

Exploring the Biology of Deep Underground Mining



Thomas Merritt

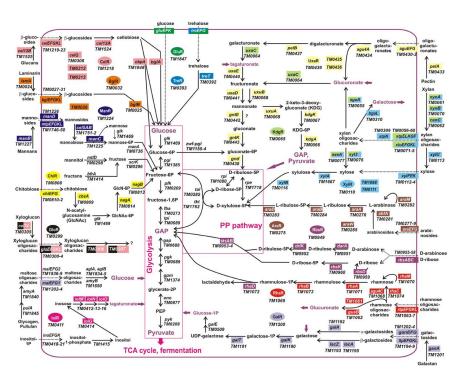
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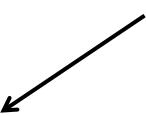




Mining Environment

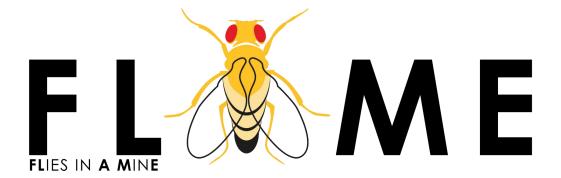
- Miners have to work in depths as great as 3.9 km.
- Atmospheric pressure over 20% greater than surface.
- Physically Strenuous Job







(Koven, 2010)





Explore the physiology of working deep underground environment

Important differences is likely pressure – but is it?



Won a Nobel Prize



Allowed others to win 7 Nobel Prizes

Why flies



Miner



Fly

Similar Biochemistry and Metabolism

Up to 80% of human disease genes are found in flies

Ease and utility

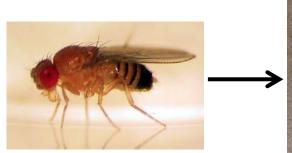




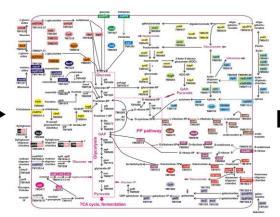




Longevity

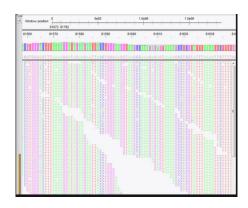






Metabolism



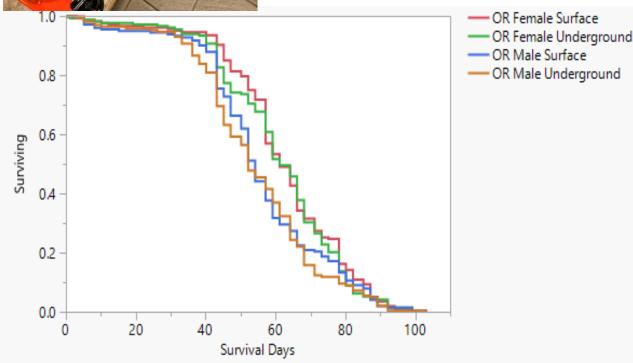


Gene activity



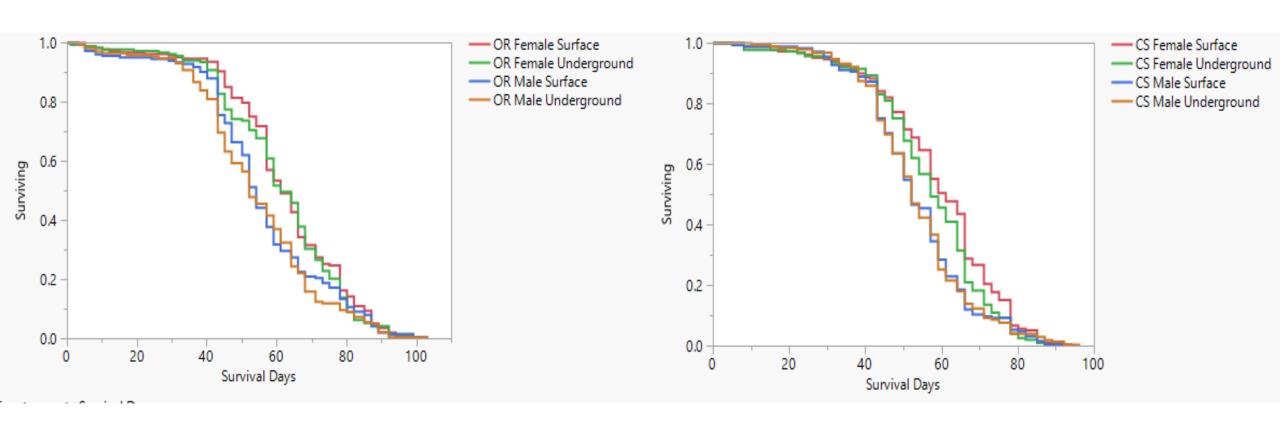
Longevity assay

Count the dead

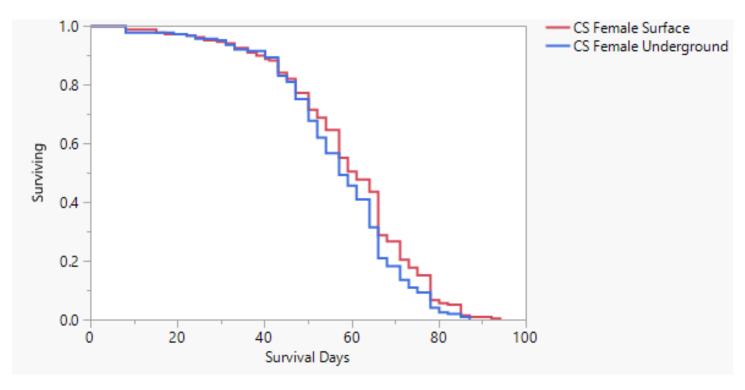


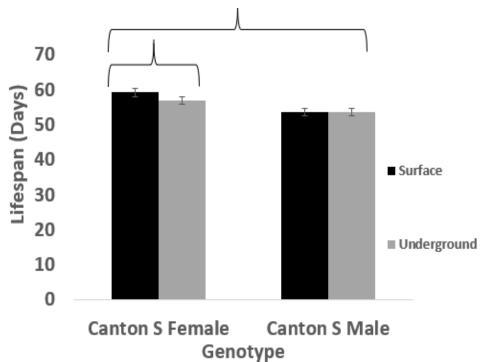
- How long do flies live on surface and underground?
- A very holistic measure of health
- 200 flies stayed on surface,
 200 flies went underground until all were dead
- Calculate % dead every 2 days

Longevity Results



Underground Treatment Significantly Decreases CS Females Lifespan





Fly Treatments

Underground Exercise



Surface Exercise

Surface Control

























Flygometer 2.0 In Action

How does one exercise a fly?

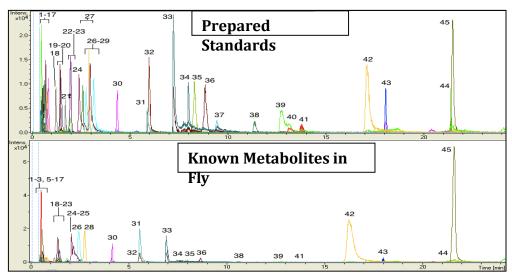
Not surprisingly, different flies take to the treadmill differently





Broad-Based Metabolomics

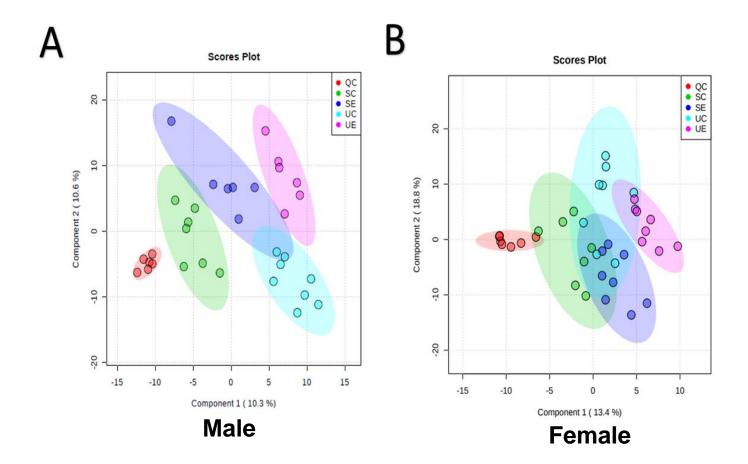
Known metabolites as standards and in fly homogenate



- Identify as many metabolites as possible
- Liquid chromatography / mass spectrometry based metabolomics
- LC separates complex biological sample into individual metabolites
- Mass spec identifies metabolites based on extremely accurate molecular weight
- No, its not that easy

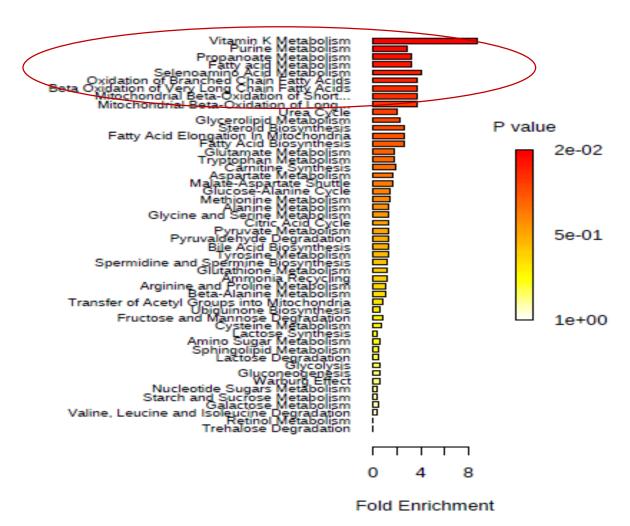
Metabolomic Shift Caused by Underground Exercise

- LC-MS almost 20,000 potential metabolites
- Almost 4% of metabolites differ between Surface and Control
- PLS-DA Plot



Vitamin K, Fatty Acid, and Nucleic Acid Metabolism Enriched

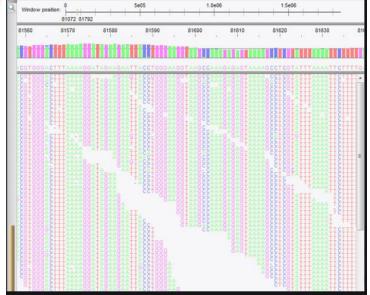
Enrichment Overview (top 50)





Broad-Based Transcriptomics

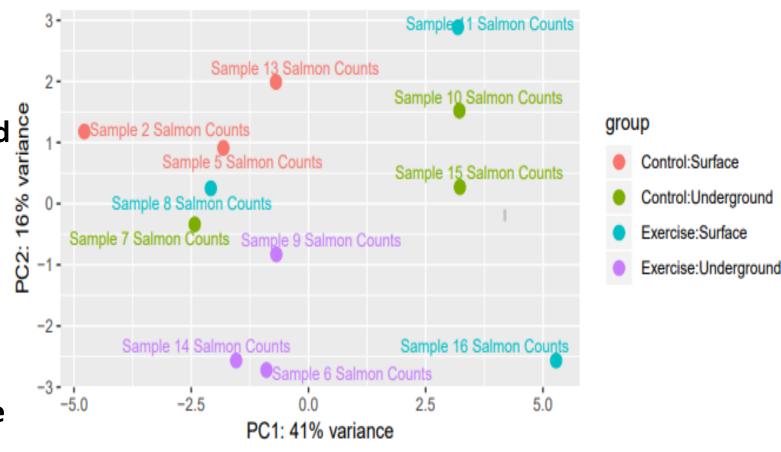
Compare with established databases



- Identify as many expressed genes as possible
- "Next Generation" sequencing of transcripts
- Computationally identify expressed genes
- No, its not that easy (OK it sort of is this easy)

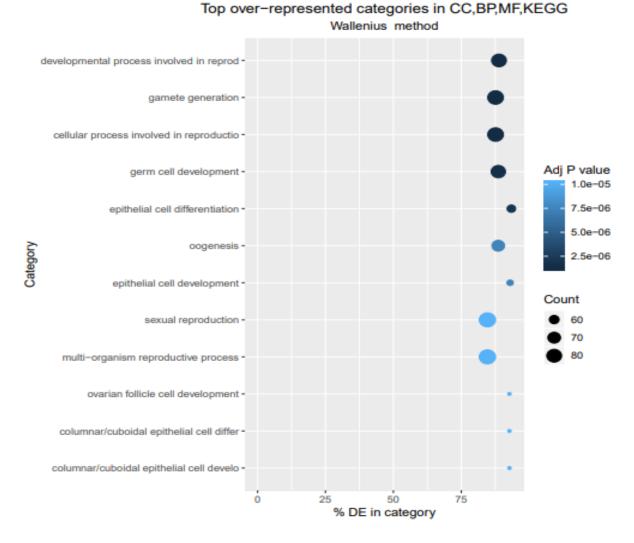
Underground Exercise Shows Transcriptomic Shift

- NovaSeq mRNAseq 100 bp reads
- 71.3 million reads, from ~13,000 genes ~800 genes differed between surface and underground (~6%)
- CS females, 1 week treatment
- PCA Plot Comparing Treatments
- Largest Shift Between Surface Control and Underground Exercise



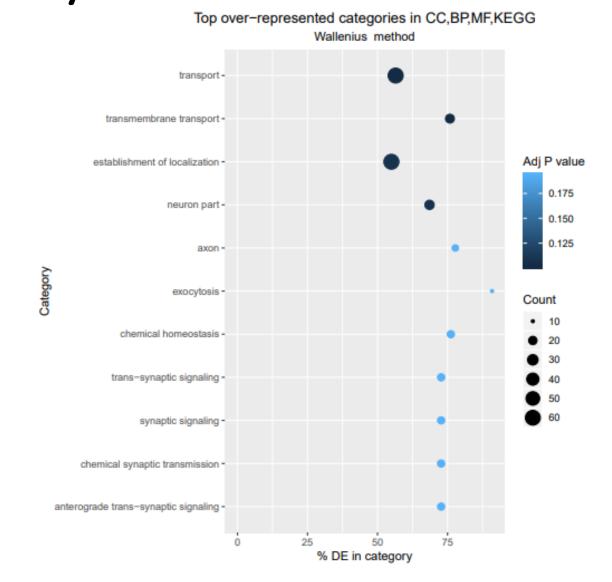
Underground Exposure Upregulates Reproductive Gene Pathways

- Gene Ontology Analysis
- Canton S Female
- Weeklong Exposure
- Surface Control vs Underground Exercise
- Over-represented
 Categories in Female
 Reproductive Processes



Underground Exposure Downregulates Gene Pathways

- Gene Ontology Analysis
- Canton S Female
- Weeklong Exposure
- Surface Control vs Underground Exercise
- Over-represented
 Categories in Transportation
 and Signalling



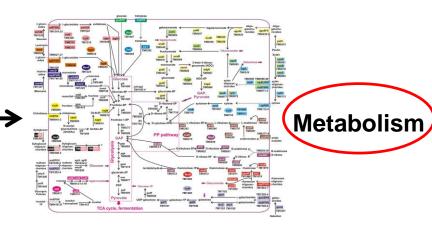


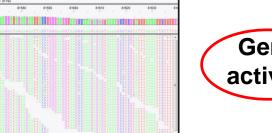












Gene activity





Don't underestimate the Gee-Whiz factor!



Cue the CBC



Art MacDonald signed Fly Vial



LAB SPACES / S1 EP7
Inside the Dark Matter Lab Buried
Over a Mile Underground



We have a responsibility to explain what we do and why its important



Acknowledgments

- Merritt Lab
- Dr. Heath MacMillan
- SNOLAB Staff









Canada Research Chairs

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