.212$ ) (Tables S6 16-17).

For the mating delay treatment, singular fits were obtained for models assuming a quadratic or cubic effect of mother age and including a random intercept and slope, therefore there was not sufficient data to assess for this treatment whether differences among individual mothers carried through to emerged offspring starvation tolerance. There was also insufficient information in the data to provide evidence for among-mother variation in the control treatment, with the difference in AIC weights between models with and without random

effects  $<0.15$  (Tables S6 16 - 18). However, there was evidence for an effect of individual mother on starvation tolerance in the nutritional stress treatment (all models without random effects  $\omega \leq 0.04$ ) (Tables S6 16 - 18), but not for variation among individual mothers in the effect of maternal age on offspring starvation tolerance ( $\omega \leq 0.005$ ).

The results of the above analyses for the probability of abortion, offspring wet weight and offspring starvation tolerance were unaffected by removing from the analysis mothers in the nutritional stress treatment that had died before the end of the experiment. The analyses can be found in S10 File and compared with results in S6 File.

## Discussion

While there is substantial evidence of actuarial senescence in insects and some evidence for reproductive senescence (Nussey *et al.* 2013; Zajitschek *et al.* 2019), the drivers of variation in senescence patterns are not fully understood. Here we manipulated both age at mating and nutrition to quantify reproductive senescence in tsetse, a viviparous fly with high maternal allocation and iteroparous reproduction. Both offspring weight and starvation tolerance declined with maternal age, after a peak, yet these patterns of senescence were similar across treatments. Contrary to predictions from life history theory, therefore, neither changes in maternal allocation nor resources affected the timing and rate of reproductive senescence, in terms of offspring quality.

We did observe a steeper increase in the hazard of mortality with age for nutritionally stressed mothers (S5 File). This suggests that females do not have a survival benefit from reduced reproductive effort, as nutritionally stressed mothers had higher probability of abortion and produced smaller offspring at any age compared with mothers in the control and mating delay groups. This contrasts with findings from studies of other insects where nutritional stress both reduced reproductive output and either maintained or even extended lifespan relative to a control group (De Sousa Santos & Begon 1987; Ernsting & Isaaks 1991; Kaitala 1991). Taken together, it may be that factors other than the direct costs of reproduction through physiological damage, or indirectly through resource allocation trade-offs, impose higher mortality rates. Alternatively it may be that, given the extreme maternal allocation in tsetse, even though females produce relatively smaller offspring they still pay a high cost of reproduction in terms of physiological damage; and females on a poor quality diet experience this cost to a greater extent in terms of impact on mortality. This is supported

by data from field-caught tsetse, where smaller females invest relatively more of their fat in their offspring, even though their offspring were smaller (Hargrove *et al.* 2018).

We find that females experiencing nutritional stress have a relatively higher rate of abortion. Hargrove and Muzari (Hargrove & Muzari 2015), using field collected *G. pallidipes*, showed that transfer of the majority of fat to the larva occurs only after c. 80% pregnancy has been completed. Therefore, a female could potentially abort a larva if there are not enough fat reserves for a full-term pregnancy. Evolutionary models tailored to tsetse life-history, with high investment in single offspring across multiple reproductive bouts, could yield insights into whether such spontaneous abortion is an adaptive strategy to retain reserves for future reproduction, or a result of physiological constraints that limit the reserves available (McNamara *et al.* 2009).

Our study highlights the benefits gained from individual-level data to understand senescence. The bell-shaped relationship of offspring quality with age may have contributed to the relatively small effects of age evident in previous studies where grouped ages and mean values were used, rather than tracking reproductive output from individual females (Langley & Clutton-Brock 1998; McIntyre & Gooding 1998). The bell-shaped pattern observed here is strikingly similar in shape across treatments and reflects the general bell-shaped pattern of reproductive senescence observed across diverse taxonomic groups e.g. (Velando *et al.* 2006; Sharp & Clutton-Brock 2010). As summarised by (Monaghan *et al.* 2020), a bell-shaped relationship between maternal age and reproductive output, or offspring quality, can arise from population-level effects, due to selective disappearance, but also from individual-level effects. By tracking individual mothers, here we provide evidence that, for tsetse, the bell-shaped relationship between offspring wet weight and maternal age is a consequence of individual effects. The initial increase in offspring wet weight could be a consequence of mothers accruing reserves each time she takes a bloodmeal, so that at each sequential reproductive event she has more energy reserves that can be provided to the offspring. The subsequent decline could then be explained by senescence.

Tracking individual mothers provided insights into individual variation in maternal allocation. There was marked variation in offspring wet weight among mothers, particularly for the mating delay treatment and variation in senescence patterns for wet weight, particularly for nutritionally stressed mothers. Some females in the mating delay treatment

consistently produced smaller than average offspring, across all ages, and these females contributed more to the individual heterogeneity than those producing consistently heavier offspring. For nutritionally stressed mothers, variation between individuals in offspring wet weight increased as mothers aged. These observations suggest that variation in offspring quality is affected not only by mother size but also unmeasured aspects of her condition. The large amount of variation between individuals, in offspring weight and changes in the extent of variation with maternal age in this study was unexpected, suggesting that future studies quantifying the relative roles of mother size and physiological condition on offspring size across different ages would be valuable.

Offspring from young mothers that were nutritionally stressed had the lowest starvation tolerance. We also found that maternal age affected offspring starvation tolerance independently of age-effects mediated through wet weight, suggesting that there may be other factors associated with maternal age that influence the quality of offspring. More subtle effects of maternal age on the quality of resources transferred to offspring warrant further investigation in tsetse and other species. For tsetse, during late stages of pregnancy females not only transfer fat but also amino acids. It may be that young nutritionally stressed females are limited in these amino acids. (Cmelik *et al.* 1969). The authors reasoned that the tyrosine and phenylalanine obtained from a single bloodmeal is unlikely to be sufficient to meet the amount required by offspring and that a surplus stored from previous bloodmeals may be required. This suggests that size may be limited when nutrients are limited, irrespective of the amount of energy available. It also demonstrates that resource allocation processes are likely more complex, and more nuanced studies on the effects of the quality of resources as well as quantity may be required to understand the ageing process.

We focused on reproductive senescence in this study. One limitation is that we did not continue the experiment beyond 100 days to quantify more fully mother survival for all three treatments. Our evidence of actuarial senescence in the nutritional stress treatment supports analysis of mark-recapture studies of *G. m. morsitans* in the field (Hargrove *et al.* 2011) which also showed an increase in mortality as a function of age. Considering this, an additional experiment to test whether females mated later experience a delayed onset in actuarial senescence would be informative, given the similar rates of reproductive senescence observed across treatments.

We also note that our conclusions with respect to reproductive senescence are constrained to this 100-day period. While, in principle, it would be optimal to run the experiment until all females had died, the >1/4 of offspring being aborted and fewer females surviving would result in relatively small sample sizes beyond this time point. However, we are confident that our study timeframe captures an ecologically relevant period to study age-dependent allocation in tsetse, given that field studies show that 90% of females have died before 100 days (Hargrove *et al.* 2011). Lastly, we acknowledge that our conclusions may not necessarily extend to wild flies, and further study of tsetse in the field would need to be carried out to confirm this.

To conclude, our results provide evidence of a bell-shaped relationship between maternal age and offspring quality at the individual-level for the iteroparous tsetse fly. This complements the existing body of work from other species, that has predominantly shown a similar curve for the number of offspring produced as mothers age (Monaghan *et al.* 2020). Contrary to predictions from life history theory, however, our study did not find evidence of a decline in offspring quality as a function of direct costs of reproduction or resource allocation trade-offs.

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604

605 **Figure 1 Overview of experiments.** M – mating. Colour-coded silhouette of pregnant tsetse  
606 with larva, timeline shows a silhouette of a pupa and tubes containing blood, the dark red  
607 indicates red blood cell and beige indicates serum.

608

609 **Figure 2 Predicted probability of larval abortion as a function of maternal age, by**  
610 **treatment.** Predicted probabilities from generalised linear mixed effects model fits to the  
611 data and 95% prediction intervals. Plots of raw data are provided in S7 File.

612

613 **Figure 3 Offspring wet weight as a function of maternal age and treatment.** Showing  
614 model fits to the data: thick line – population level, thinner lines – individual level. Points –

615 average wet weights for 10-day intervals and 95% confidence intervals. Plots of raw data are  
616 provided in S7 File.

617

618 **Figure 4 Effect of sex, wet weight and maternal age on starvation tolerance (the number**  
619 **of days a newly emerged fly can survive starvation).** a) Wet weight as a function of  
620 offspring sex by treatment; b) Predicted survival time based on linear mixed effects model.  
621 Days adults survived starvation is plotted against maternal age. Prediction for each wet  
622 weight quartile shown. Plots of raw data are provided in S7 File.

623

624 **Supplementary Files**

625 **S1 File. Pilot data from the tsetse colony at LSTM.**

626 **S2 File. Additional methods.**

627 **S3 File. Nutritional stress pilot.**

628 **S4 Figure. Histogram of offspring survival time by treatment.**

629 **S5 File. Analysis of mother survival.**

630 **S6 File. Statistics.**

631 **S7 File. Raw data plots.**

632 **S8 File. Generalised additive models.**

633 **S9 File. Offspring fat as a function of wet weight.**

634 **S10 File. Statistics excluding mothers in the nutritional stress treatment that died before**  
635 **the end of the experiment.**

636

