

Using the fruitfly, Drosophila melanogaster to Understand the Genetic Basis of Aging

Dr. Jeff Leips¹, Dr. Michelle Starz-Gaiano¹, Dr. Peter Abadir, MD²

¹Dept of Biological Sciences, University of Maryland Baltimore County, Baltimore MD 21250 ² Johns Hopkins Bayview Medical Center, Baltimore, MD 21224

Introduction

- All organisms deteriorate with age
- Aging **decreases the quality of life** of the elderly
- In human populations, individuals differ in rates of physiological decline with age
- Individual differences in aging is partially attributable to genetic differences among individuals
- The actual genes that are responsible for these differences among individuals are largely unknown and so limiting our ability to design treatments

Why Use Drosophila?

- Flies show similar age-related declines to that of humans
- decline in **physical strength** with age
- decline in walking speed and endurance with age
- decline in **ability to fight infection** with age
- Many fully sequenced genomes, - > 60% of fly genes shared with humans
- Efficient and Economical Model Organism: - Short life span (~50 days) allows measurement of genetic
- influences on age-related decline across entire life span
- Inexpensive to observe and maintain large populations (maximizes statistical power to detect genetic effects)
- **Genomic mapping techniques** to identify genes contributing to variation in aging **well developed** - 1000's of genetic resources allow **experimental manipulation** genes to validate genetic effects on aging
- Ideal organism for training the next generation of scientists



Drosophila as a model to understand the genetics of age-related decline in immune function







• Age-related decline in immunity places a substantial burden on the healthcare system: infection-related hospital admission is among the most costly and the most common in the elderly¹

Innate	immune	response	2
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- Two main components - Phagocytosis - Production of **Antimicrobial Proteins**
- Conservation of Genes/Signaling Pathways in flies and humans

Measuring the Immune Response: Infection Assay

Microinject Bacteria

Measure Infection Level 24 hours after infection



Effect of aging on the ability to clear infection differs among genotypes

Age (in weeks)

Genome wide expression experiment identified ~ 250 genes that contribute to individual variation in age -specific ability to fight infection,



- in humans
- Genotypes treatment
- to individual differences

References

Review, 2004. 3: p. 1-14.

Funding

This research was supported by National Institutes of Health grants 1R03 AG023339-01 and 5R01 DK084219-02.

Using Drosophila to understand individual variation in response to drug treatment

• Individual differences in response to drug treatment is often related to genetic differences among individuals³

• We are examining the genetic basis of the response to Lisinopril (a commonly prescribed drug to treat age-related changes in blood pressure - drug not effective for all patients.

• Effects of Lisinopril Control treatment in flies generally mimics effects 0 20 40 60 80 100 120 **RAL_229** differ in their response to drug 80 100 120 20 40 60 Mapping efforts Day underway to identify RAL_304 genes that contribute Control in drug response. T reatm ent 80 100 120 40 The effect of Lisinopril on A) Short Dav distance climbing speed, B) Long Lisinopril treatment extends life span distance climbing speed depended on in genotype specific manner A: genotype. C) Lisinopril improved RAL_73, B: RAL_229, C: RAL_304 strength in all lines.

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