

Rapid and costly ageing in wild male flies

Short-lived insects surprisingly still suffer senescence under natural conditions.

Ageing (senescence) has never been demonstrated convincingly in any insect in the wild, where mean lifespans are probably much shorter than in the laboratory¹, and most evidence for senescence in other wild animals (such as mammals) is limited to their reduced survival with age². Here we show that ageing is detectable in wild populations of a very

short-lived insect, the antler fly (*Protophila litigata*), and causes debilitating and costly effects that force a decline not only in survival probability, but also in the reproductive rate of males. Our findings argue against the possibility of a trade-off between fitness components, whereby survival may decline without senescence if investment in reproduction increases with age³, and indicate that ageing rates are subject to intense selection in the wild.

Although theory predicts the evolution of rapid senescence in organisms that experience high extrinsic (age-independent) mortality rates⁴, it has been suggested that very few individuals in these groups (such as insects or small mammals) survive long enough in the wild to exhibit detectable senescence^{5,6}.

We tested for senescence in a wild population of the antler fly, a small dipteran that breeds exclusively on discarded antlers of moose and deer. The tendency of adult flies to spend their lives on a single antler, as well as the long duration of their mating (2.3 h; ref. 7), facilitate the acquisition of field data on mating success and survival. We surveyed mating aggregations on nine moose antlers every 2 h over 72 days, and recorded the presence and mating status (single or coupled) of each of 609 individually marked males⁸.

The daily probability of mortality in males increased with age (Fig. 1a; likelihood ratio comparisons with constant model: Gompertz, $\chi^2_1 = 4.293$, $P = 0.0383$; two-parameter Weibull, $\chi^2_1 = 3.931$, $P = 0.0474$; three-parameter Weibull, $\chi^2_1 = 3.950$, $P > 0.1$). The Gompertz model fitted marginally better than the two-parameter Weibull, on the basis of Akaike's information criterion (AIC; Gompertz, 78.11; Weibull, 78.29) and bootstrap analysis (Gompertz chosen in 5,838 of 10,000 iterations).

The daily mating rate decreased linearly with age (Fig. 1b; linear AIC, 2,337.94; two-parameter Weibull AIC, 2,338.72; constant AIC, 2,345.03). Although the mean mating rate may decline with age because males that mate less frequently live longer, rather than because of senescence, we found no evidence of a trade-off between mean daily mating rate in early life (age, 2–5 days) and lifespan ($n = 15$ lifespan classes, $r^2 = 0.05$, $F = 0.74$, $P > 0.4$). Mating rate is probably a reliable indicator of male reproductive rate in this species, as take-overs are rare and females lay eggs immediately after copulation⁷. Thus, as predicted by theory⁴, we found that both survival

and reproductive rate declined with age.

To assess the relative fitness costs of senescent declines in survival and reproduction, we compared the reproductive value (that is, the expected lifetime mating success) of males at age 0 days with hypothetical reproductive values based on constant initial (presenescent) daily rates of survival or reproduction. Males lost 1.7 times more fitness as a result of declining reproductive rate than from declining survival rate, but this difference was not significant (bootstrap $P = 0.0992$).

However, a numerical analysis comparing the relative sensitivity of fitness to small, additive reductions in survival and mating rates at different ages of onset⁹ indicated that at younger ages (< 20 days), selection was stronger on survival than on mating rate, whereas at older ages (> 20 days), selection was stronger on mating rate than on survival. Net fitness losses from senescence in both traits amounted to about 20% (95% confidence limit: 13–26%) of potential fitness. However, our analysis may underestimate both total fitness losses and relative losses from senescence in reproductive rate if low-quality individuals are under-represented in older age classes¹⁰ and if male fertility declines with age¹¹.

We have shown that senescence can be detected in wild insects, despite a high extrinsic mortality rate (about 13% per day) and brief median lifespan (6 days). Because there is a simultaneous decline in both survival and reproduction, our results indicate unambiguously that senescence occurs in this wild insect population. The high net fitness costs of senescence confirm a basic assumption of evolutionary theories — that senescence rates are under strong selection in wild animals.

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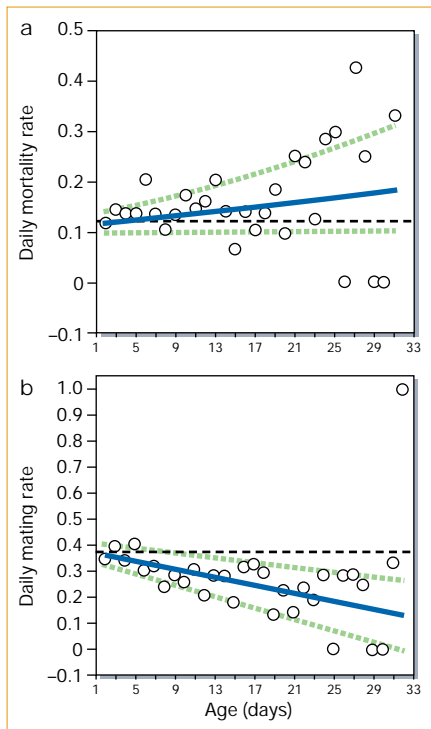


Figure 1 Evidence of senescence in male antler flies (*Protophila litigata*) in the wild. **a**, Mean daily probability of mortality (hazard rate; circles), $h(t)$, with the fitted Gompertz model (solid blue line) and initial hazard rate (dashed black line; instantaneous $h_0 = 0.1359$; $\gamma = 0.0166$). **b**, Mean daily mating success (circles), $m(t)$, with the fitted linear model (solid blue line) and initial mating rate (dashed black line; $\lambda = 0.3741$; $\kappa = -0.0076$, likelihood ratio $P < 0.0001$). Dotted green lines indicate 95% confidence limits. Marked males were released at antlers 24–48 h after adult emergence and were presumed to have died on the day after their last sighting, as mean daily sighting probability was high (63%) and the probability of immigration was very low. Mean lifespan, mating success and mating rate did not change over the season ($P > 0.2$ for all tests). Senescence models were fitted by maximum likelihood, with daily survival and mating rates as random binomial and Poisson processes with age-specific probabilities. Confidence limits were generated by 10,000 bootstraps. Hazard-rate models compared: constant, $h(t) = \lambda$; Gompertz, $h(t) = h_0 \exp(\gamma t)$, fitted as $N_{i+1}/N_i = \exp[-(h_0/\gamma)(\exp(\gamma(t+1)) - \exp(\gamma t))]$; three-parameter Weibull, $h(t) = h_0 + \alpha t^\beta$, fitted as $N_{i+1}/N_i = \exp[-h_0 - (\alpha(\beta+1))((t+1)^{\beta+1} - t^{\beta+1})]$; two-parameter Weibull ($h_0 = 0$). Mating-rate models compared: constant; Weibull; linear, $m(t) = \lambda + \kappa t$. Because data for the mark–release day were incomplete, these data (and the 75 males that survived for less than 1 day after release) were excluded from the analyses.

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Competing financial interests: declared none.