Diet significantly affected starvation resistance (as defined by time until death with no access to food) among an **experimentally evolved** population of *D.*

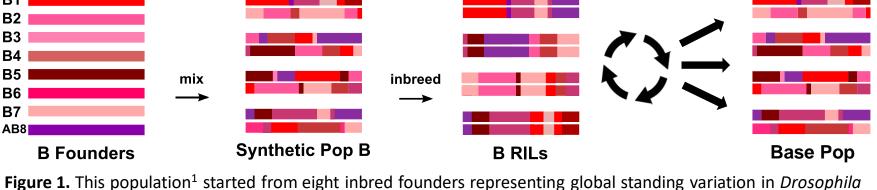
melanogaster.

Dietary impact on starvation resistance in an evolved multiparent population of *Drosophila melanogaster* Authors: ZF Elkins, Jordyn Moaton, Libby King

Division of Biological Sciences, University of Missouri

The base population

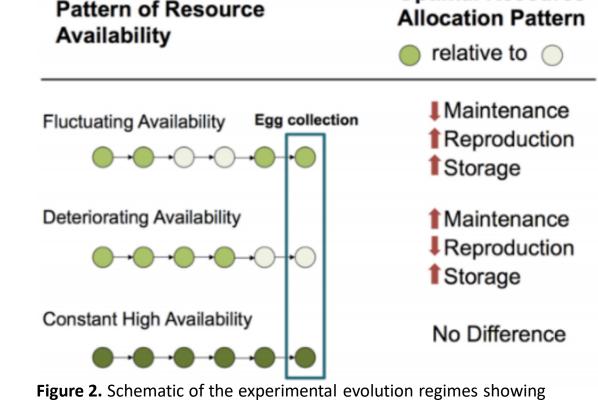
B1



melanogaster. They were mixed en masse for 50 generations and about 800 recombinant inbred lines (RILs) were inbred from this population. The RILs were then re-mixed to create a base population for experimental evolution.

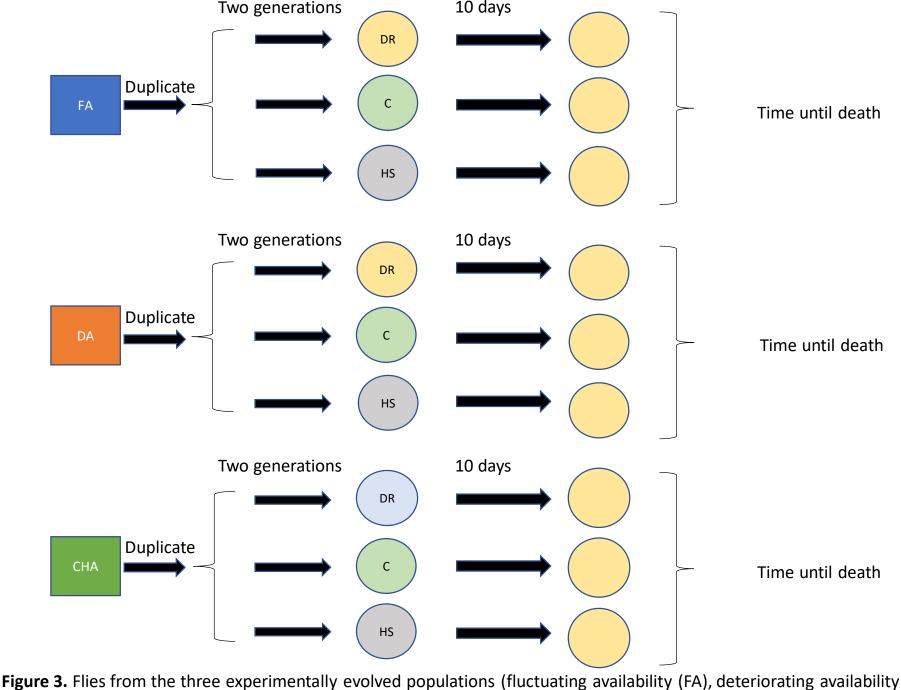
Optimal Resource

Methods



resource allocation. Dark green circles indicate high sugar food, green circles indicate standard food, and light tan circles indicate low yeast Starvation resistance assay

changing diets and predictions for the evolved pattern of plasticity in



324; n_{vigl} = 10) for 10 days before being placed on nutrition-less agar. Starvation resistance was quantified as time until death starting from the moment flies were placed on nutrition-less agar. Results Survival probability over time predicted by diet Strata + diet=C + diet=DR + diet=HS

p = 0.00095

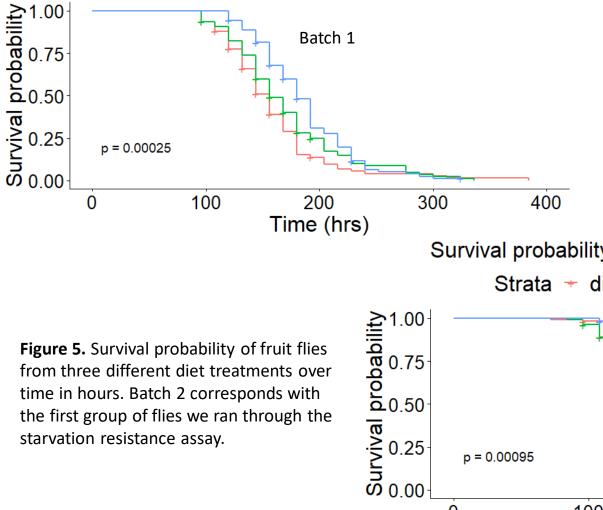
CH

300

200

(DA), and constant high availability (CHA)) were duplicated and fed a standard diet for two generations. The following generation was placed on three different diets (dietary restriction (low yeast), control, and high sugar; $n_{Batch1, Batch2}$ = 323,

Batch 1



Ċ

Survival probability over time predicted by diet Strata + diet=C + diet=DR + diet=HS

Figure 4. Survival probability of fruit flies from three different diet treatments over time in hours. Batch 1 corresponds with the first group of flies we ran through the

starvation resistance assay.

Batch 2

Time to starvation in Batch 1 of old flies Time to starvation (hours) CH

DR

Figure 7. Each vial's time until starvation

separated by diet treatment in the second batch of flies. Diets include control (C), low

C

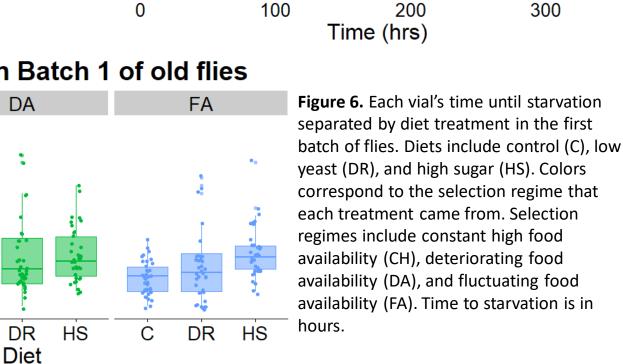
HS

starvation resistance assay.

300

200

100



Time to starvation in Batch 2 of old flies

FA

0.112

0.365

0.725

< 0.001

<0.001 ***

Fime to starvation (hours) yeast (DR), and high sugar (HS). Colors correspond to the selection regime that each treatment came from. Selection regimes include constant high food availability (CH), deteriorating food availability (DA), and fluctuating food availability (FA). Time to starvation is in hours. Extra figures C (N=108) diet reference DR (N=108) 0.72 (0.54 - 0.96)

HS (N=108)

CH (N=108)

DA (N=108)

FA (N=108)

(N=162)

0.49 (0.37 - 0.66)

reference

1.13 (0.84 - 1.50)

1.25 (0.93 - 1.68)

reference



Sex

sellD

3.31 (2.50 - 4.39) M (N=162) < 0.001 # Events: 282; Global p-value (Log-Rank): 1.6751e-17 AIC: 2616.66; Concordance Index: 0.69 Hazard ratio C (N=109) diet reference DR (N=108) 0.80 (0.60 - 1.06) HS (N=108) 0.54 (0.41 - 0.71) **Figure 7.** Cox proportional CH (N=108) hazard model for time sellD reference until death due to DA (N=108) 1.14 (0.86 - 1.52) starvation in flies from Batch 2 FA (N=109) 1.05 (0.79 - 1.40) F (N=162) Sex reference (N=163) # Events: 300; Global p-value (Log-Rank): 2.1882e-36 0.5 AIC: 2694.53; Concordance Index: 0.76

Future directions

 Incorporate genetic data in analysis Search for regions of the genome contributing to higher starvation resistance Use previously published lipid content data to compare starvation resistance with lipid composition of

SCAN ME

flies within the three selection regimes References

Acknowledgements

¹ http://FlyRILs.org

This work is supported by the University of Missouri Life Sciences Fellowship, and by NIH grant R01 GM117135 to E.G.K. We would also like to thank Elizabeth Jones for lending technical expertise and time that aided in conducting this experiment.