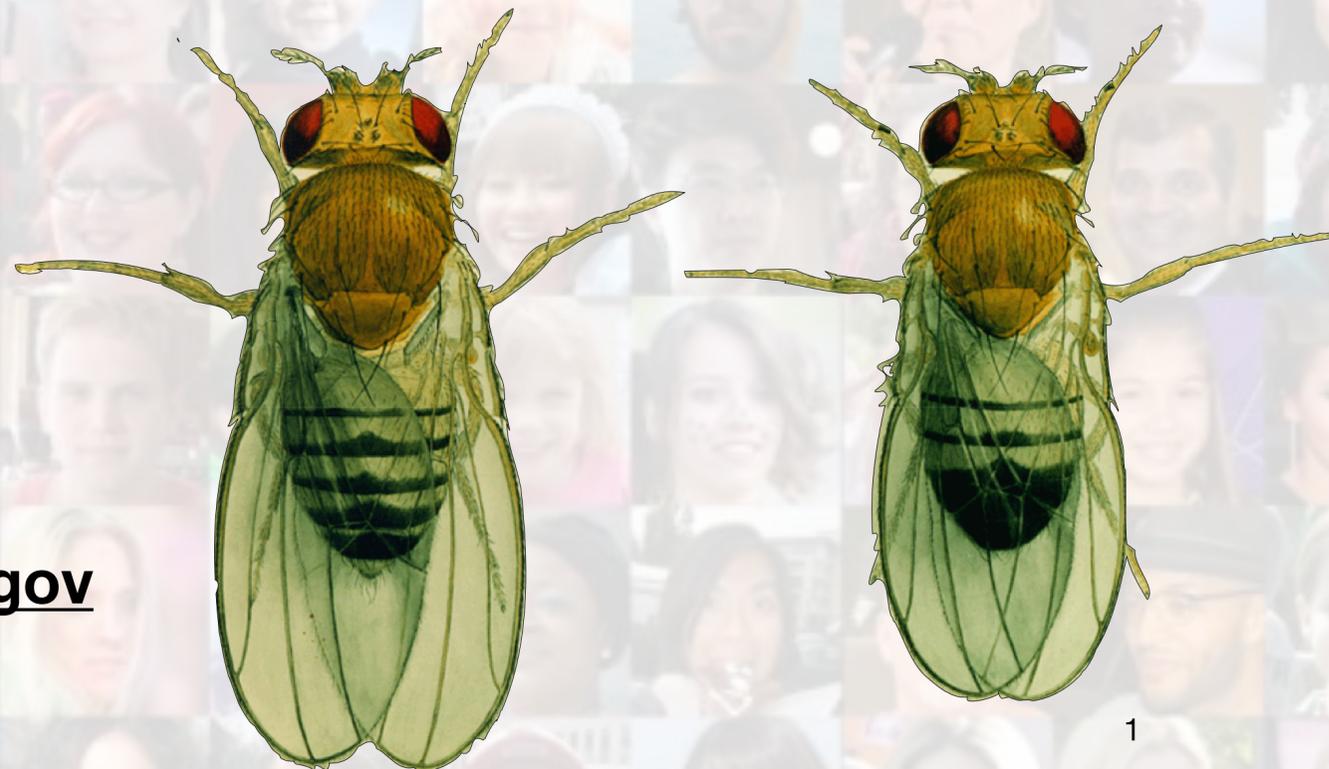


# Comparative Genomics for Precision Toxicology

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Disclaimer

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# **Registration, Evaluation, Authorisation and Restriction of Chemicals (REACH)**

**“No data, no market”**

# ASPIS: effects of chemicals on health

€60 million, 3 projects, 5 years



Prof. John Colbourne



Prof. Mathieu Vinken



Prof. Bob van de Water



Comparative

Expression &  
Metabolite Profiling

Initiating events

Adverse outcome



Data mining

Artificial Intelligence

Adverse outcome  
pathways

Focused *in silico* &  
*in vitro* testing



Human Cell lines

Pluripotent Stem  
Cells

Organs on a chip

Metabolism,  
Distribution, &  
Excretion

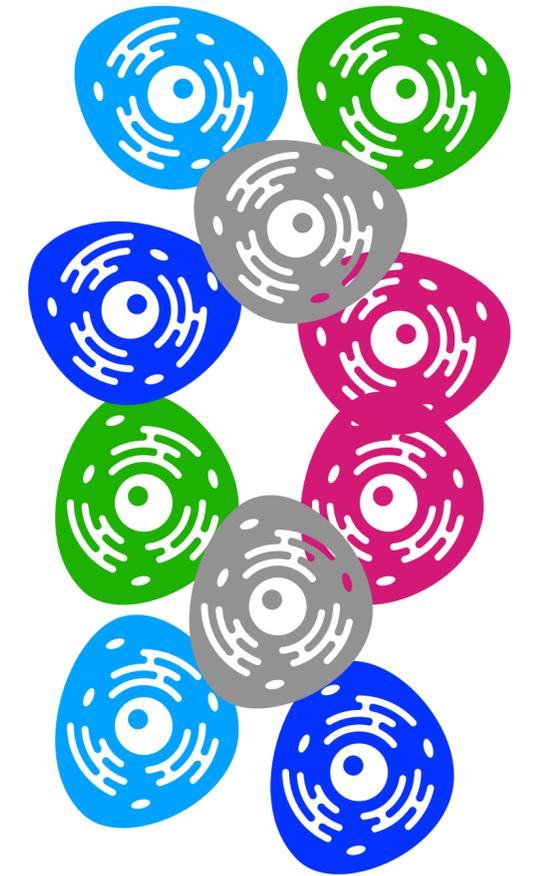
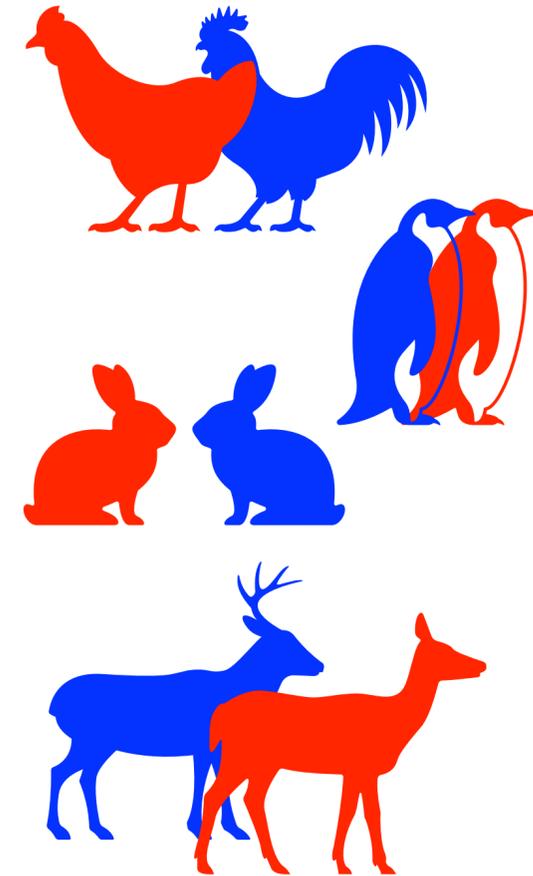
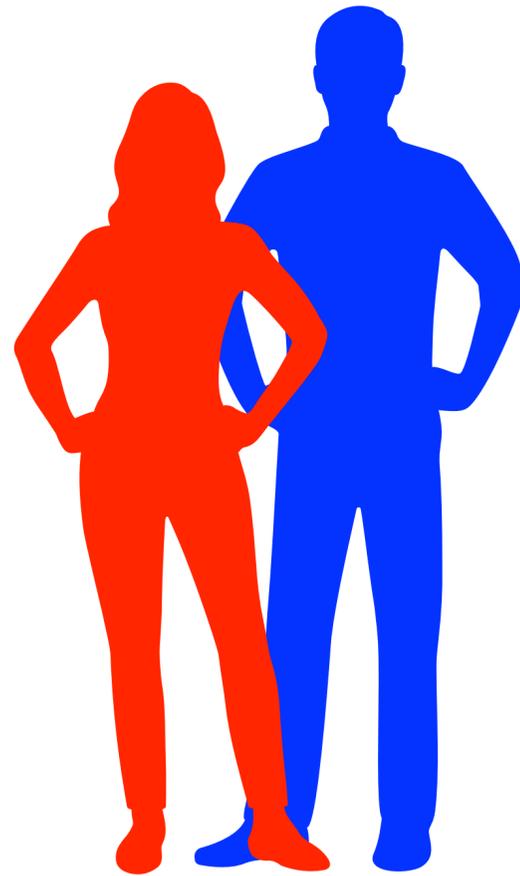
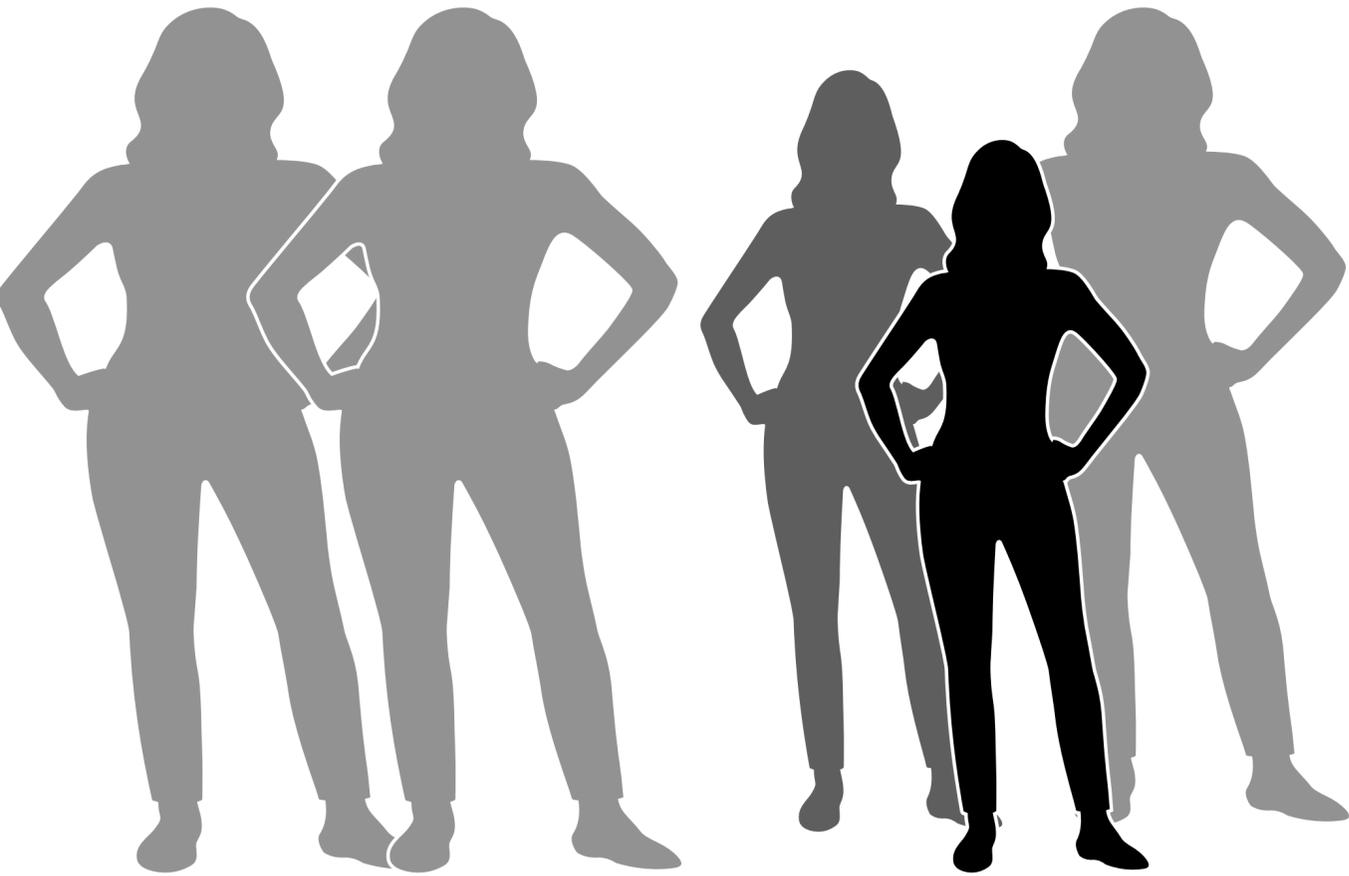
# ASPIS covers the NAMs



	Pathways	Tissues	Organs	Organ systems	Embryo > Adult	Complex behavior	Through-put	Expt. Control	Human Genetics	Ethics
Data Mining	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Low	Yes	Testing people retrospectively
Cells	Yes	No	No	No	No	No	Ultra	High	Yes	Origin of cells Reagents
On a chip	Yes	Yes	No	No	No	No	Yes	Can be Delicate	Yes	Origin of cells Reagents
organoids et al.	Yes	Yes	Yes	No	No	No	Moderate	Can be Delicate	Human	Origin of cells Reagents
Model Organisms	Yes	Yes	Yes	Yes	Yes	Yes	Moderate	High	Orthologs Humanization	Sentience

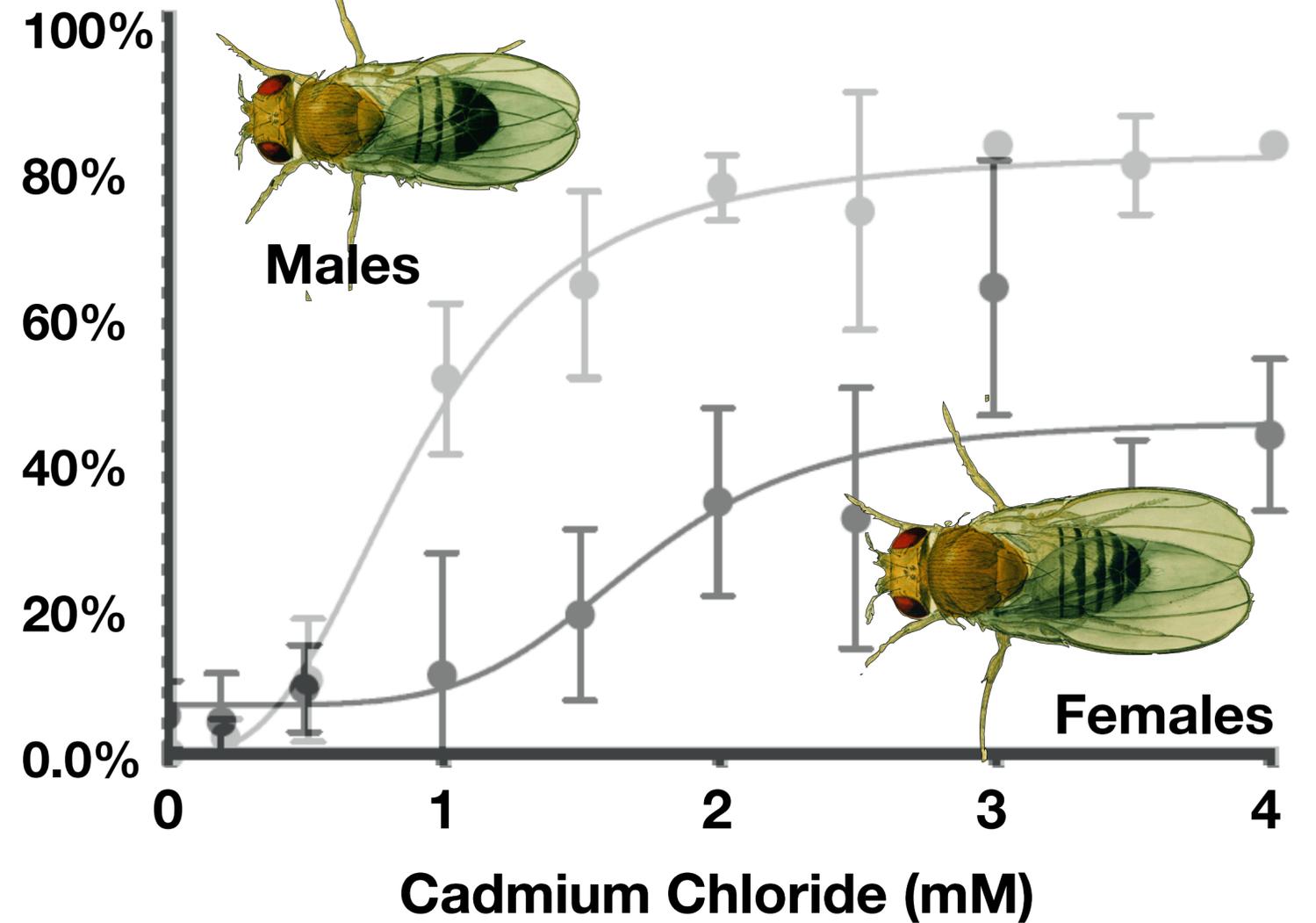
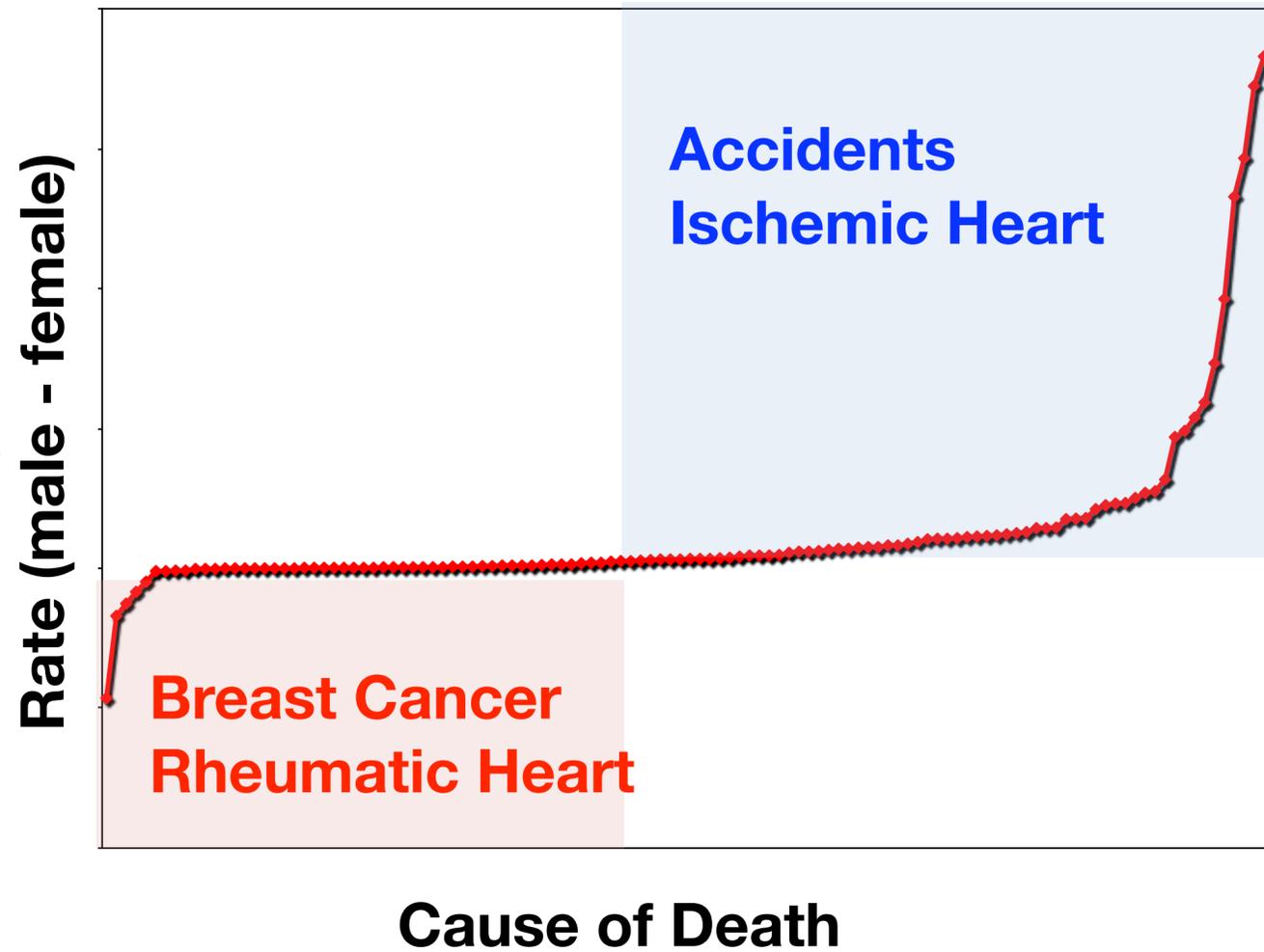
# Functional Variance:

**Genotype < Genotypes < Sexes < Species < Cell type**



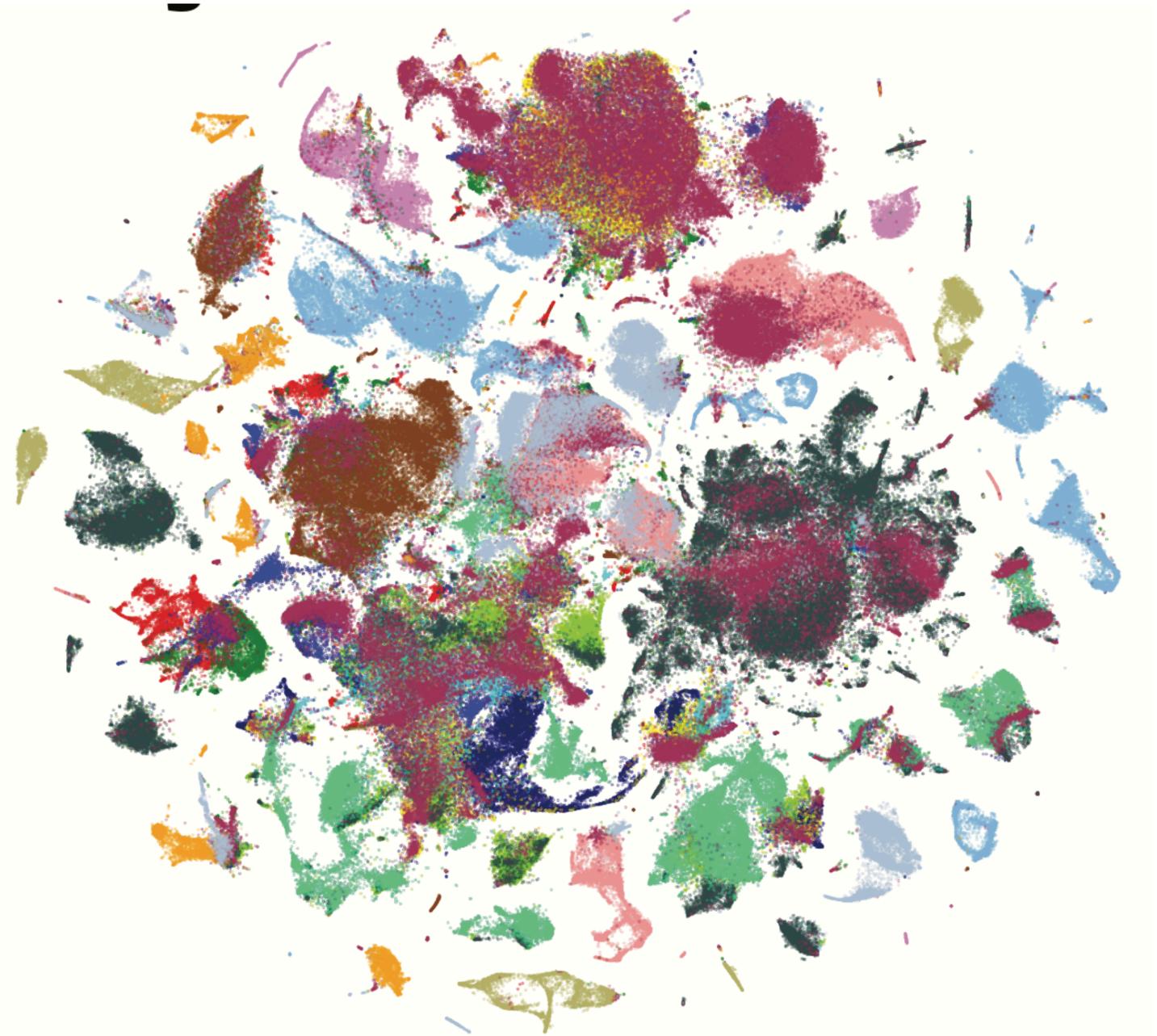
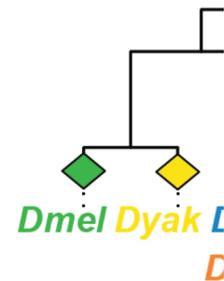
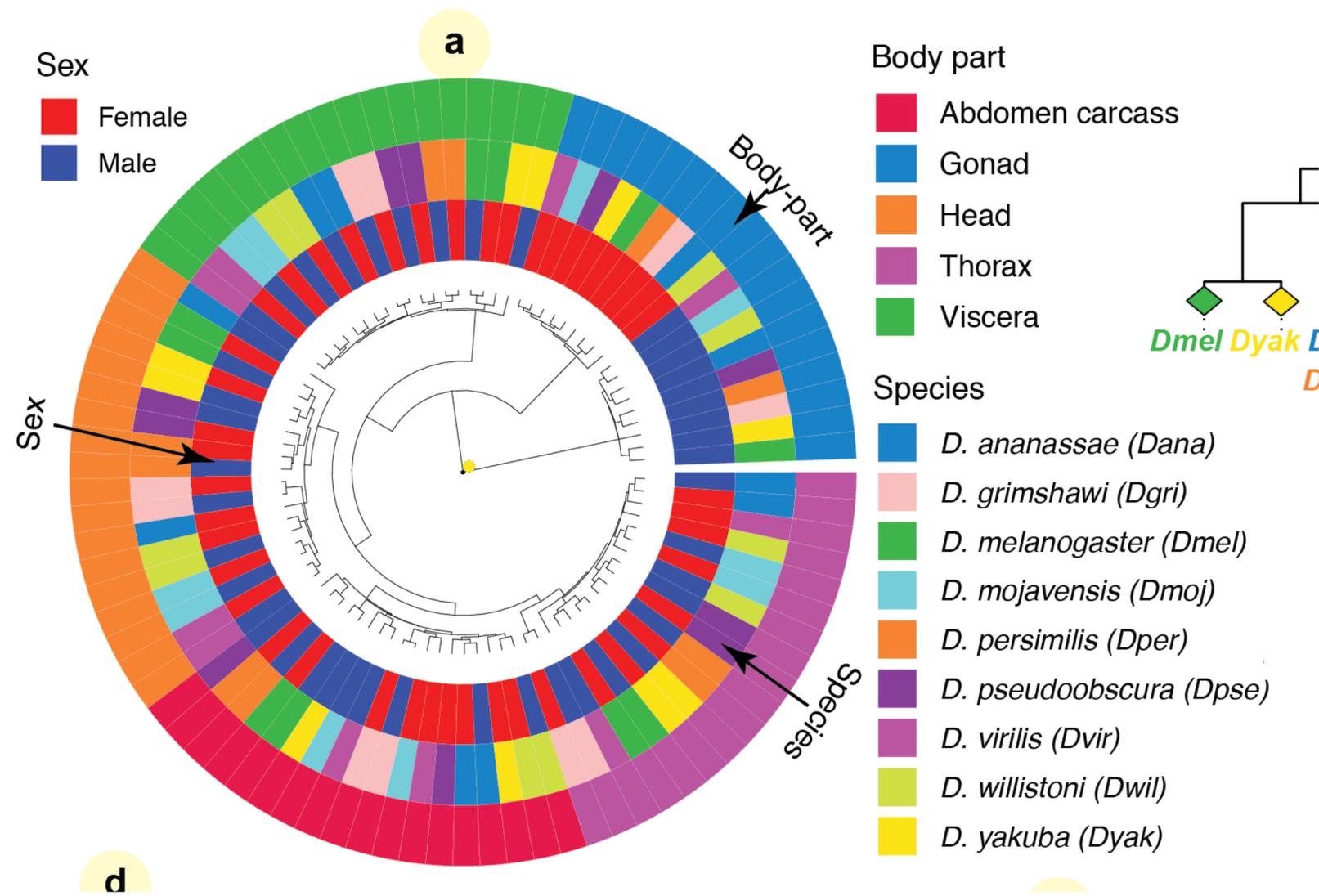
# Stratify by sex. Please...

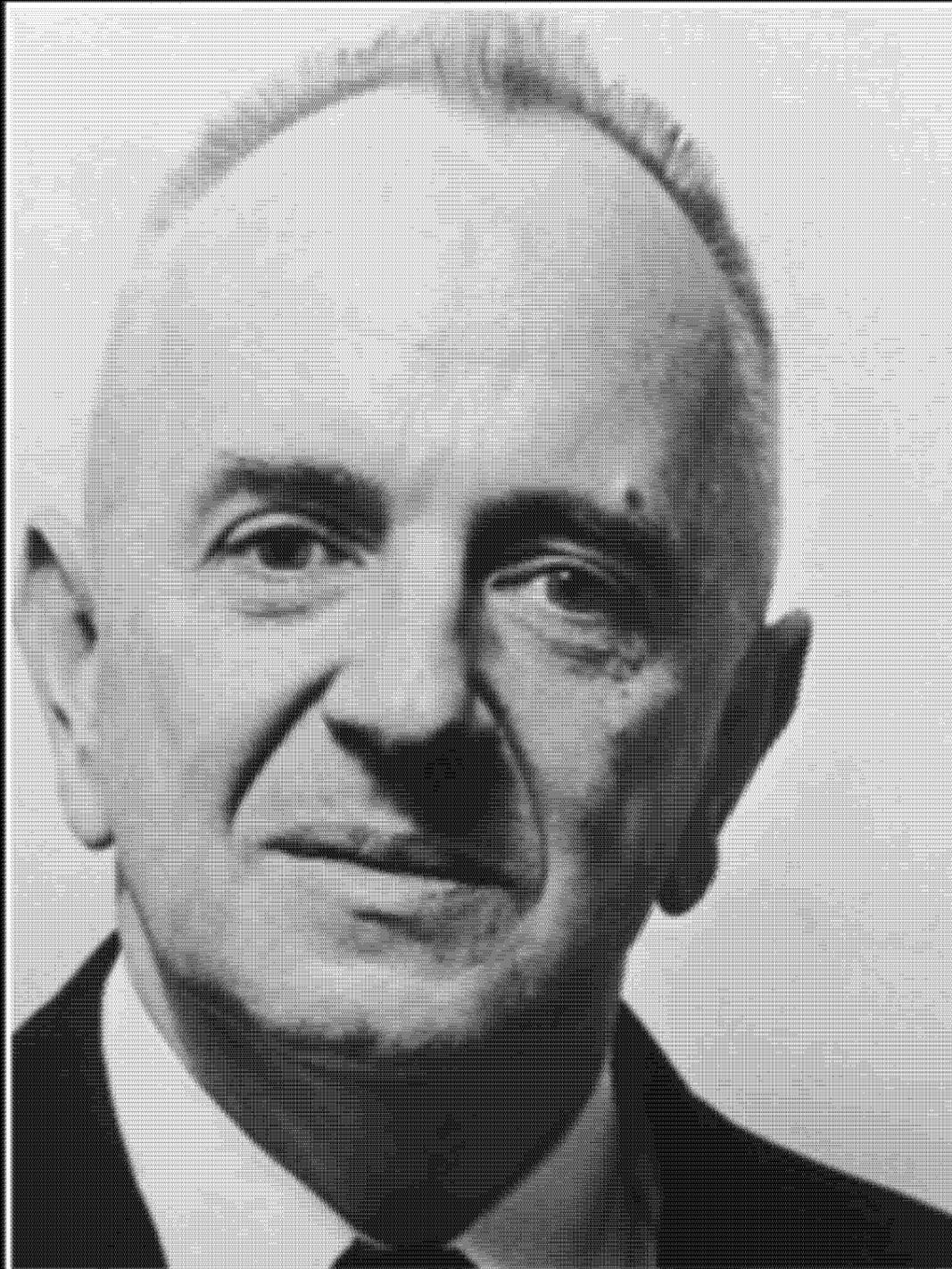
## USA Mortality 2006





# Organ/Cell expression differences are greater than those based on genetic diversity.





Nothing in biology makes sense  
except in the light of evolution.

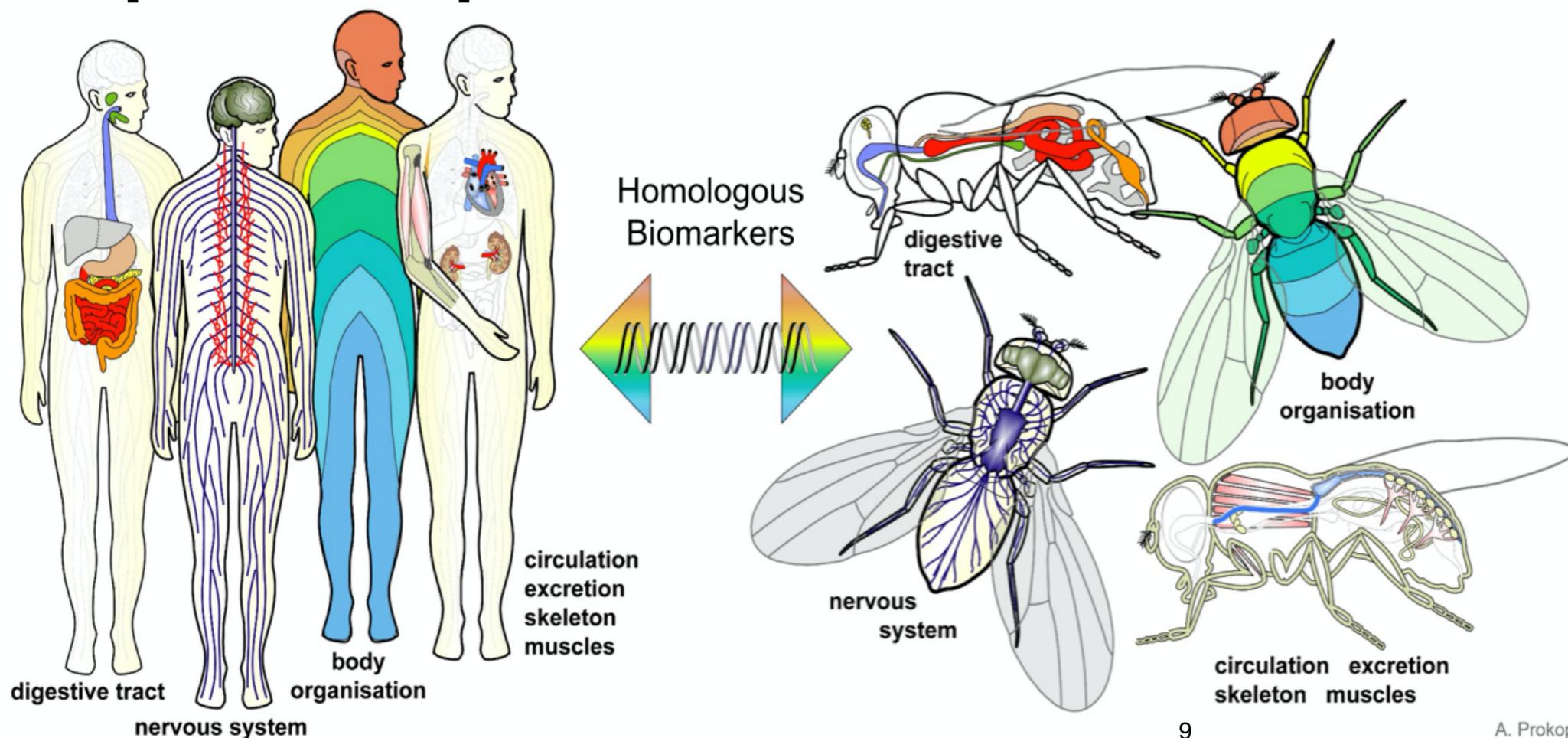
— *Theodosius Dobzhansky* —

AZ QUOTES



# Studying human variants in flies

- **85% of OMIM genes have fly homologs**
- **Genes and variants function cross-species in patient studies**



ATAD3A (Harel et al. 2016)

TM2D3 (Jakobsdottir et al. 2016)

EBF3 (Chao et al. 2017)

OGDHL (Yoon et al. 2017)

CACNA1A (Luo et al. 2017)

ARIH1 (Tan et al. 2018)

ATP5F1D (Oláhová et al. 2018)

WDR37 (Kanca et al. 2018)

MARK3 (Ansar et al. 2018)

DNM1L (Assia Batzir et al. 2019)

DROSHA (Barish & Senturk et al. 2022)

IQSEC1 (Chung et al. 2019)

OXR1 (Wang et al. 2019)

ACOX1 (Chung et al. 2020)

TOMM70 (Dutta et al. 2020)

CDK19 (Chung et al. 2020)

CAPZA2 (Huang et al. 2021)

RNF2 (Luo et al. 2021)

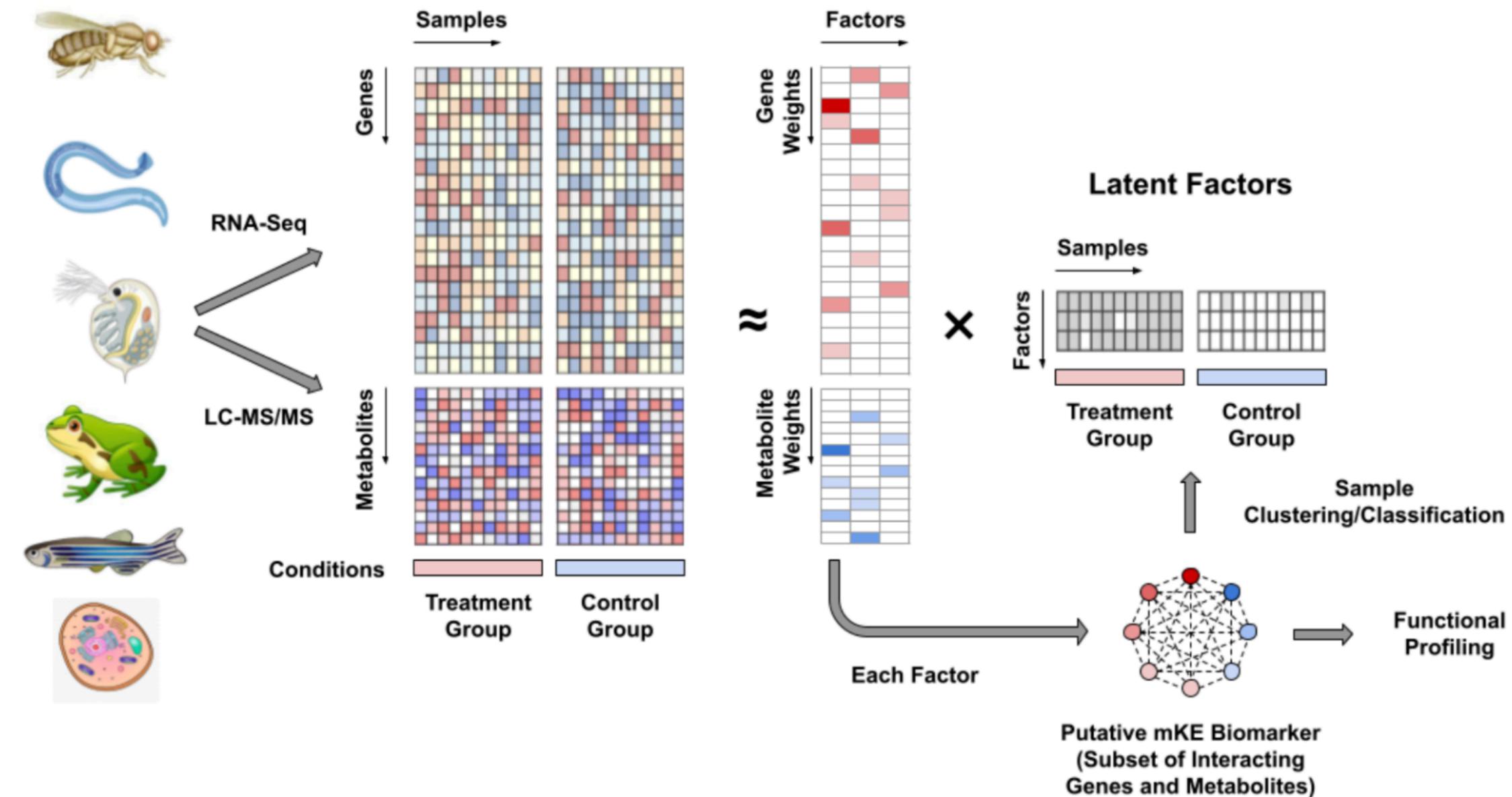
GBA1 (Wang et al. 2022)

TIAM1 (Lu et al. 2022)

FRMD5 (Lu et al. 2022)

SPTSSA (Srivastava et al. 2022)

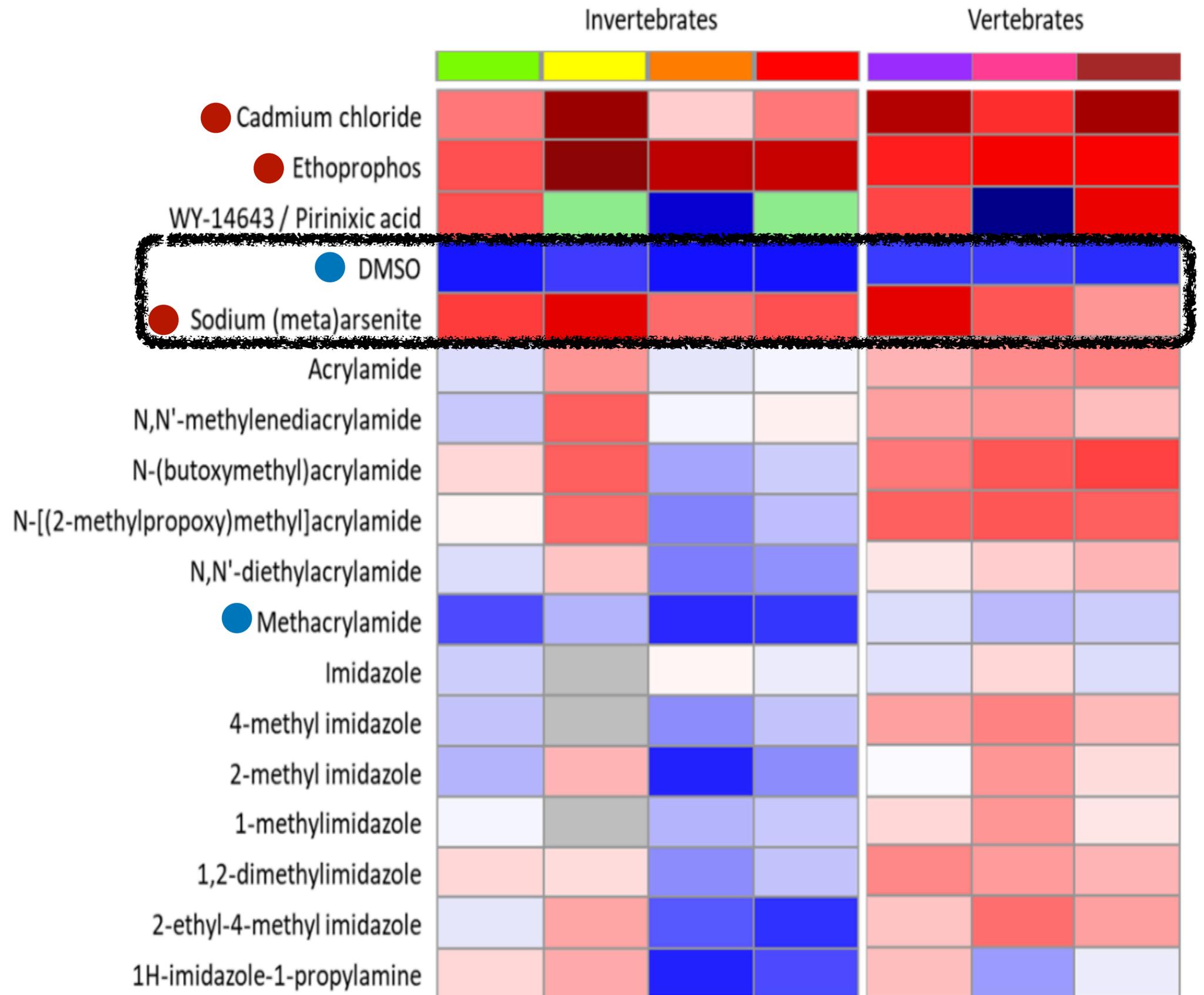
# Comparative Strategy



- Range finding to determine phenotypic dose responses (lethality & behavior differences)
- Time and dose normalized across species based on range finding
- Perform gene expression and metabolite profiling time-course
- Identify early, potentially initiating events, pathway modeling
- Testing models using directed genetics and populations

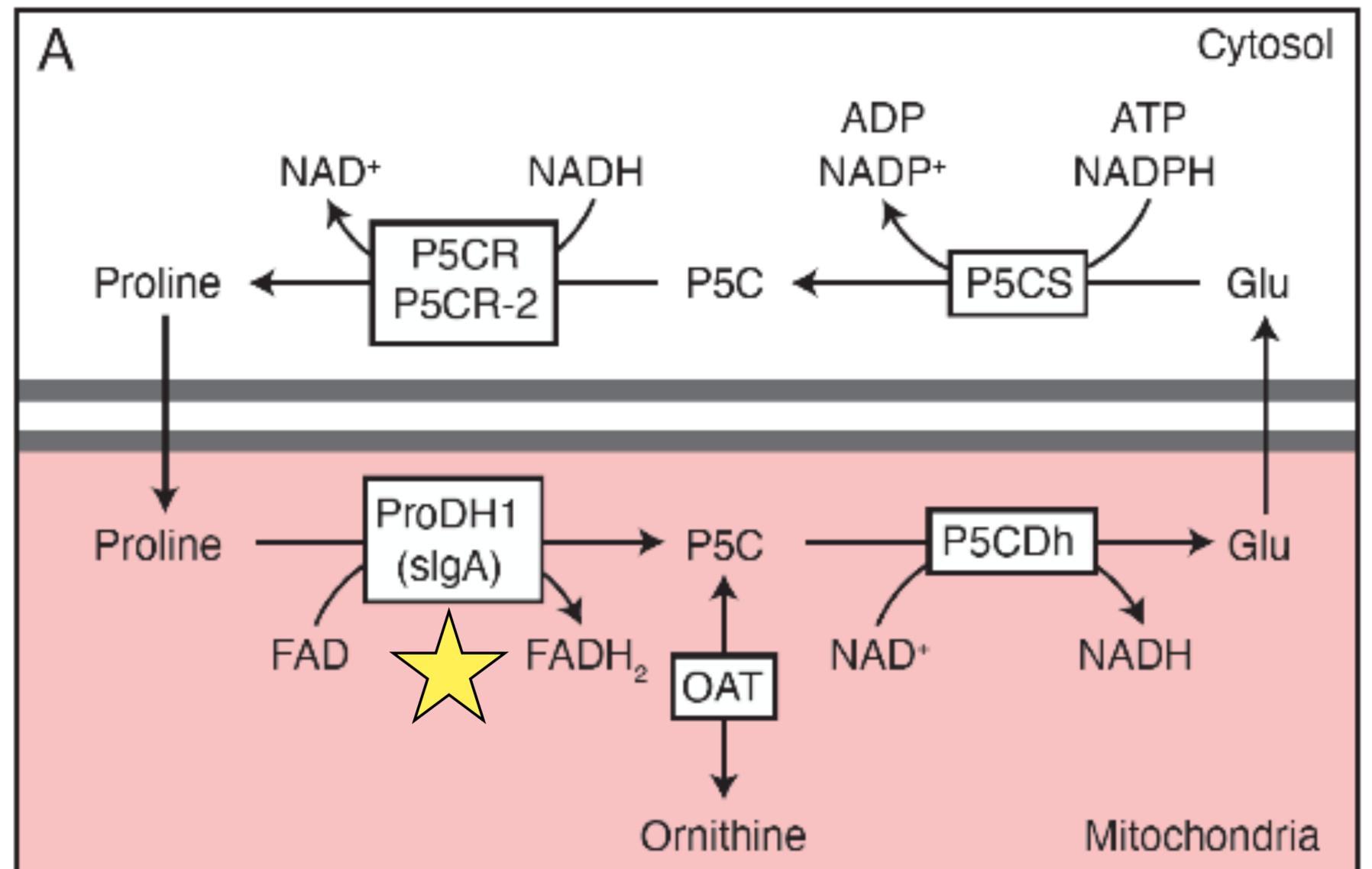
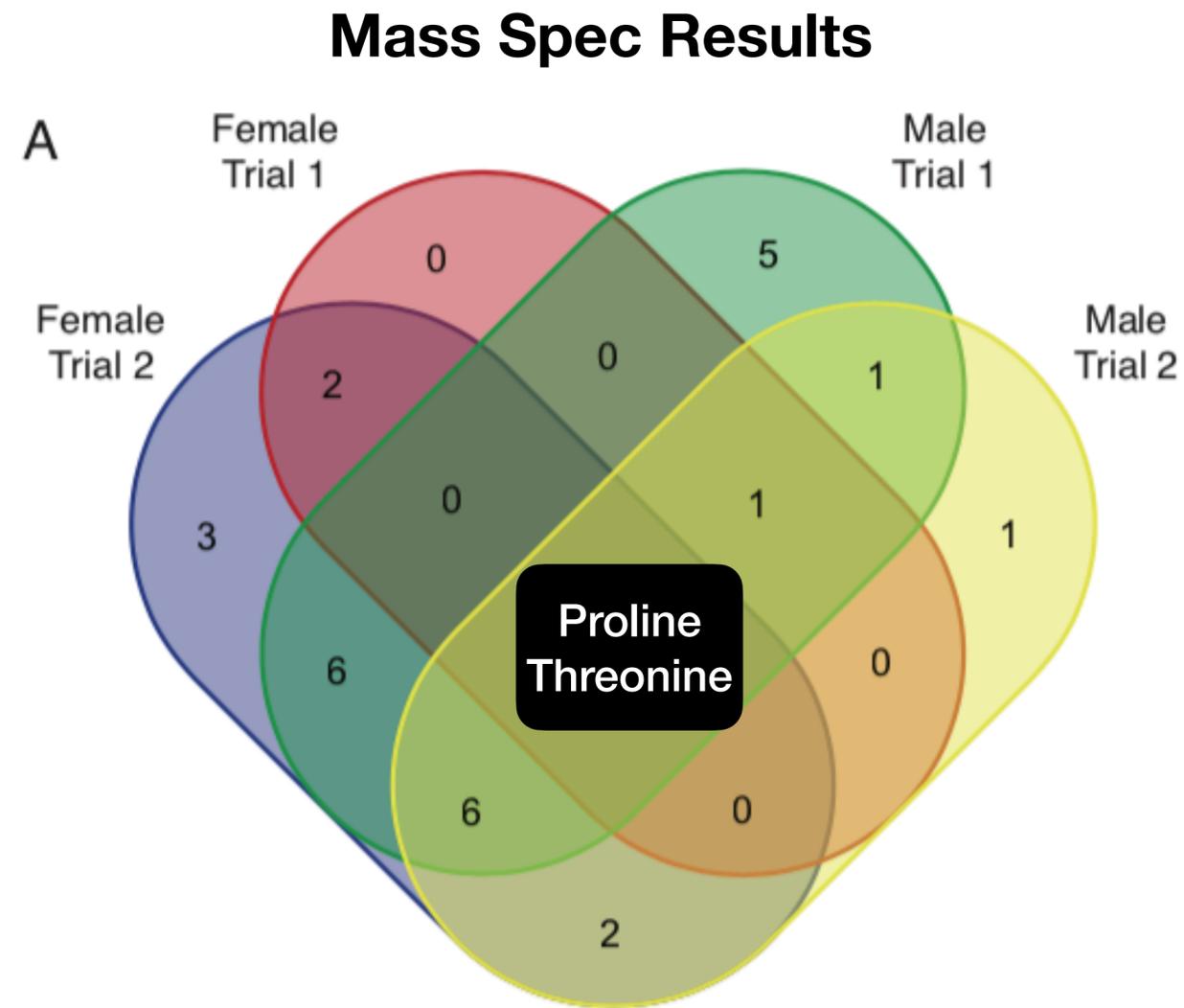
# Pilot Range Finding

- Already encouraging
- **Consistent high toxicity**
- **Consistent low toxicity**
- **Next steps:**
  - **normalize response cross-species**
  - **RNAseq**
  - **Metabolomics**

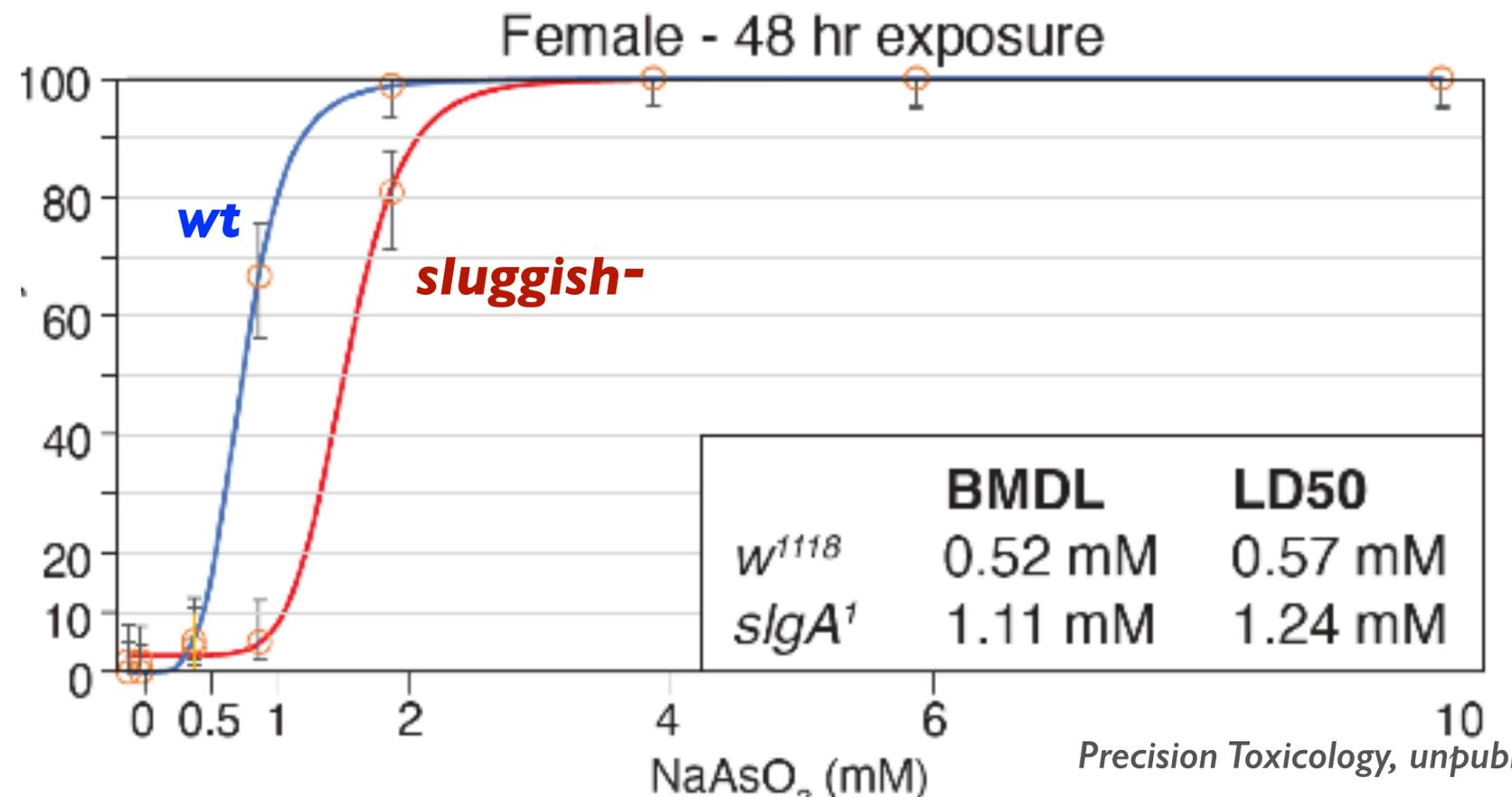
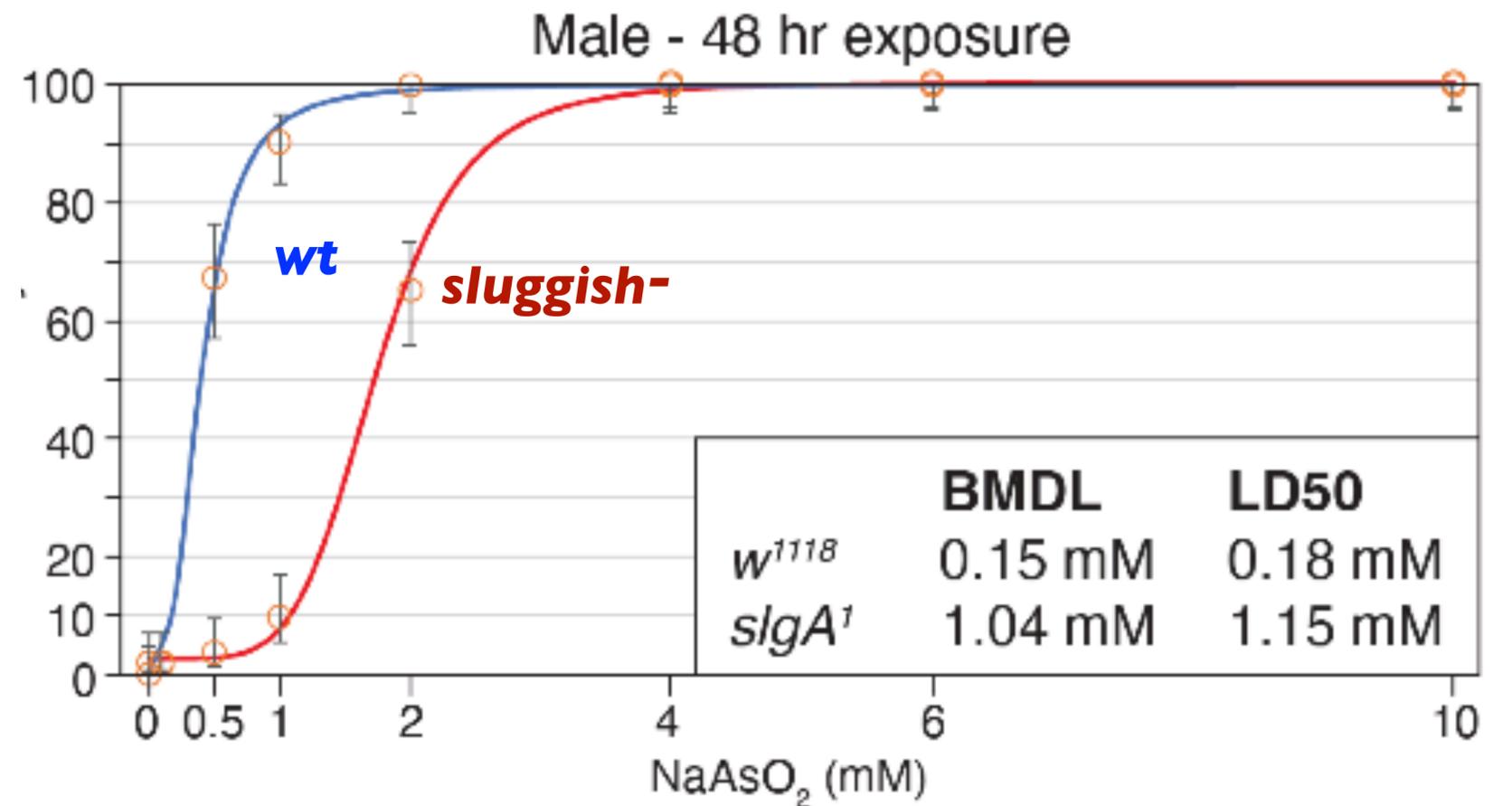


# Proline levels are up after arsenenic exposure

## Adverse or protective effect?

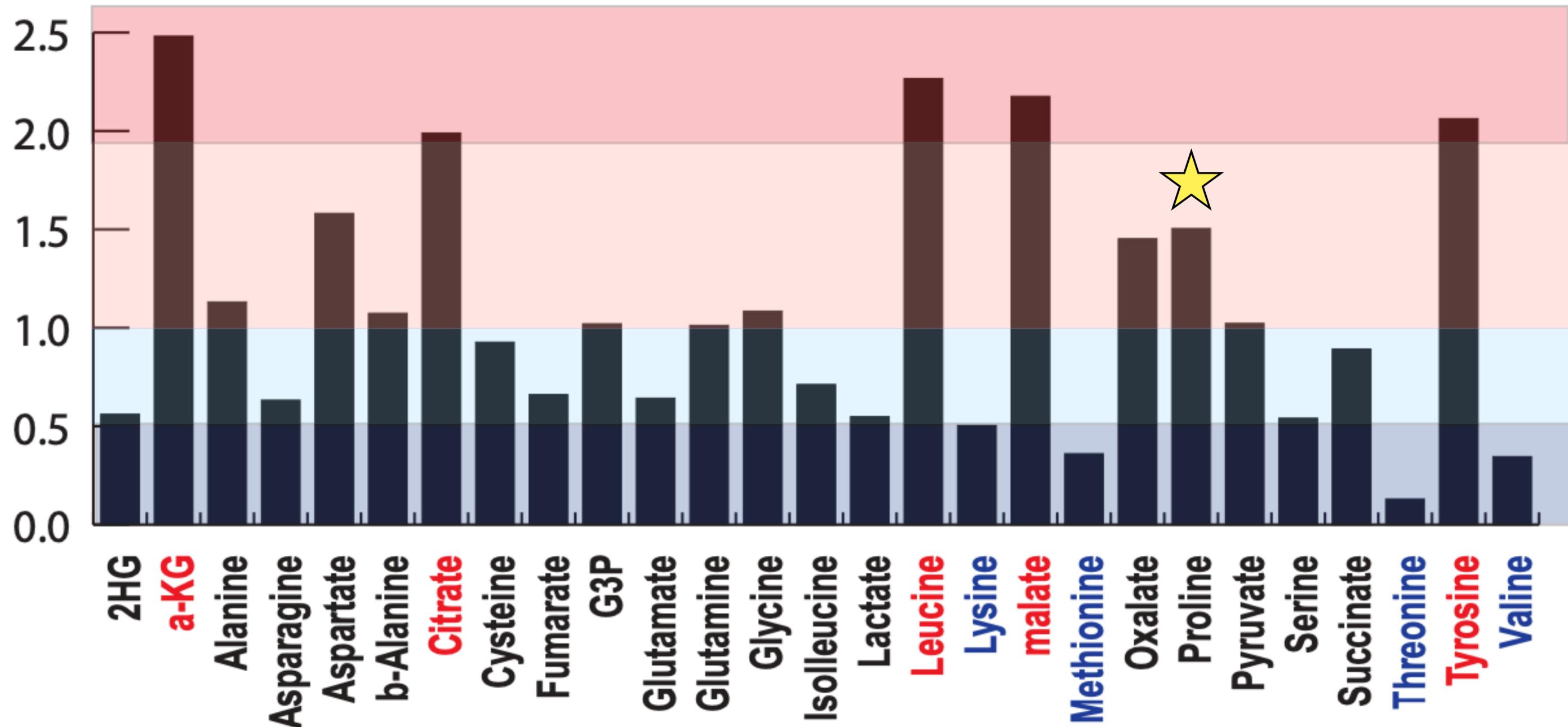


- *sluggish* suppresses toxicity.
- *sluggish* further increases Proline levels
- Consistent with Proline being protective against stress in plants
- Diminished sex-bias in *sluggish* mutants
- Sex-biased Proline levels?



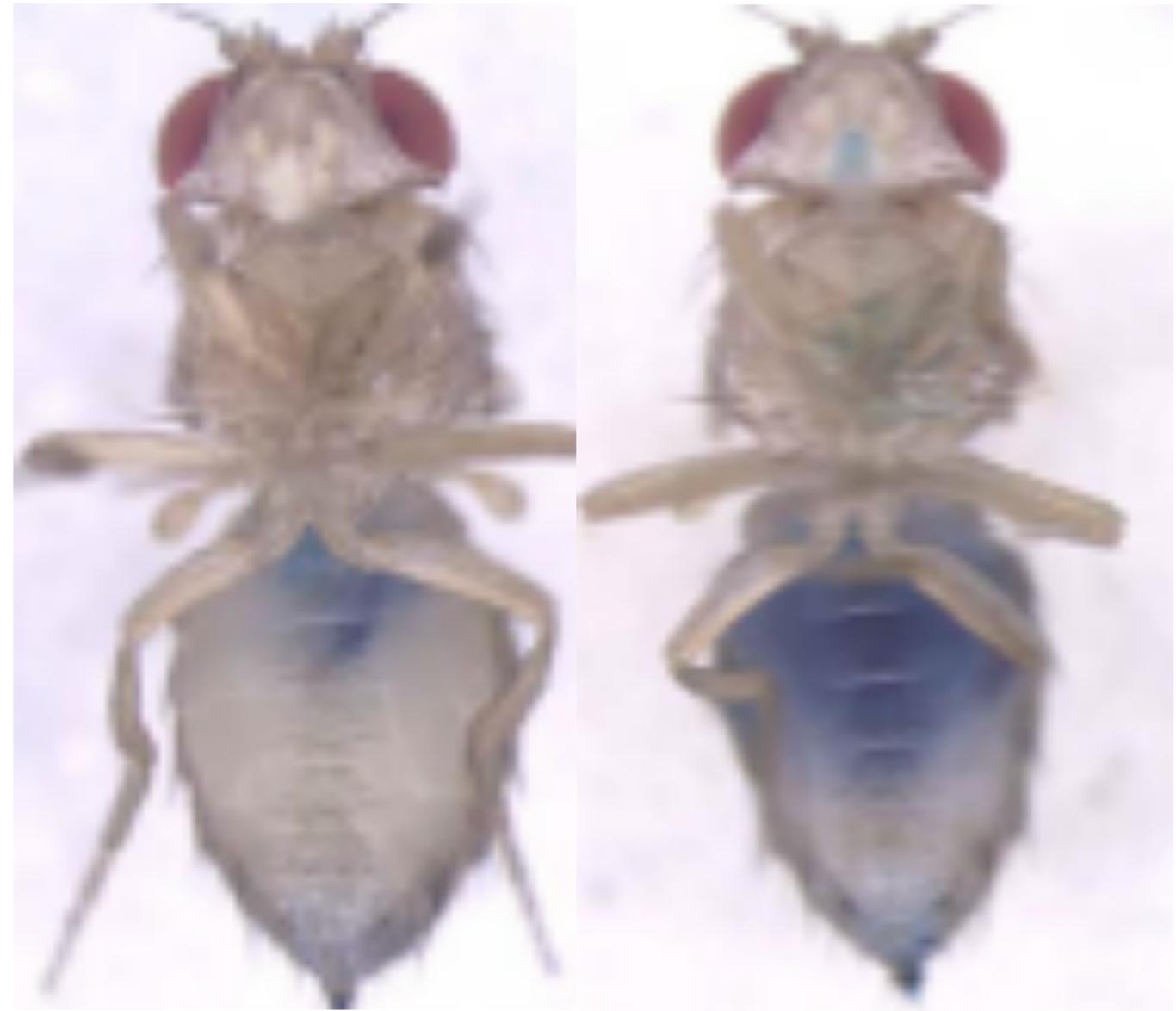
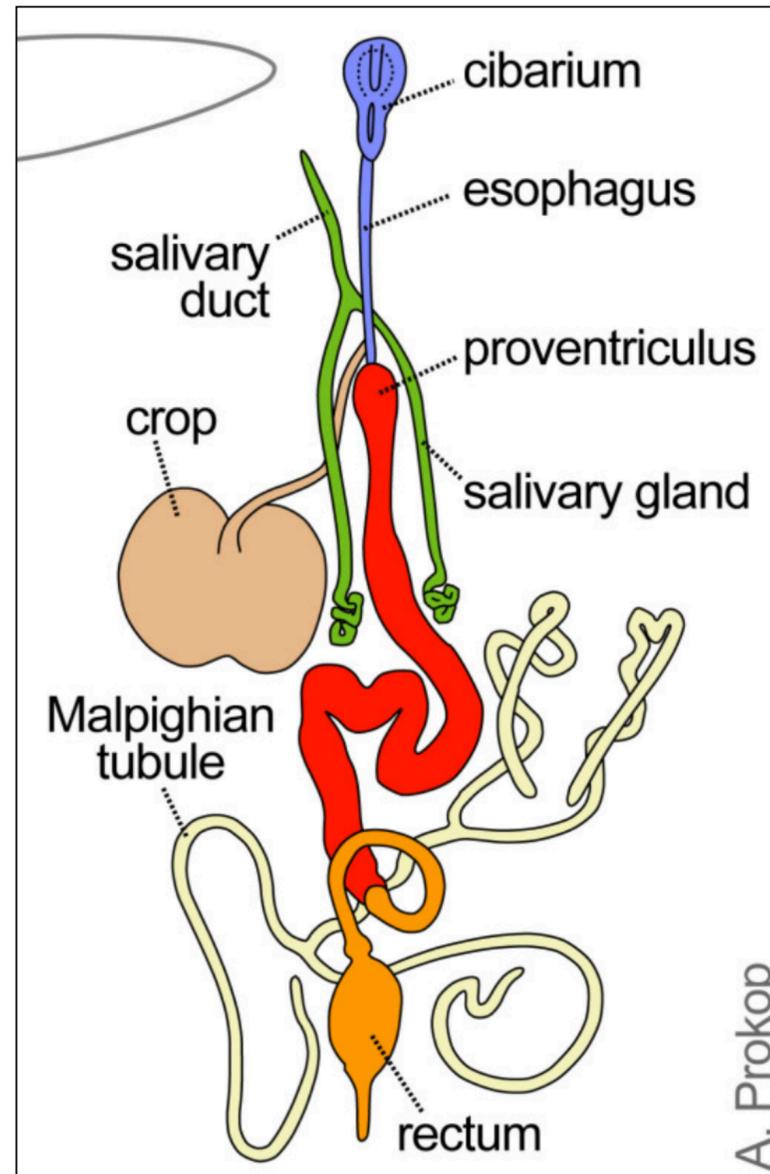
# Metabolites show sex-bias

Including 50% more Proline in females (**preliminary**)



# Even if they don't die, there's a phenotype

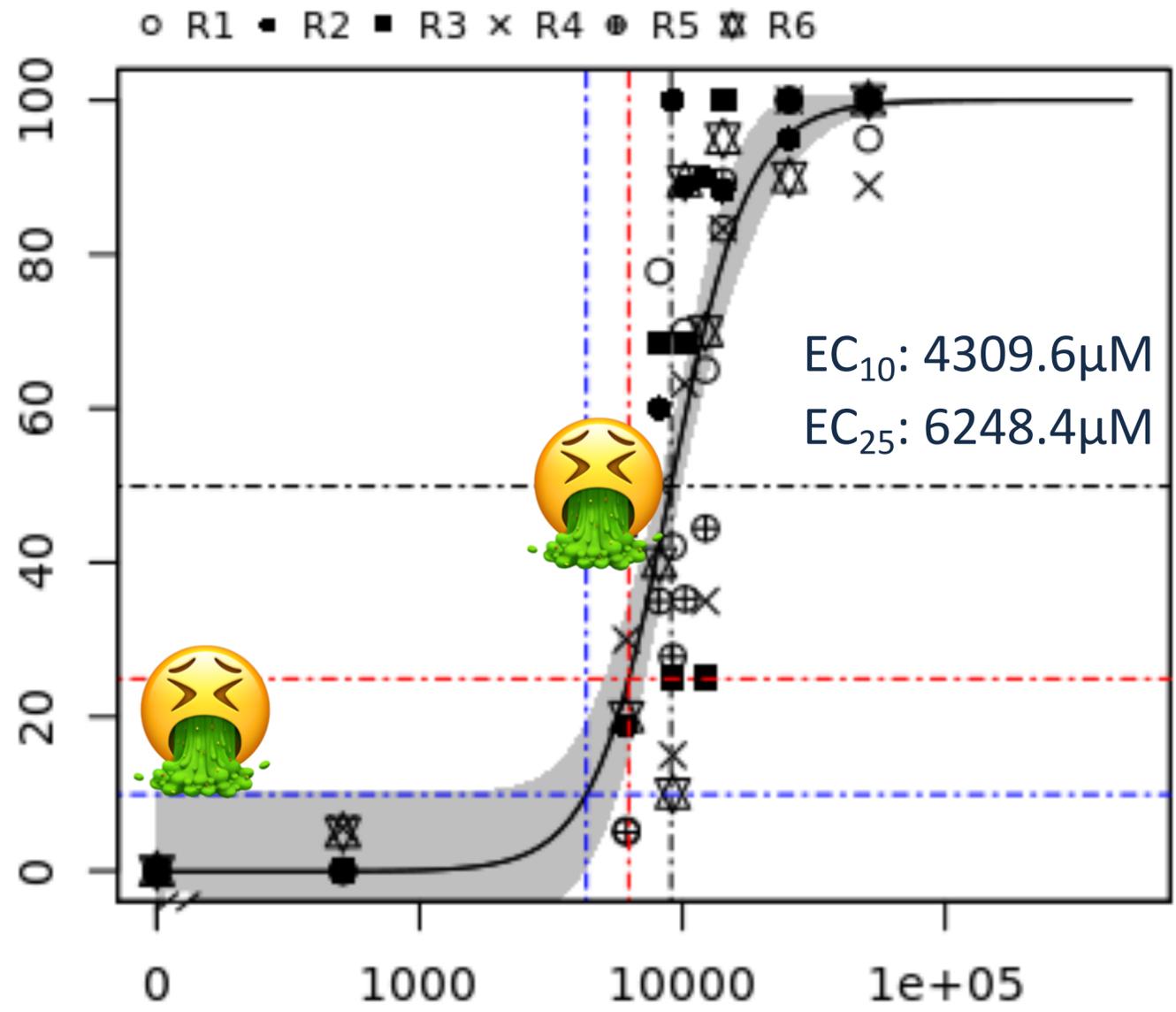
Retained food in stomach (crop). then vomit (regurgitate).



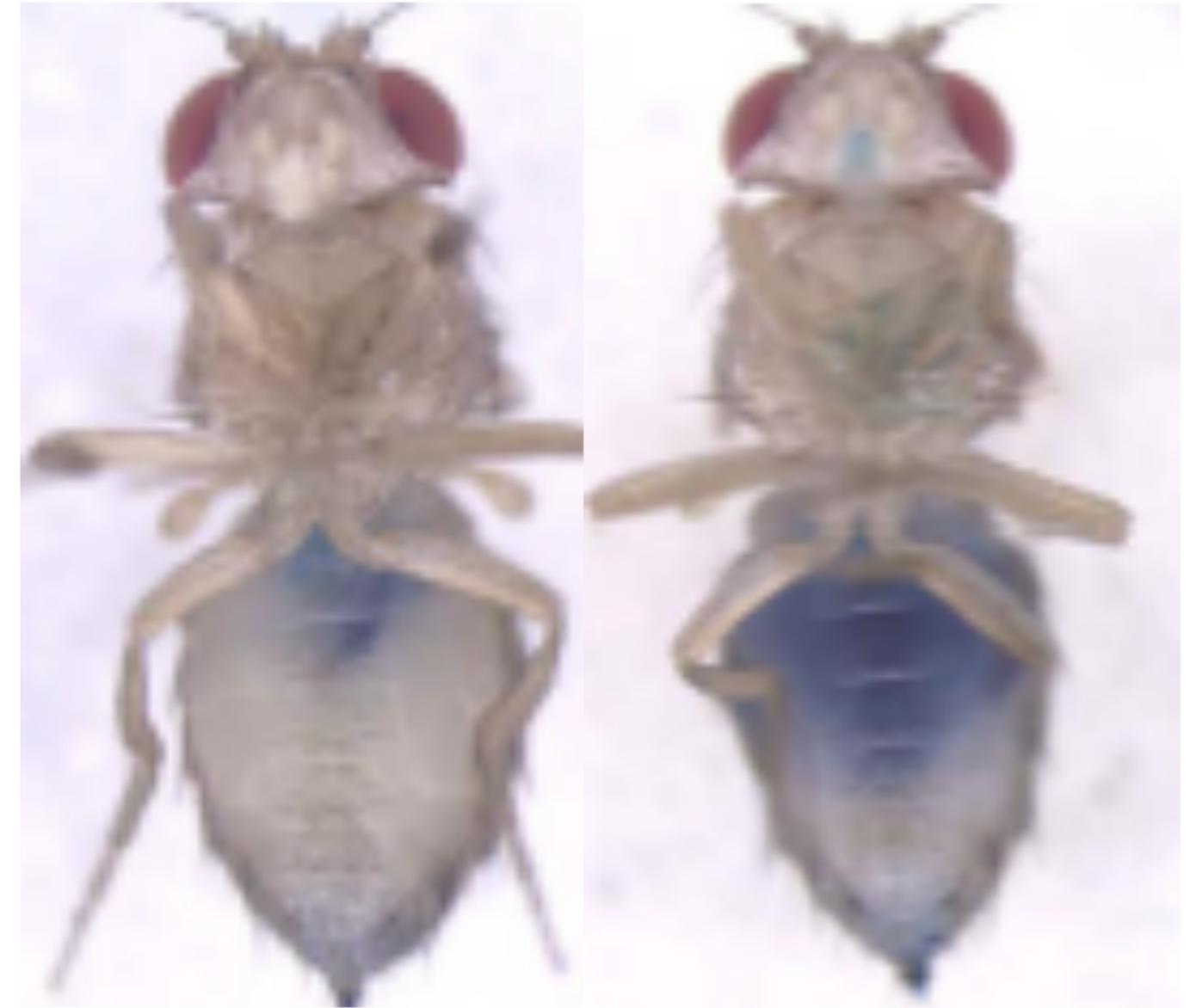
Blue food, 0% DMSO

Blue food, 2% DMSO

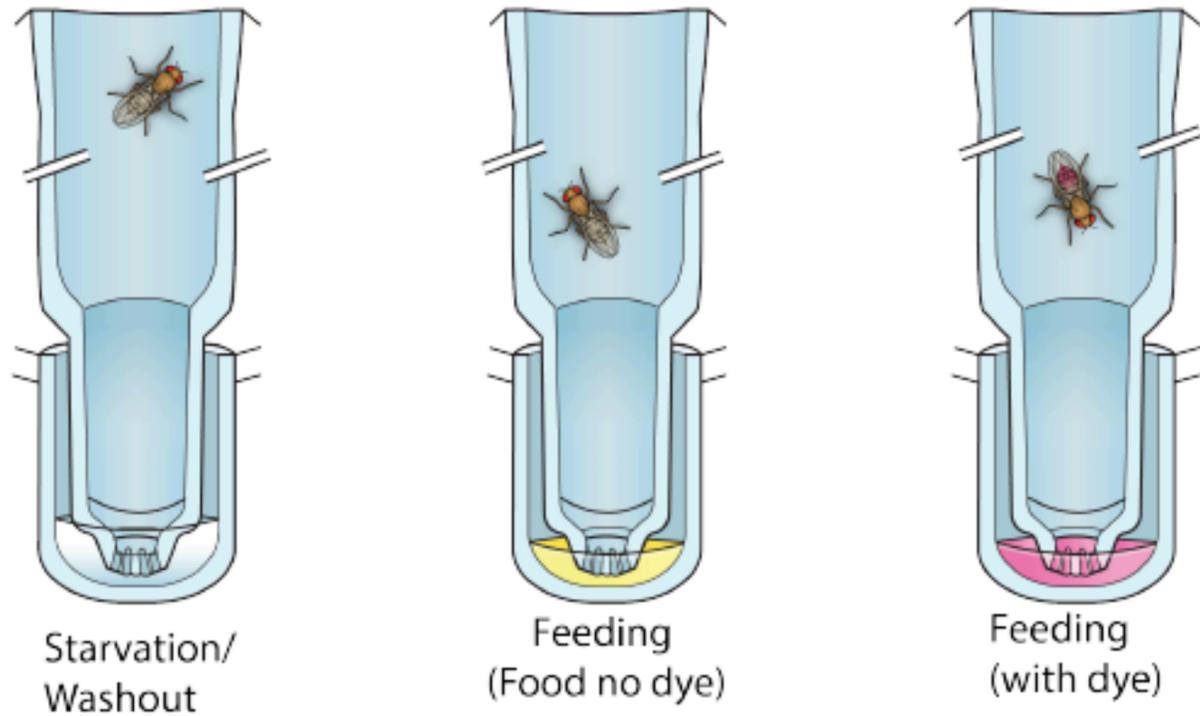
# Crop retention response often precedes death



^ some Crop Distension      ^ 100% Crop Distension

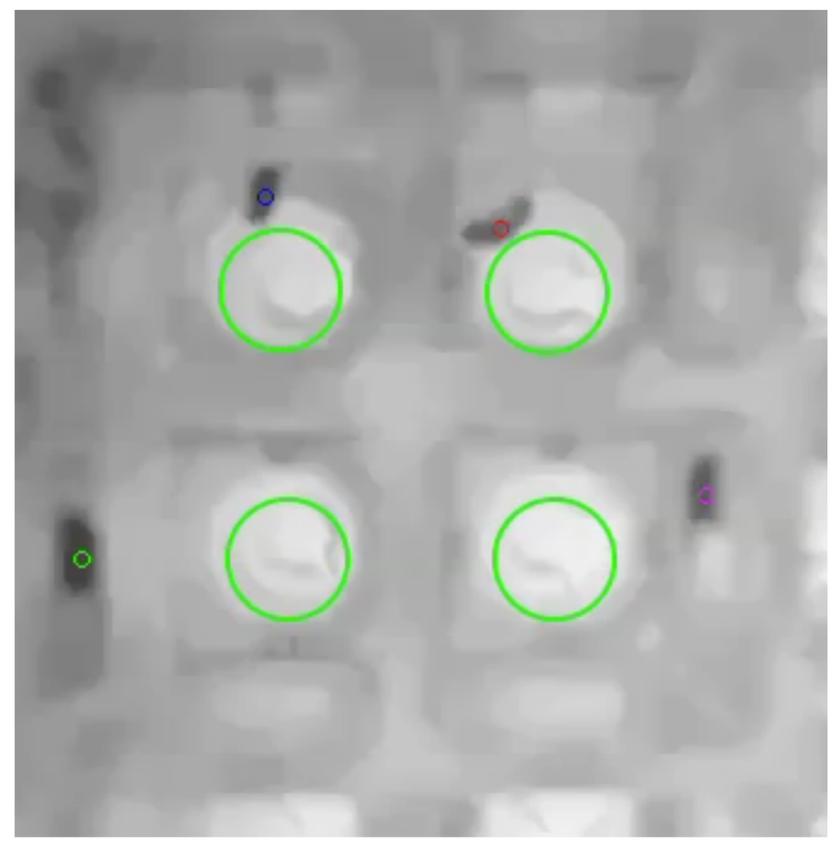
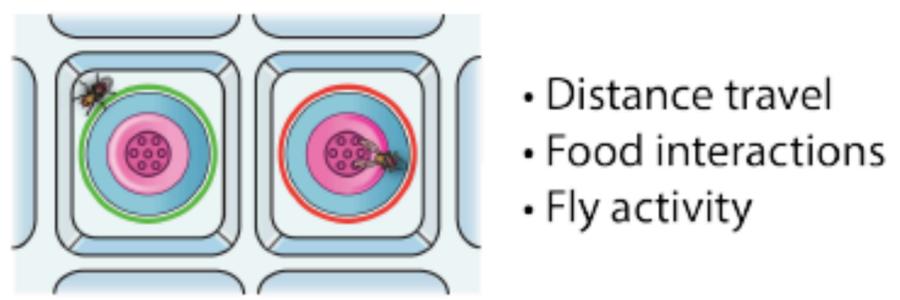


**A.** Treatment (acute bolus, chronic exposure, combinations)

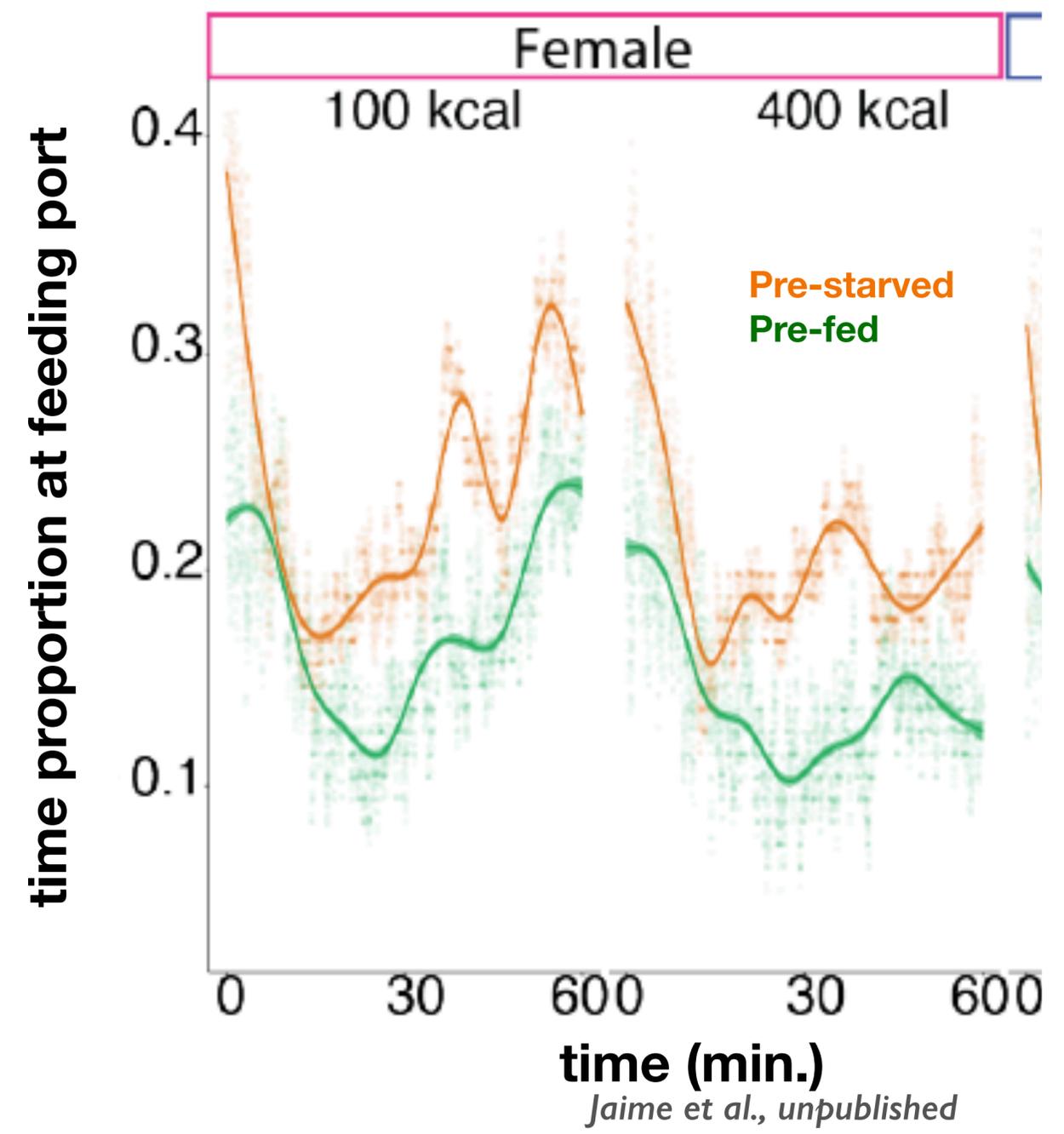
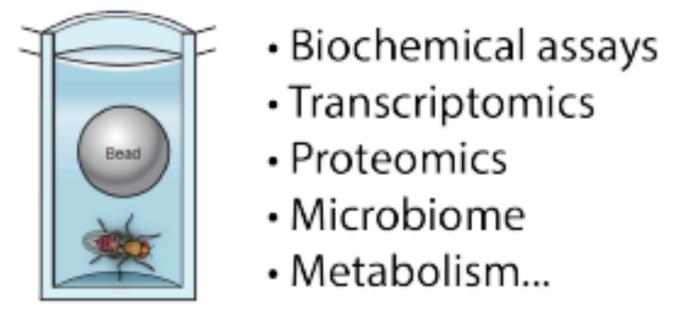


# Ministration and Industrialization Designing Flexible Plate Based Systems

**B.** Monitoring



**C.** Maceration



# My Discussion Points

- Genetic and Biochemical Pathways cut across the NAMs. Make sure to capitalize.
- Data mining requires data. Lot's of data. Lot's of organized data. FAIR (find, access, interoperate, and reuse).
- More diversity of cell types and organoids. More single cell genomics.
- Physiology and complex behavior requires whole organisms. Full stop.
- NAMs must scale. This needs to be an industry, not a research program.

**Sharvani  
Mahadevaraju**



**Leif Benner**



**Pradeep Bhaskar**



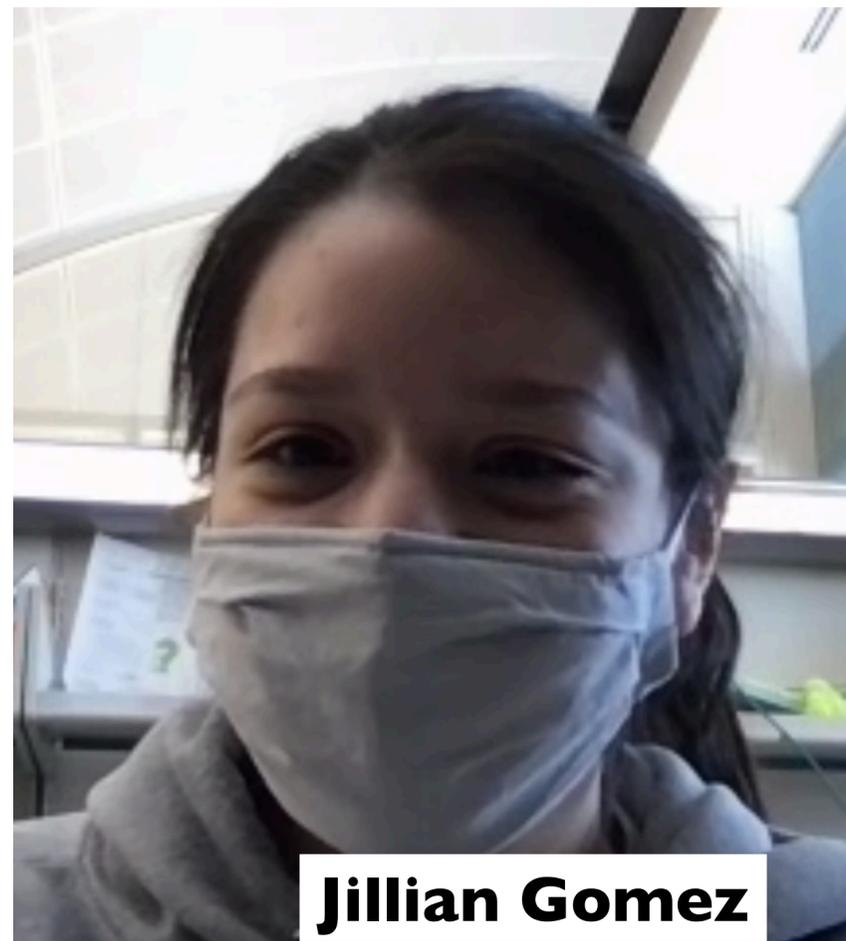
**Savannah Muron**



**Teresa Przytycka**



**Soumitra Pal**



**Jillian Gomez**



**Ibraheem Farooq**

# Sex in the FCA: Michelle Arbeitman, Michelle Bland, Sumatra Pal, Teresa Przytycka, Liz Rideout, Geo Vogler, Marianna Wolfner

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