

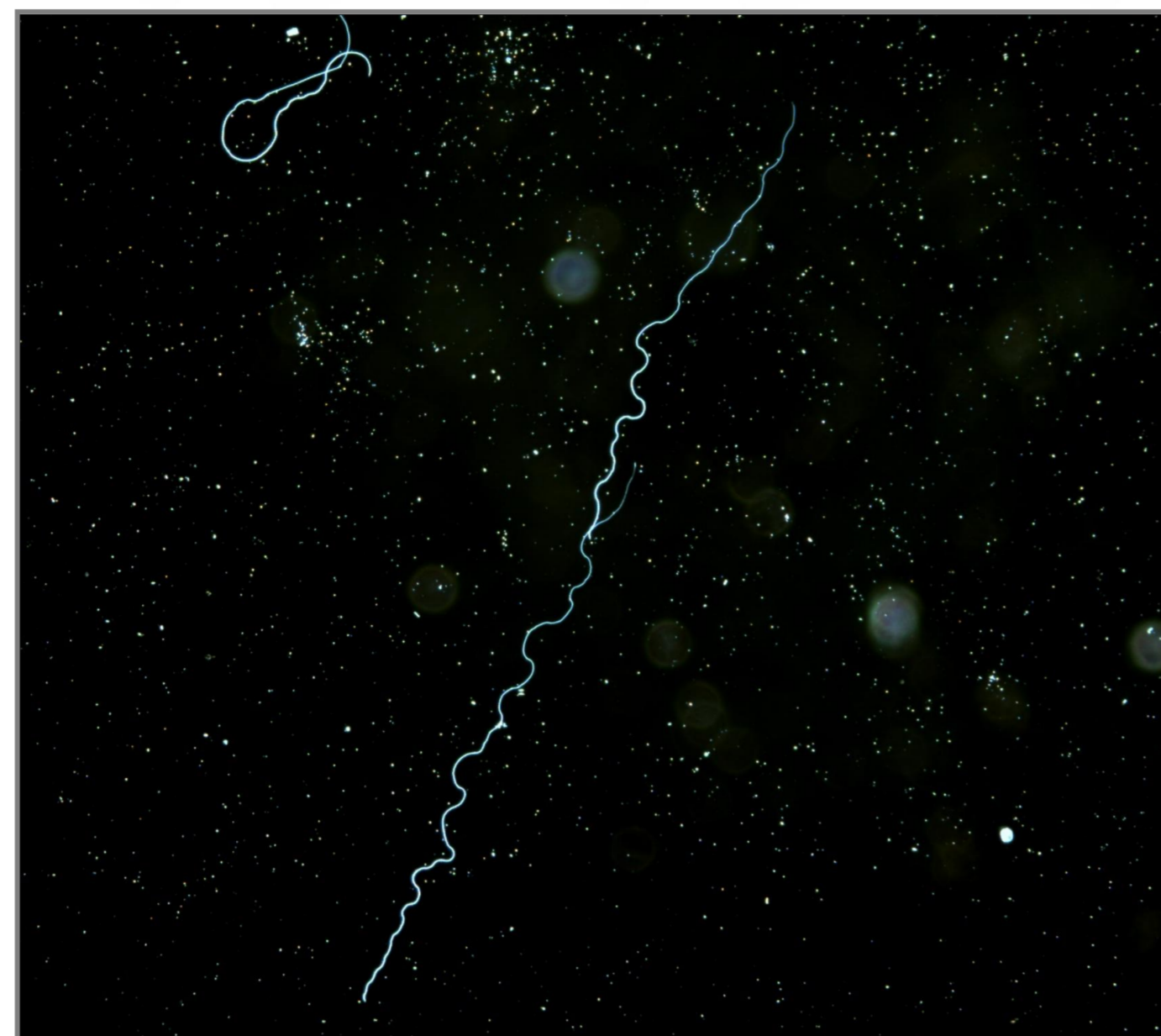
# Genotype-by-environment effects of dietary yeast on sperm traits in *Drosophila melanogaster*

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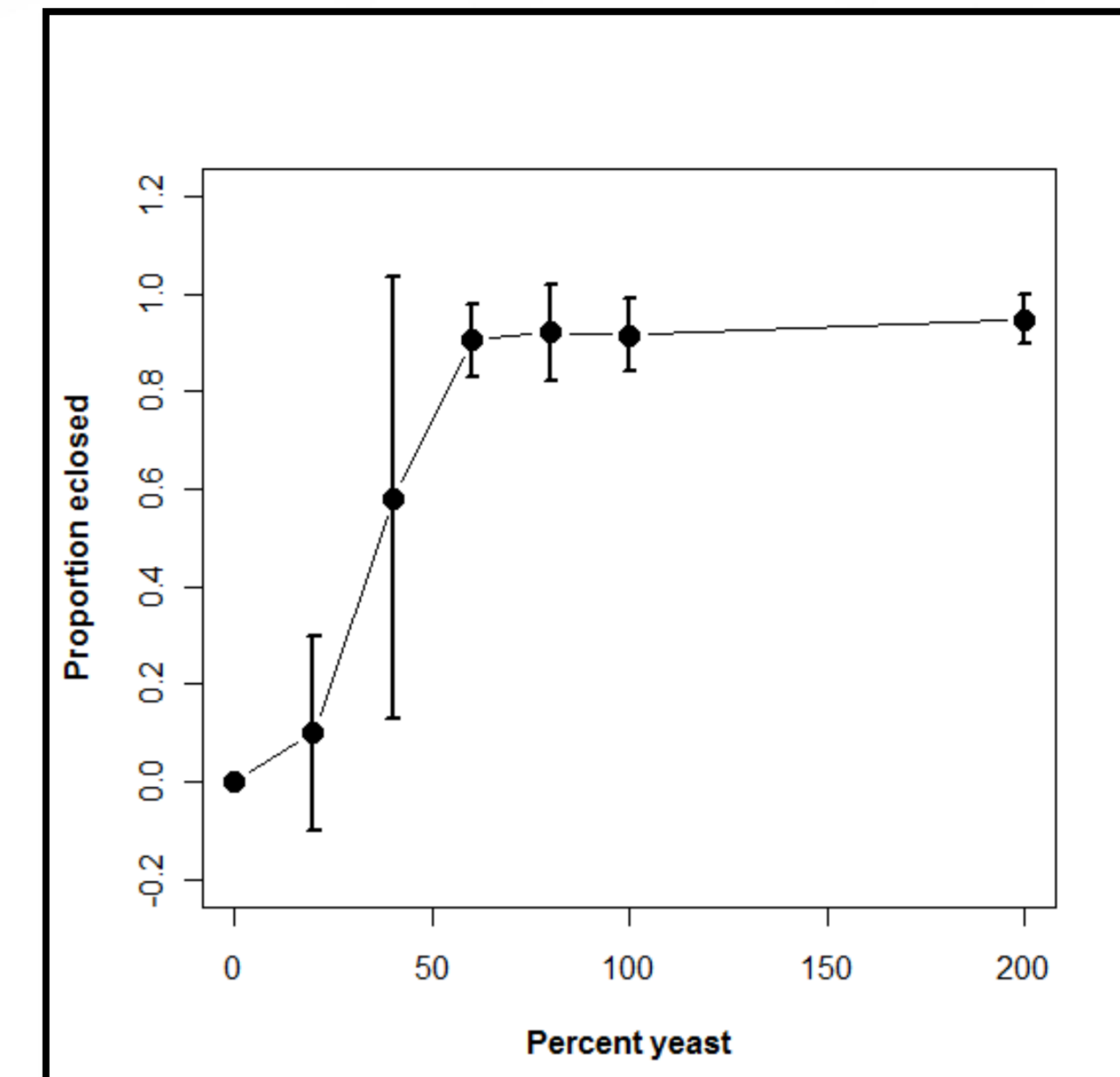
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## Introduction

Most current models of the evolution of sperm function and sperm morphology assume that changes are the result of a response to sexual selection (e.g. Immler et al. 2011). Because of this, sperm studies (whether comparative, quantitative genetic, or using experimental evolution) usually involve the measurement and/or manipulation of sexual selection parameters (e.g. Friberg et al. 2005). However the abiotic environment can also play an important role in influencing sperm physiology and morphology (Amitin & Pitnick 2007, Morrow et al. 2008). Because of this, we decided to carry out an investigation of the effect of dietary protein on sperm traits in *Drosophila melanogaster*. For this study we are using a set of sequenced inbred lines provided by the *Drosophila* Genetic Reference Panel.



A *Drosophila melanogaster* sperm. *D. melanogaster* have sperm that are approximately 1.8 mm long (300 times longer than a human sperm) which makes them easy to see under modest levels of magnification.



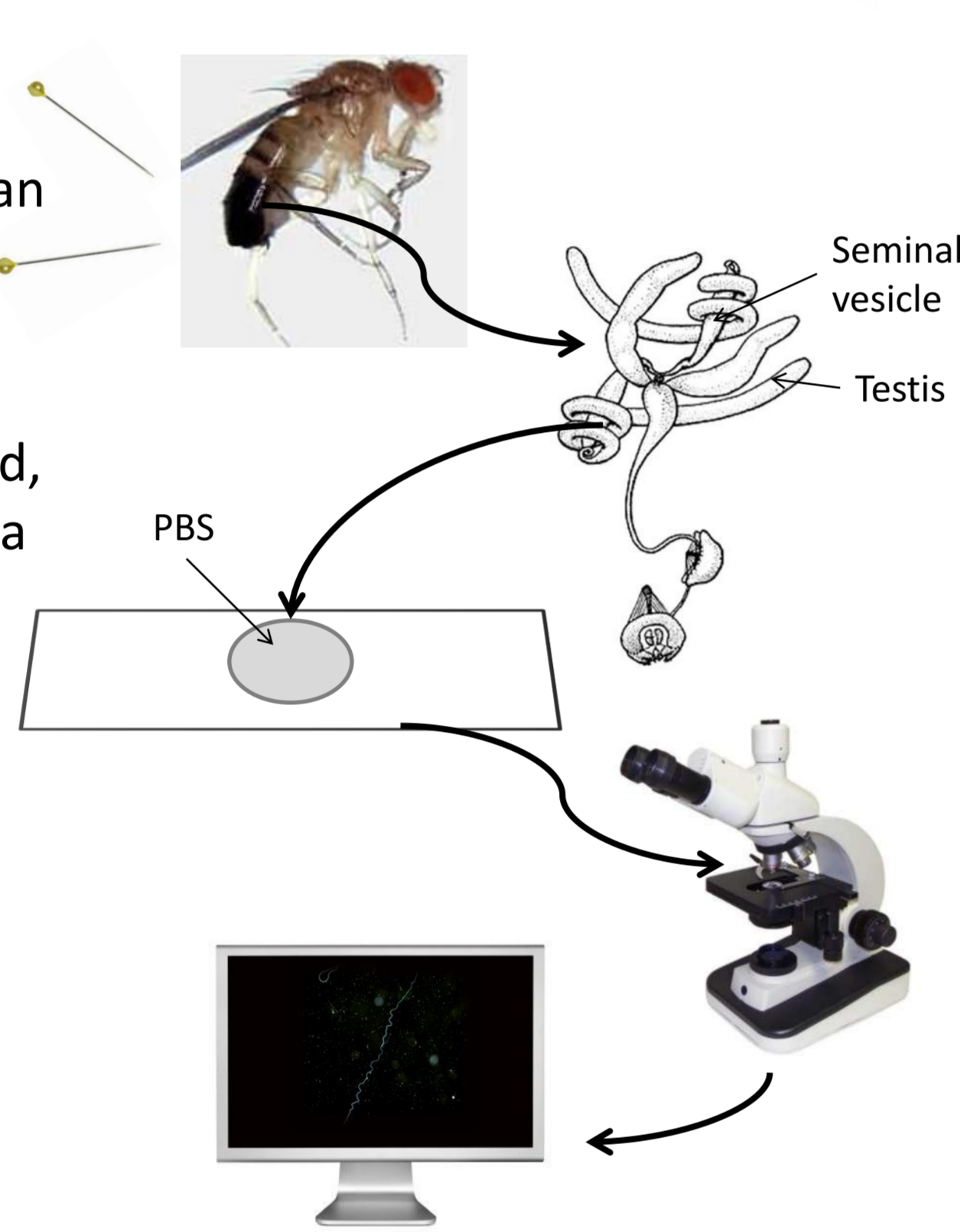
A pilot study using various yeast levels determined that development time increased at yeast levels <60%. Variance in the number of enclosed flies on day 12 was highest at 40% yeast, so this level was selected for the experiment. Error bars denote SDs.

## Project plan

1. The reproductive organs are dissected out of an adult male.

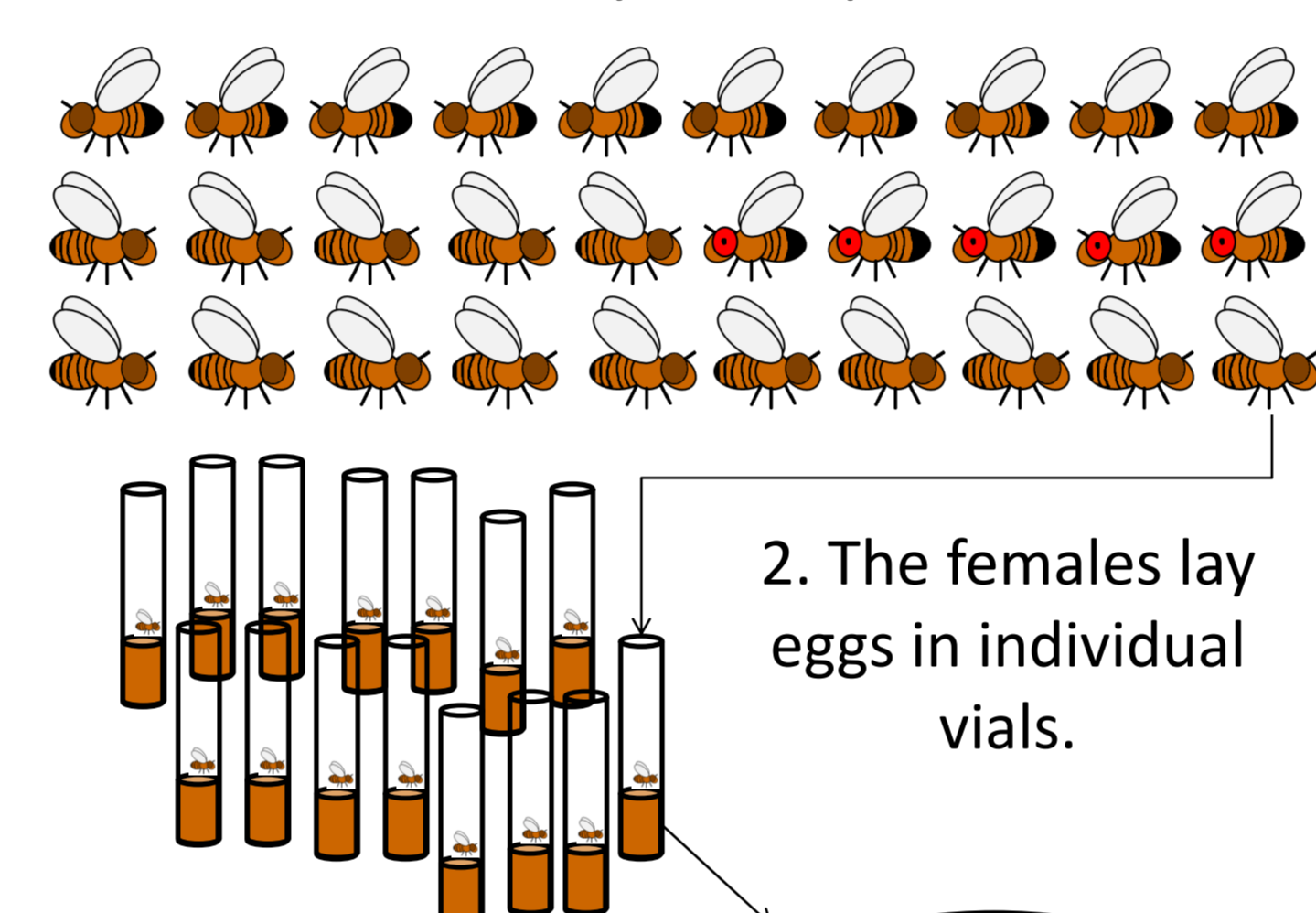
2. The seminal vesicle is removed, and opened into a drop of PBS on a slide.

3. The slide is processed and sperm are photographed through a microscope.



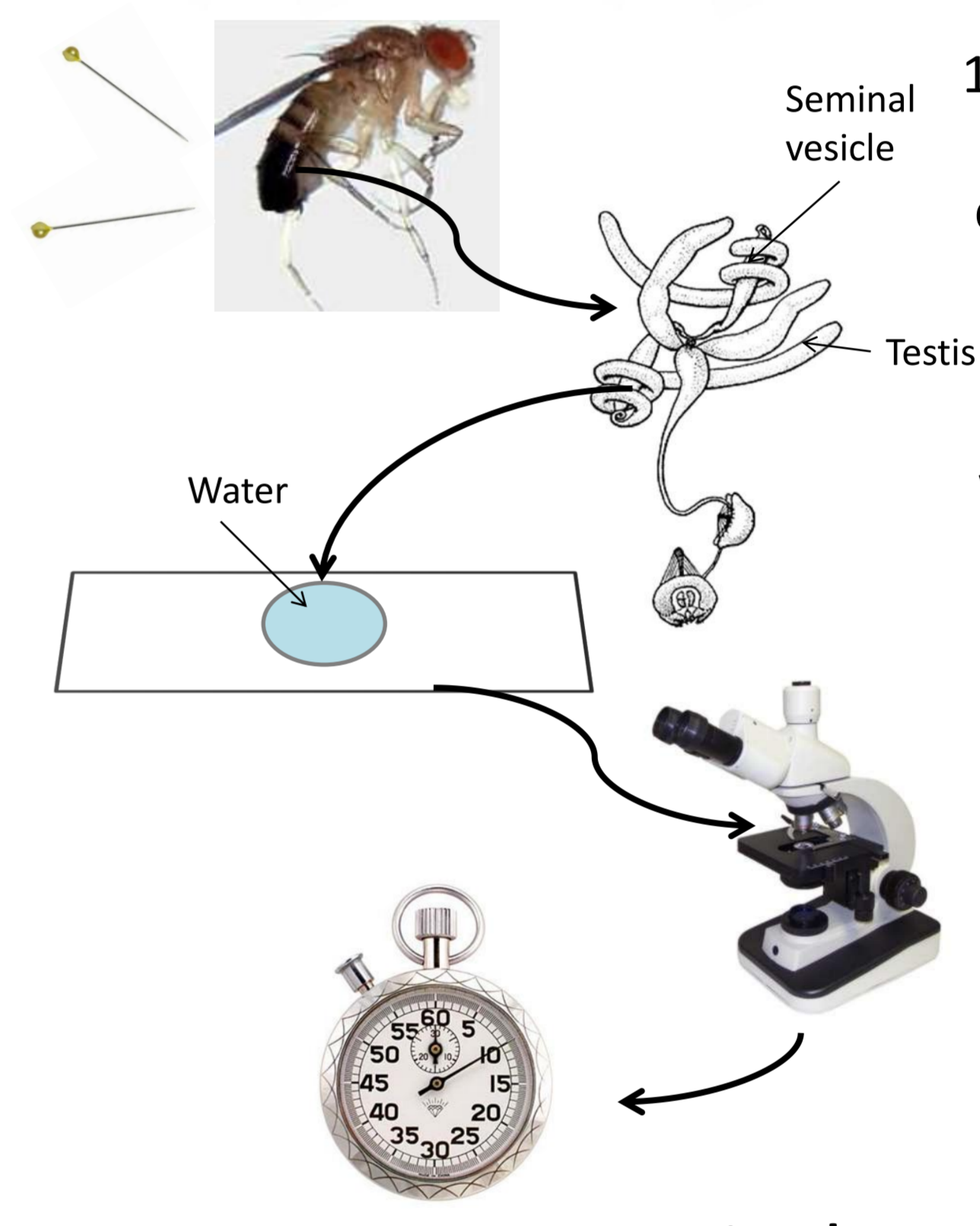
Sperm length

1. 5 red-eyed target males compete for matings with 10 brown-eyed competitor males.



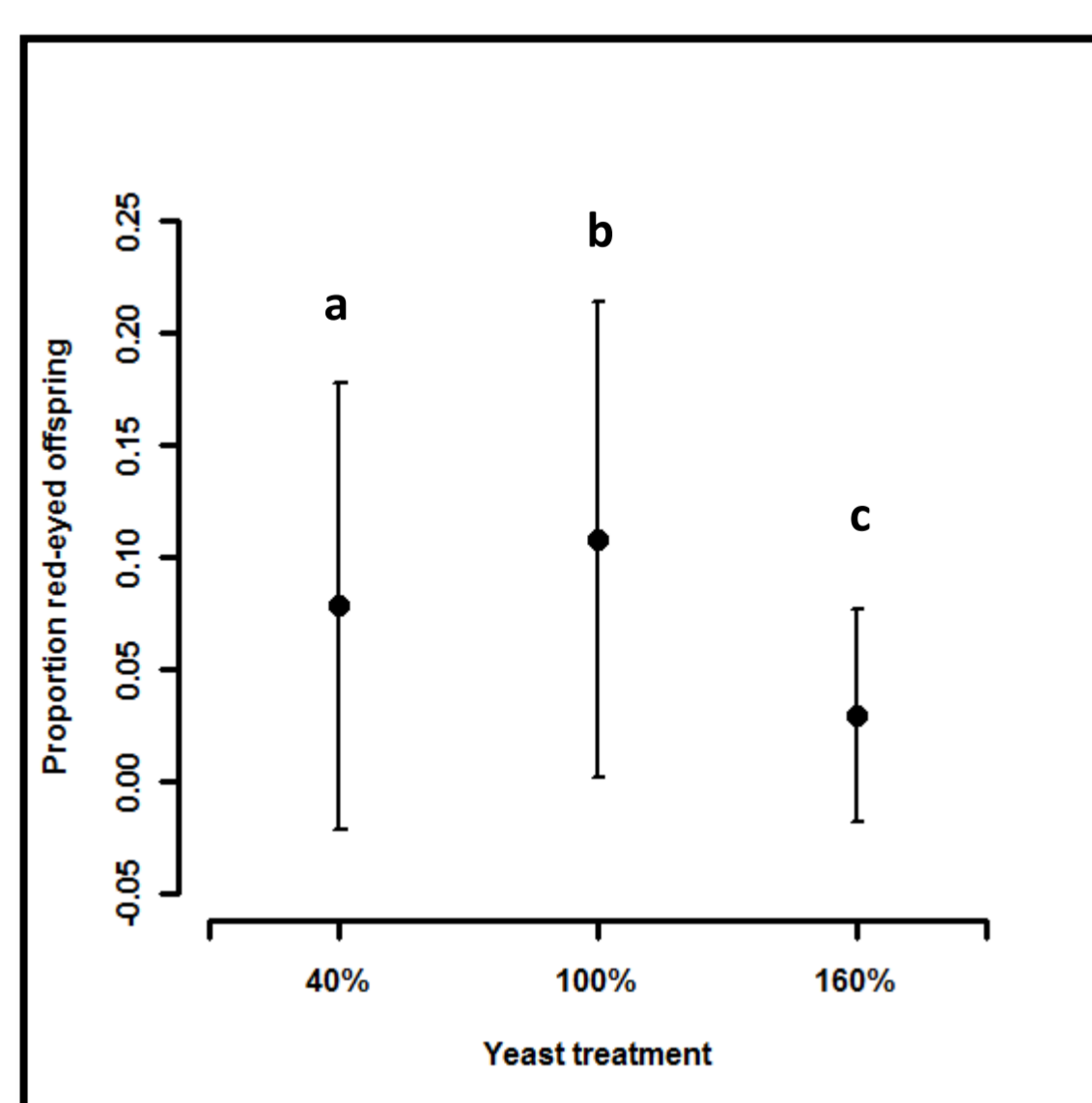
3. The proportion of adult red-eyed offspring in each clutch is measured.

Male fitness

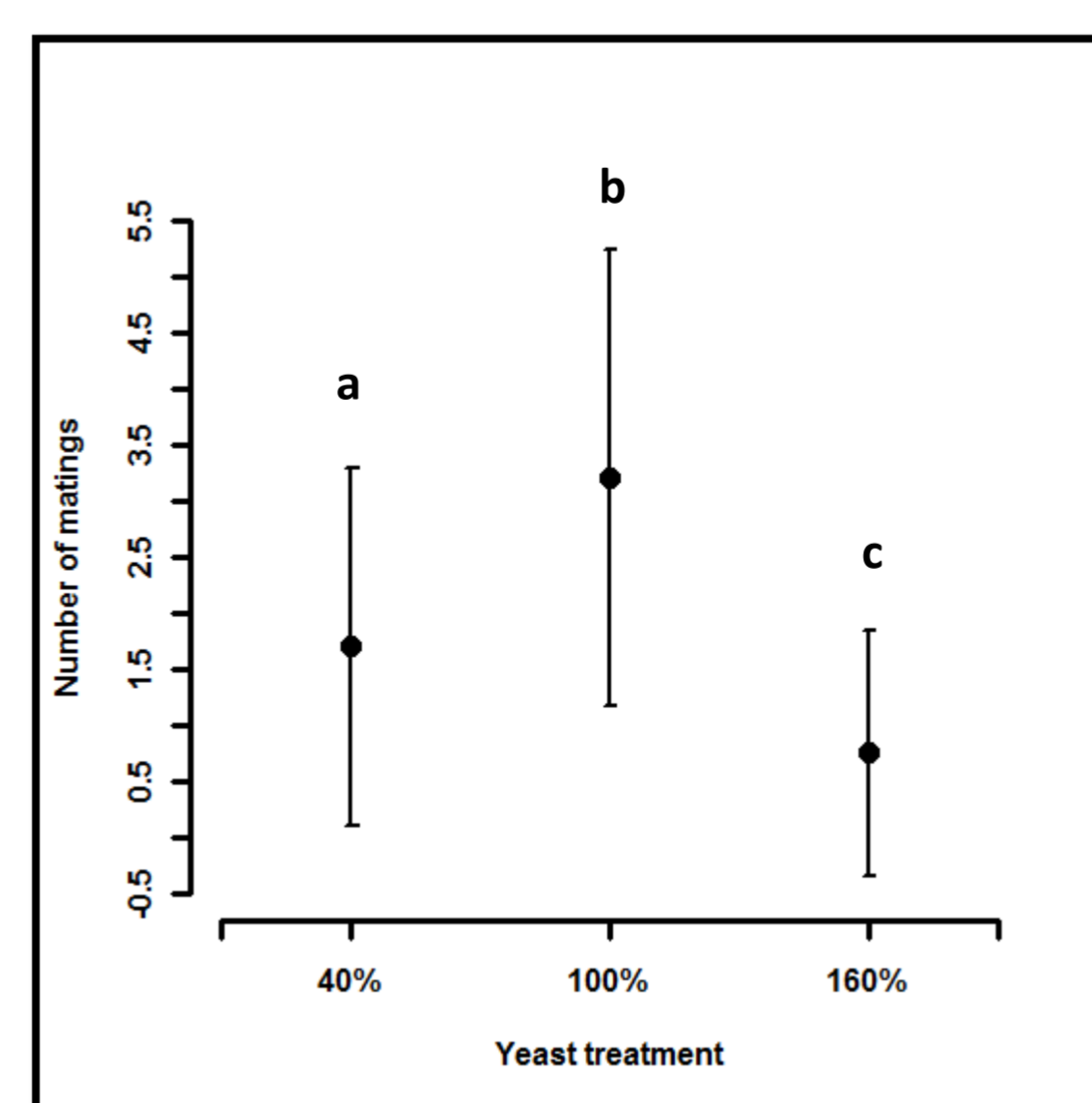


Sperm survival under osmotic stress

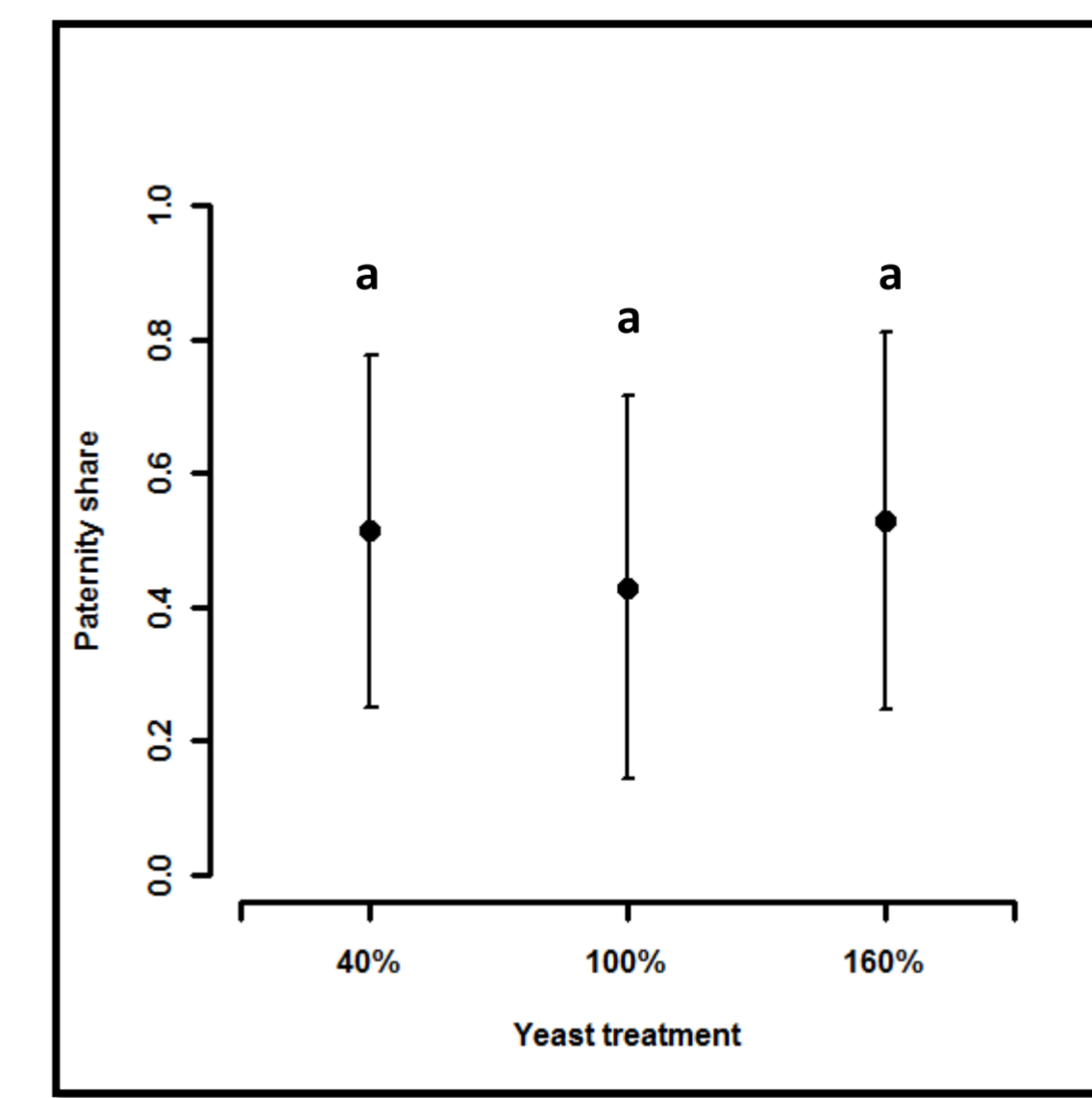
## Preliminary results



The proportion of red-eyed offspring was significantly affected by yeast treatment ( $P < 10^{-10}$ ). The large standard deviations (error bars) are the result of overall differences between the inbred lines ( $P < 10^{-15}$ ). The interaction between yeast treatment and line trended towards significance ( $P < 0.1$ ).



The number of matings obtained by the target males was significantly affected by yeast treatment ( $P < 10^{-15}$ ). Again there were large differences between the lines ( $P < 10^{-15}$ ). The interaction between yeast treatment and line was not significant ( $P > 0.1$ ).



The paternity share obtained by target males showed a trend towards a significant effect of yeast treatment ( $P < 0.1$ ). However there was no significant effect of inbred line and no interaction. More detailed analysis should help resolve whether this is a true effect or not.

## Conclusions

- As expected from previous studies on longevity (e.g. Pum Lee et al. 2008), increased yeast is not necessarily beneficial. Increased yeast resulted in reduced fitness and attractiveness in males.
- Decreased yeast was also deleterious, resulting in decreased fitness and attractiveness, but less so than in the increased yeast treatment.
- There were highly significant differences between lines, consistent with a substantial genetic component to the traits that were measured.
- Currently no indication of significant differences in plasticity among lines, although sampling is not complete.
- Some suggestion of a trade-off between attractiveness and performance in sperm competition. This requires further investigation.



## References

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