

Building a Metabolomic Bridge from  
Genotype to Phenotype  
or  
*How and Why* Do Things Fall Apart  
with Age?

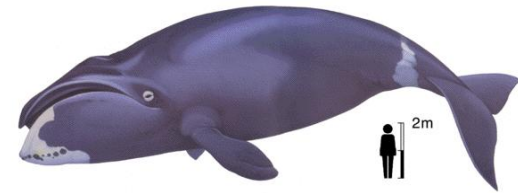
Daniel Promislow

Departments of Pathology & Biology  
University of Washington, Seattle, WA  
promislo@uw.edu

# Two messages

1. By thinking about how cells (and organisms) fall apart, we can understand some things about how they are put together.
2. Analysis of the building blocks (the metabolome) can help us accomplish this.

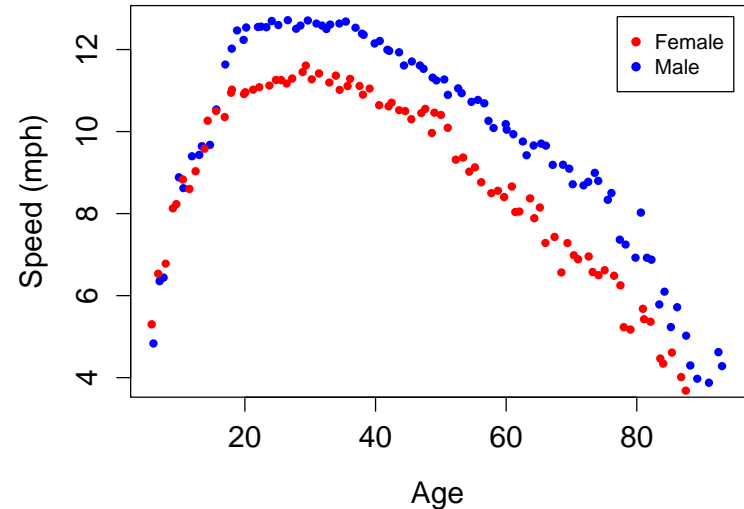
# 1. How (and why) do some species live longer than others?



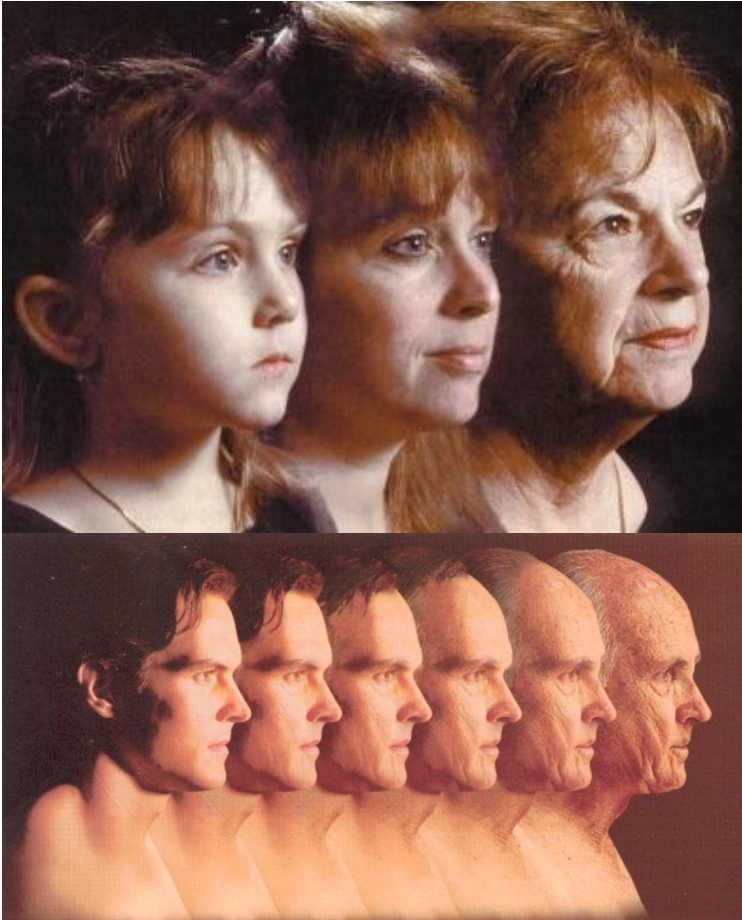
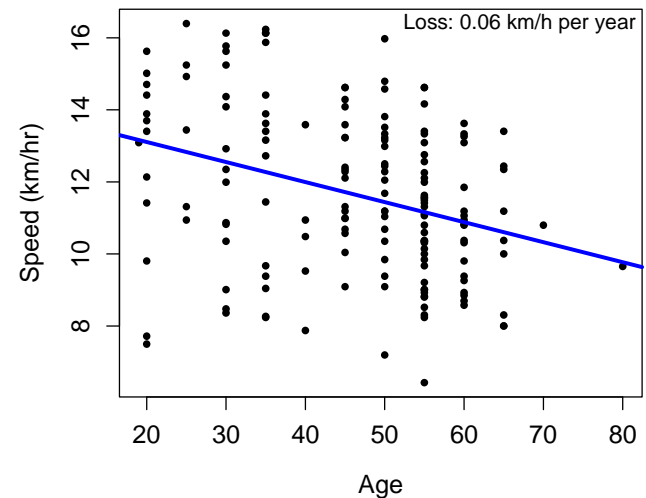
# 2. How (and why) do some individuals age faster than others?

Data source: [http://arrs.net/SA\\_Mara.htm](http://arrs.net/SA_Mara.htm)

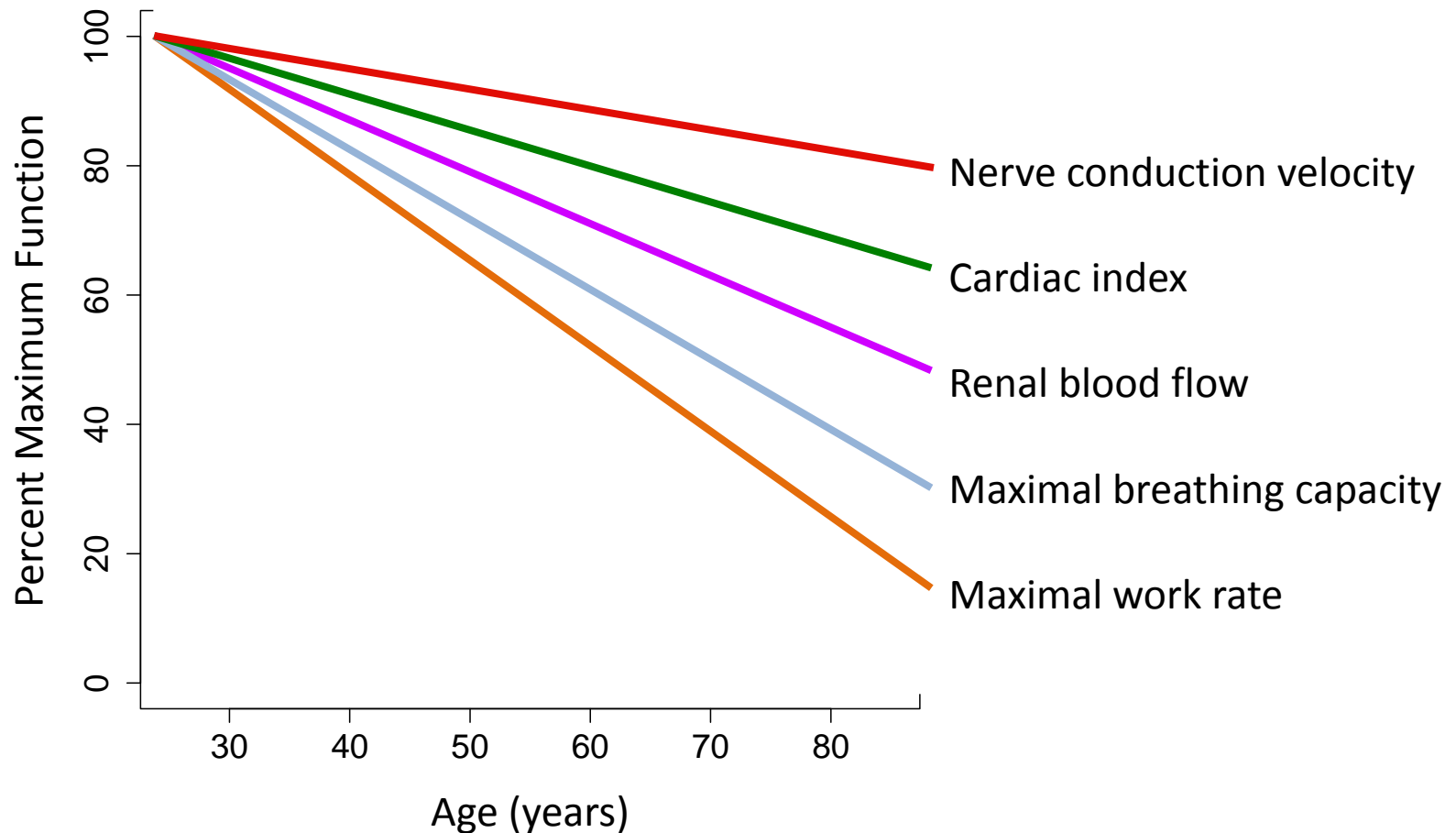
World marathon record speed, by age



Speed, 50km Noqueமான Ski Marathon, 2012

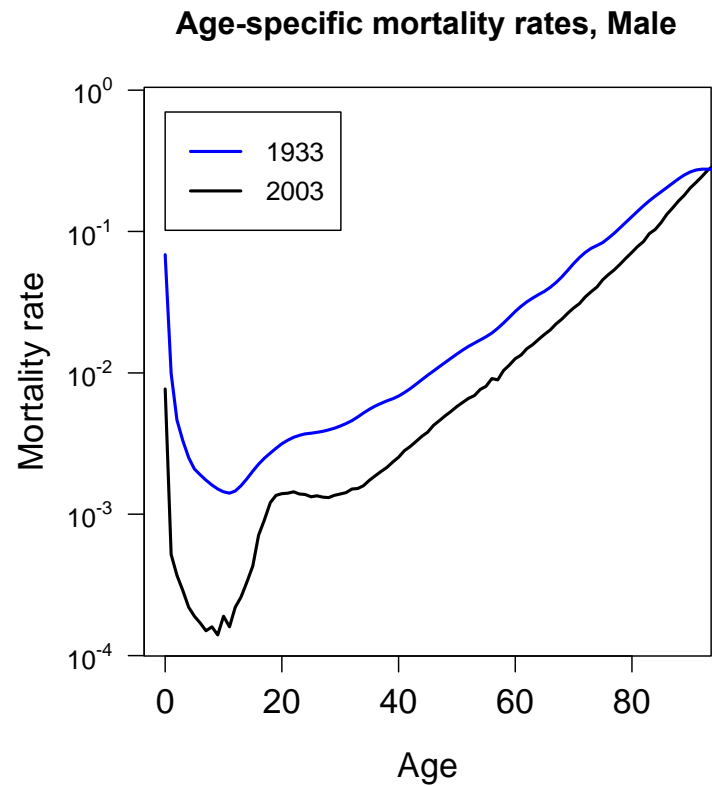
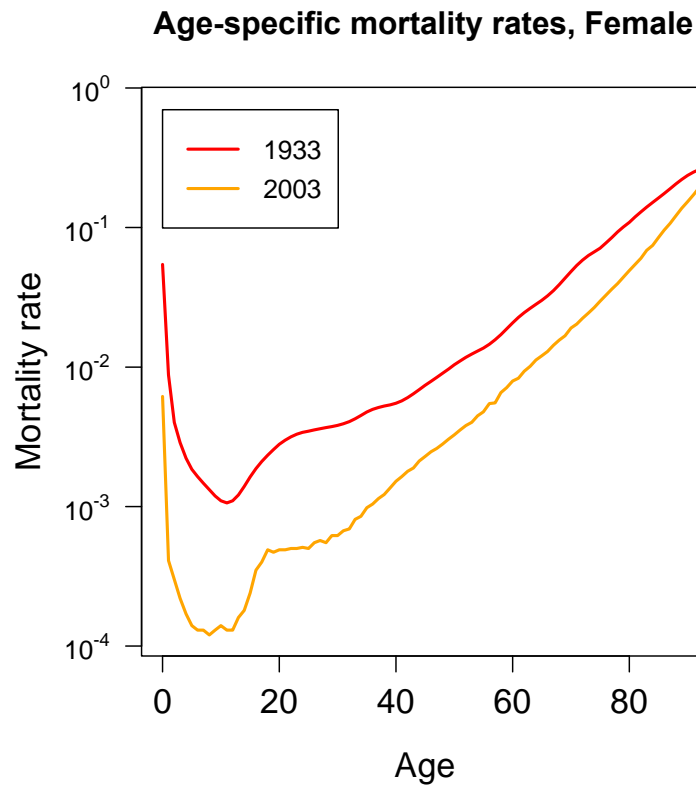


### 3. An *unasked* question: Why do some traits decline faster than others?

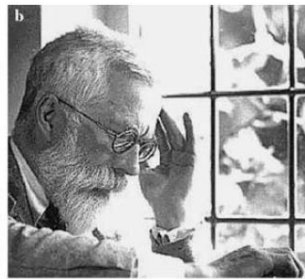
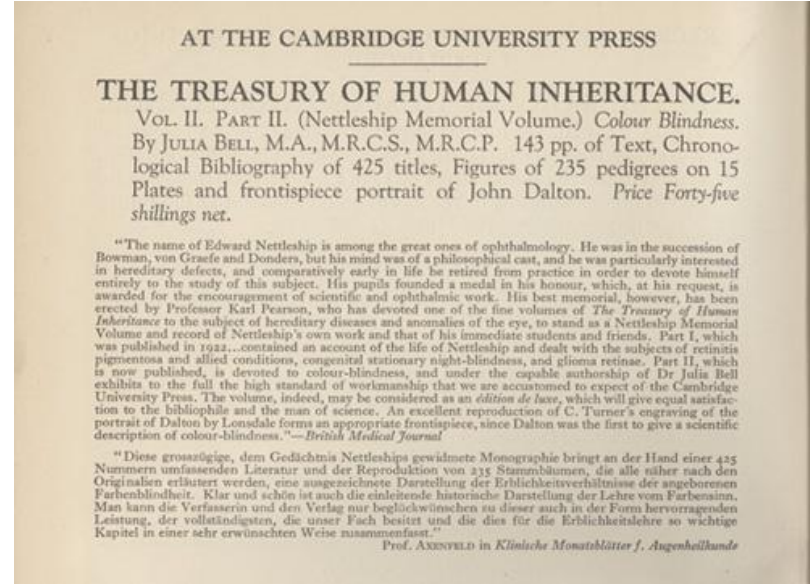


After Baker & Martin in Geriatric Medicine (ed. Cassel *et al.*, 1997)

# Senescence: age-related decline in fitness components



# Julia Bell (1879-1979)



Pearson, Fisher & Penrose



Haldane



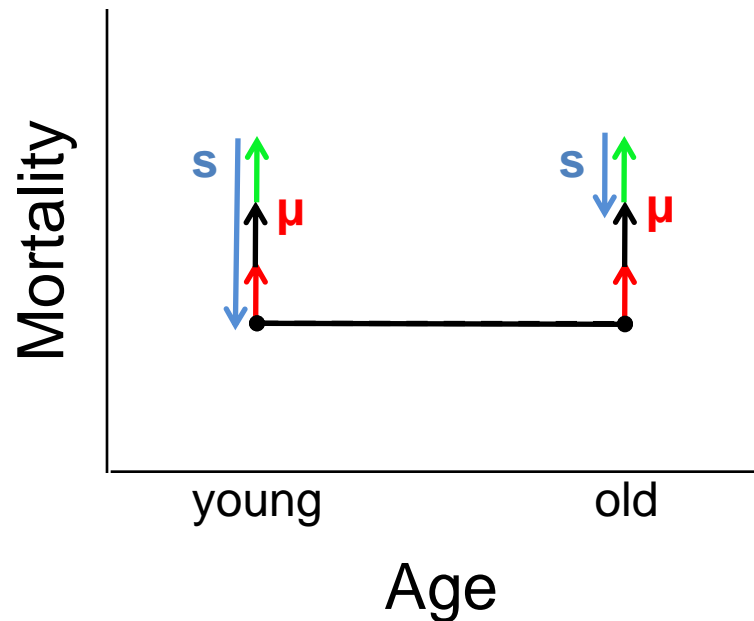
# Aging evolves due to maladaptation (Mutation Accumulation)

*“... If a genetic disaster happens late enough in individual life, its consequences may be completely unimportant”*

-An Unsolved Problem of Biology 1952



Peter Medawar 1946, 1952



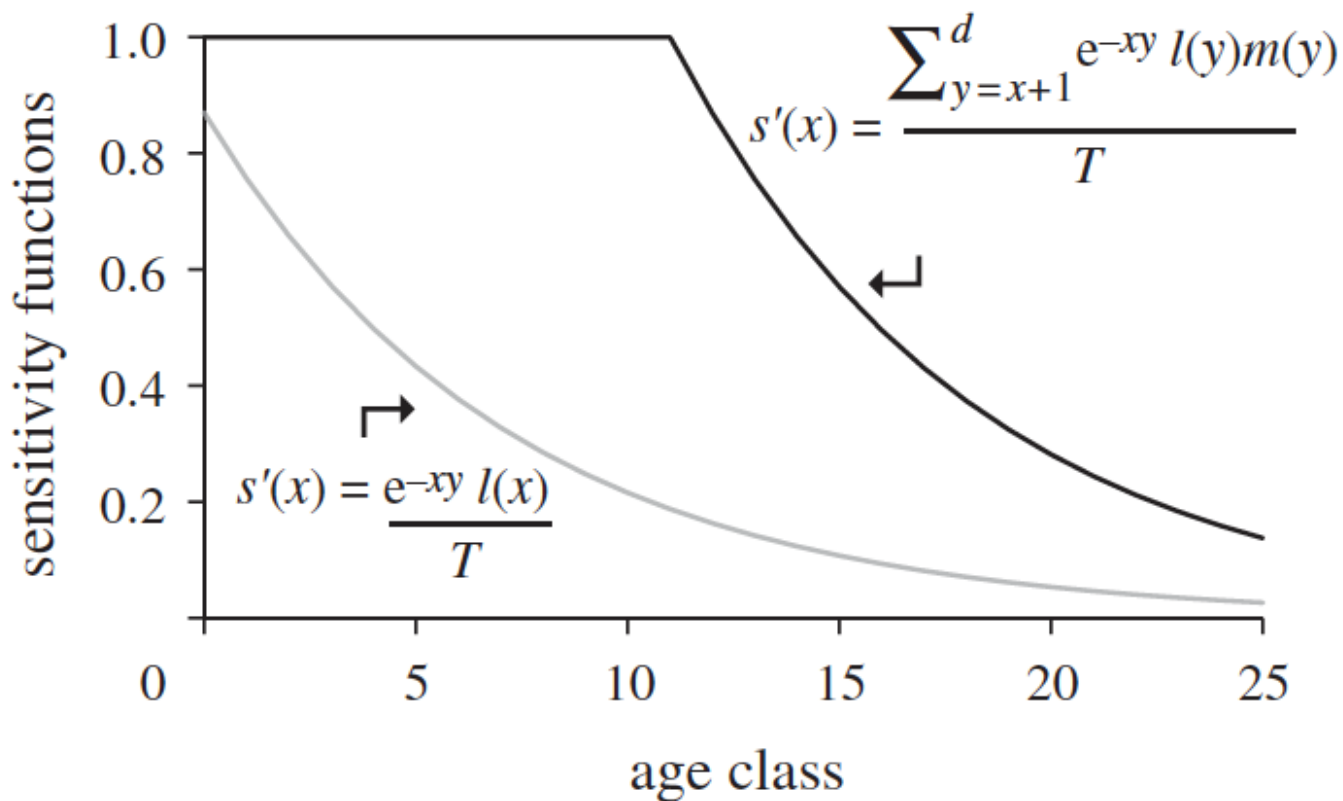
Mutations with uncorrelated or positively correlated effects across ages



*J. Theoret. Biol.* (1966) **12**, 12–45

## The Moulding of Senescence by Natural Selection

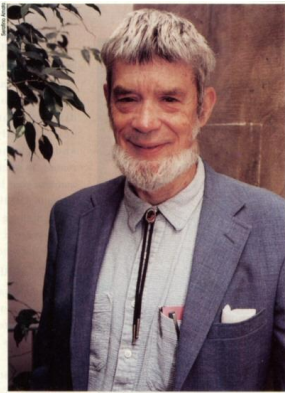
W. D. HAMILTON



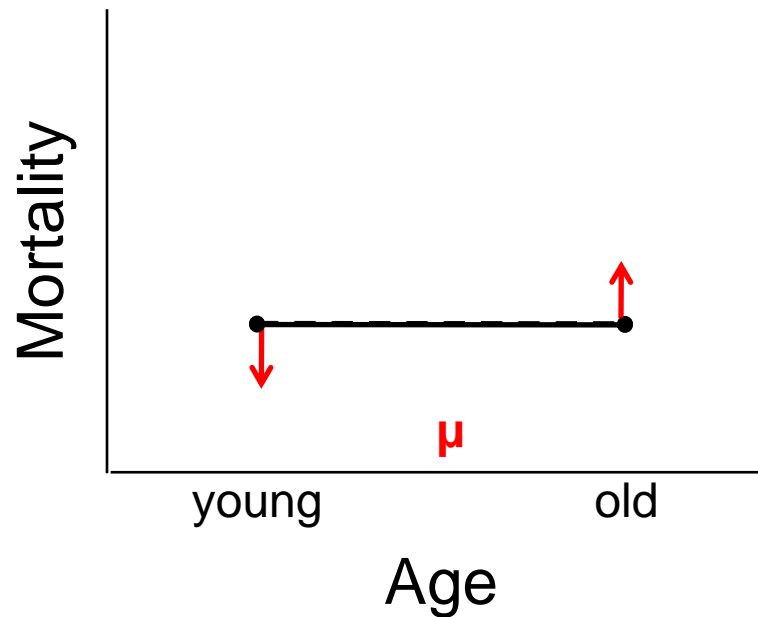
From K.A. Hughes 2010. *Phil. Trans. Roy. Soc.*

# Aging evolves due to constraint (Antagonistic Pleiotropy)

Natural selection favors alleles that increase survival at early age at the expense of survival at late age



George C Williams 1957



Mutations with negatively correlated effects across ages

# Outline

Evolution of cellular aging

1. Single-cell models of aging

Metabolomics and aging

2. Phylogenetic variation

3. Genetic variation

4. Diet Restriction and Networks

# Single-celled organisms won't senesce

## PLEIOTROPY, NATURAL SELECTION, AND THE EVOLUTION OF SENESCENCE <sup>1</sup>

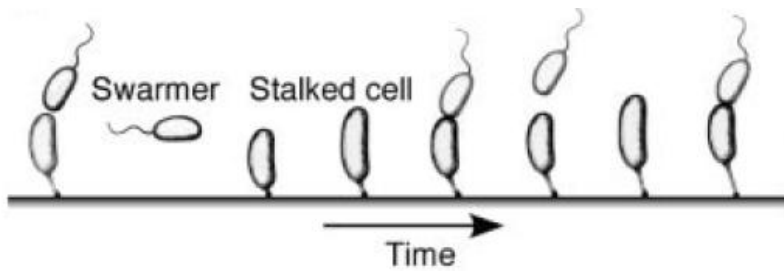
GEORGE C. WILLIAMS

*Michigan State University*

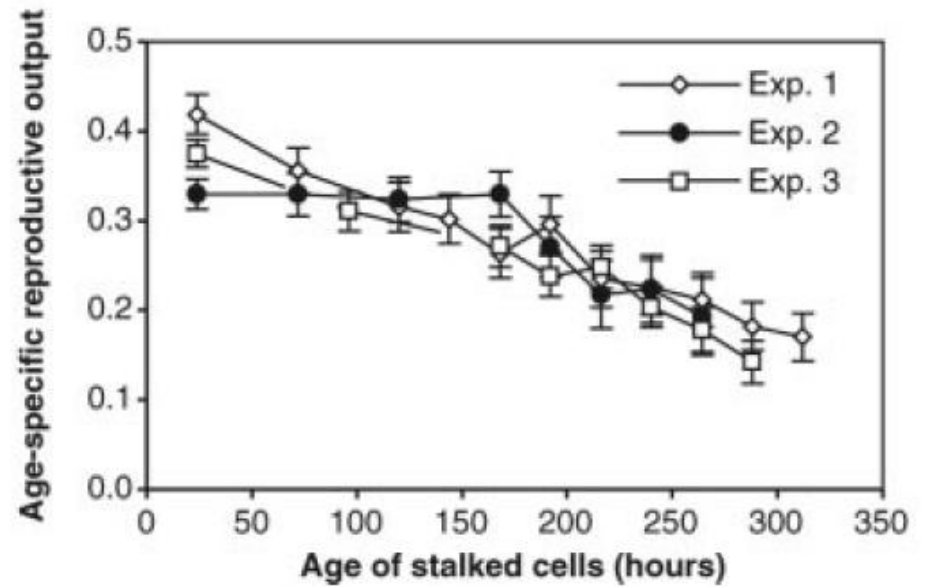
Received February 26, 1957

(1) *Senescence should be found wherever the conditions specified in the theory are met, and should not be found where these conditions are absent.* There are organisms in which the distinction between soma and germ-plasm may not exist, but the other assumptions of the theory would seem to be inevitable for any organism, at least for any that has a clear distinction between soma and germ-plasm. The theory regards senescence as an evolved characteristic of the soma. We should find it wherever a soma has been evolved, but not elsewhere.

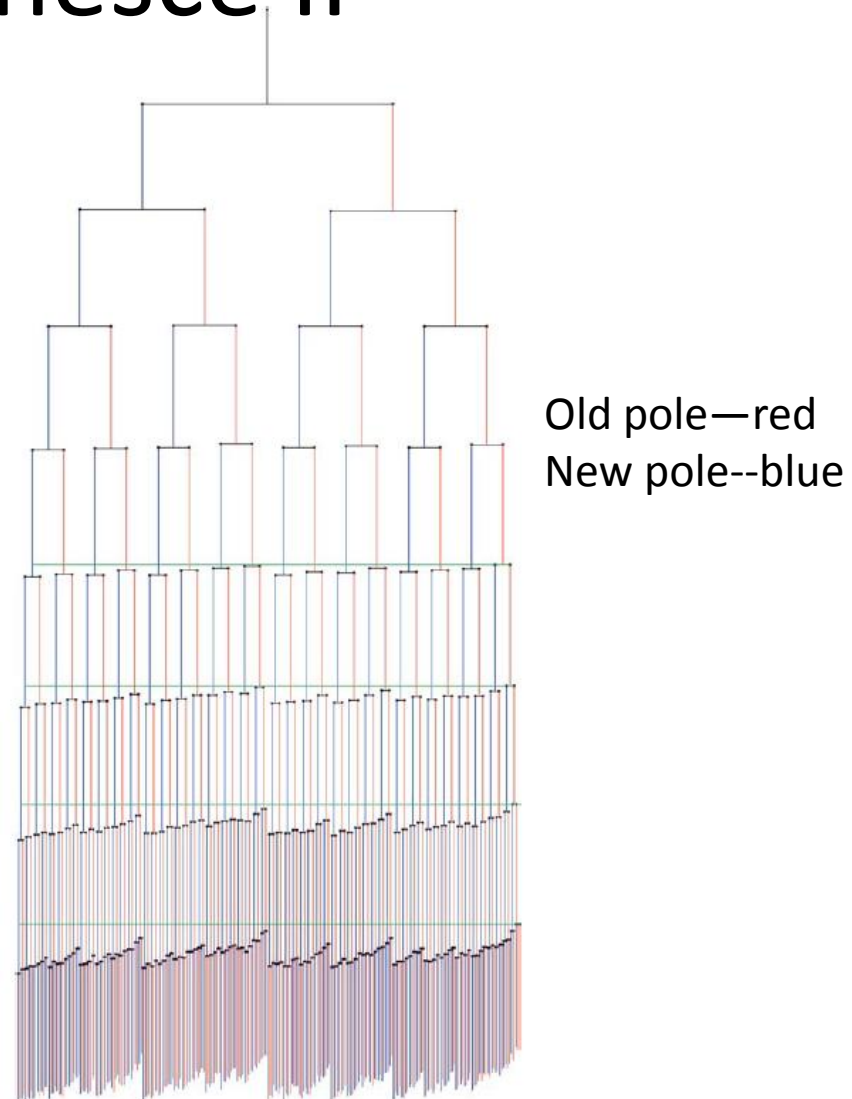
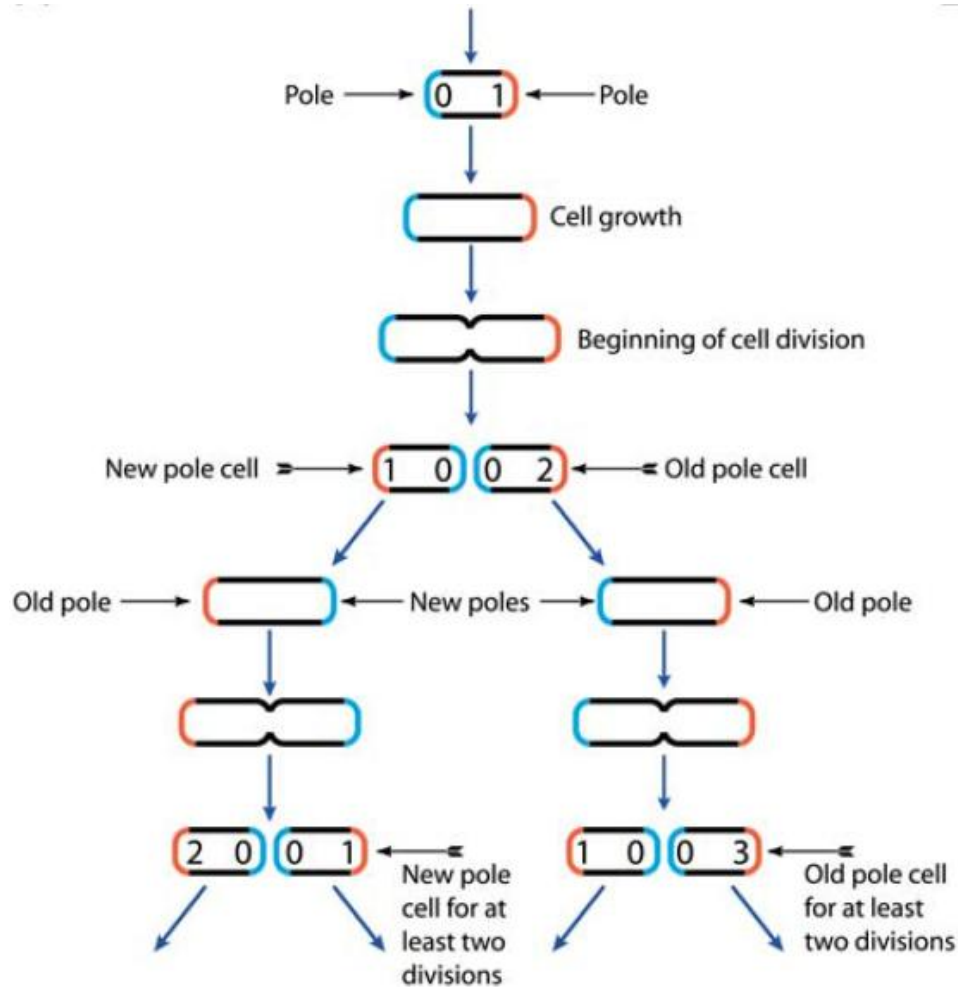
# Bacteria senesce I



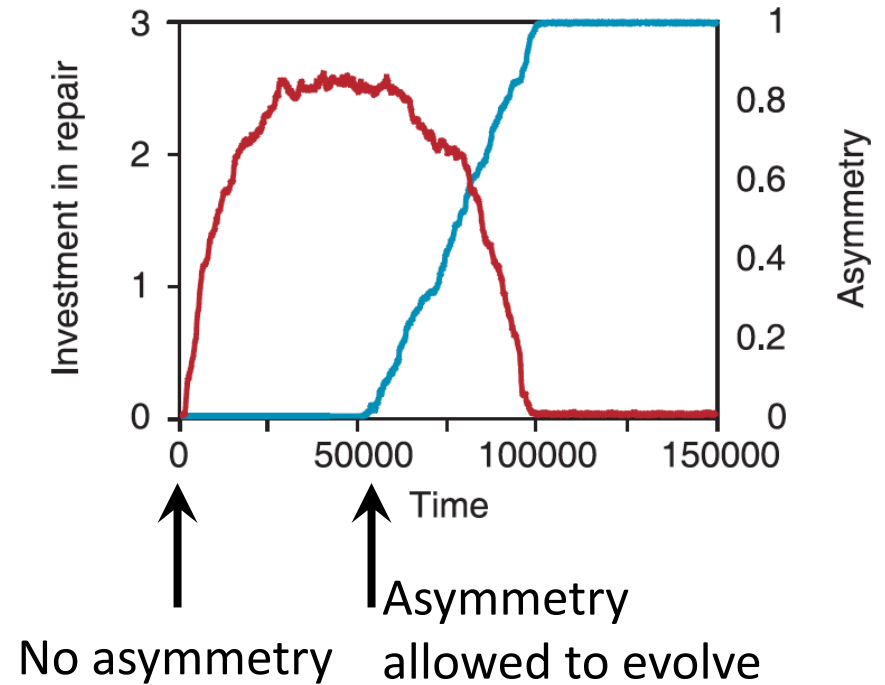
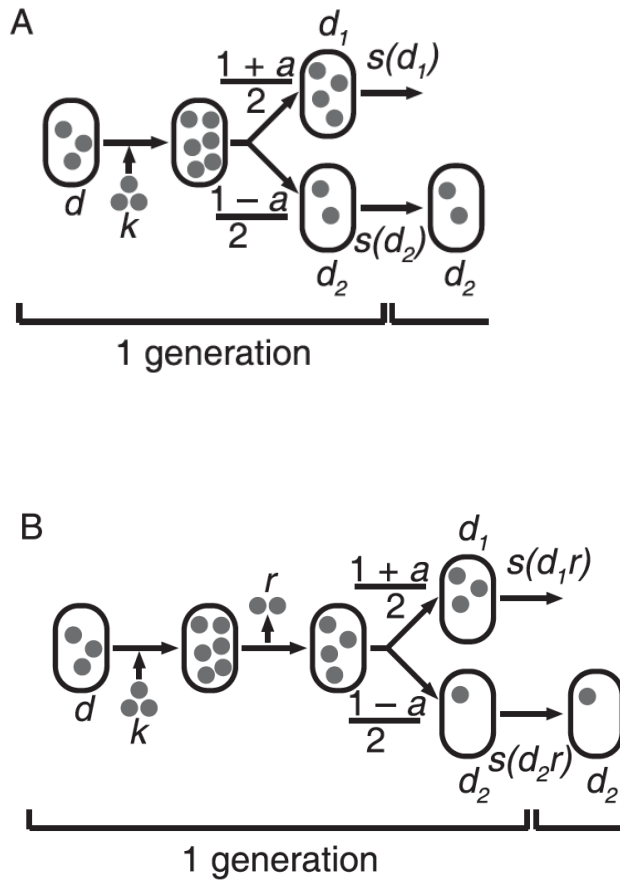
Life cycle of *Caulobacter crescentus*



# Bacteria senesce II

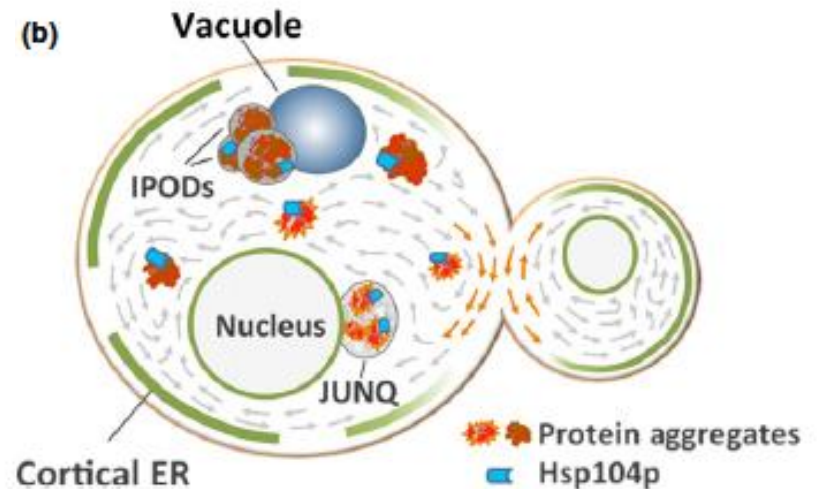
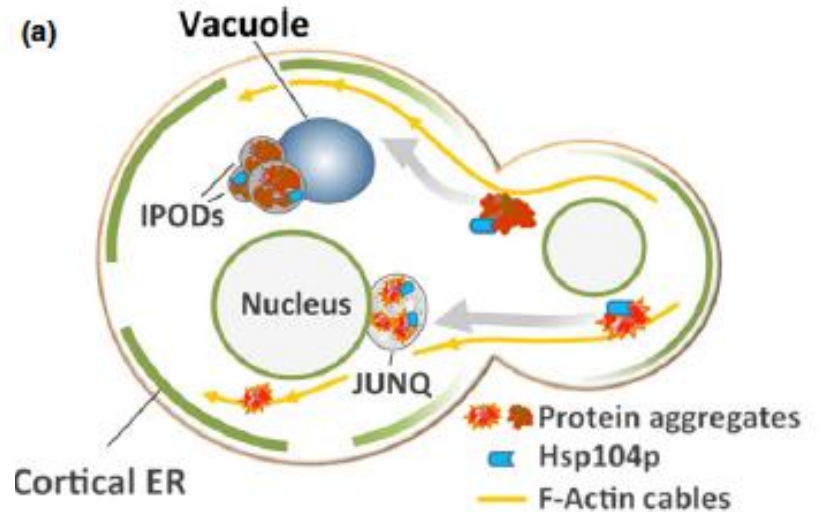
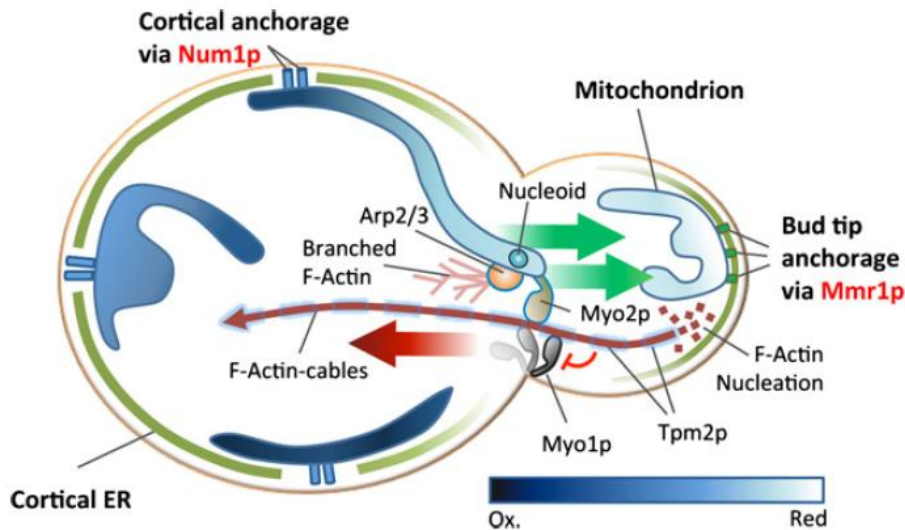


# Bacteria senesce—Theory





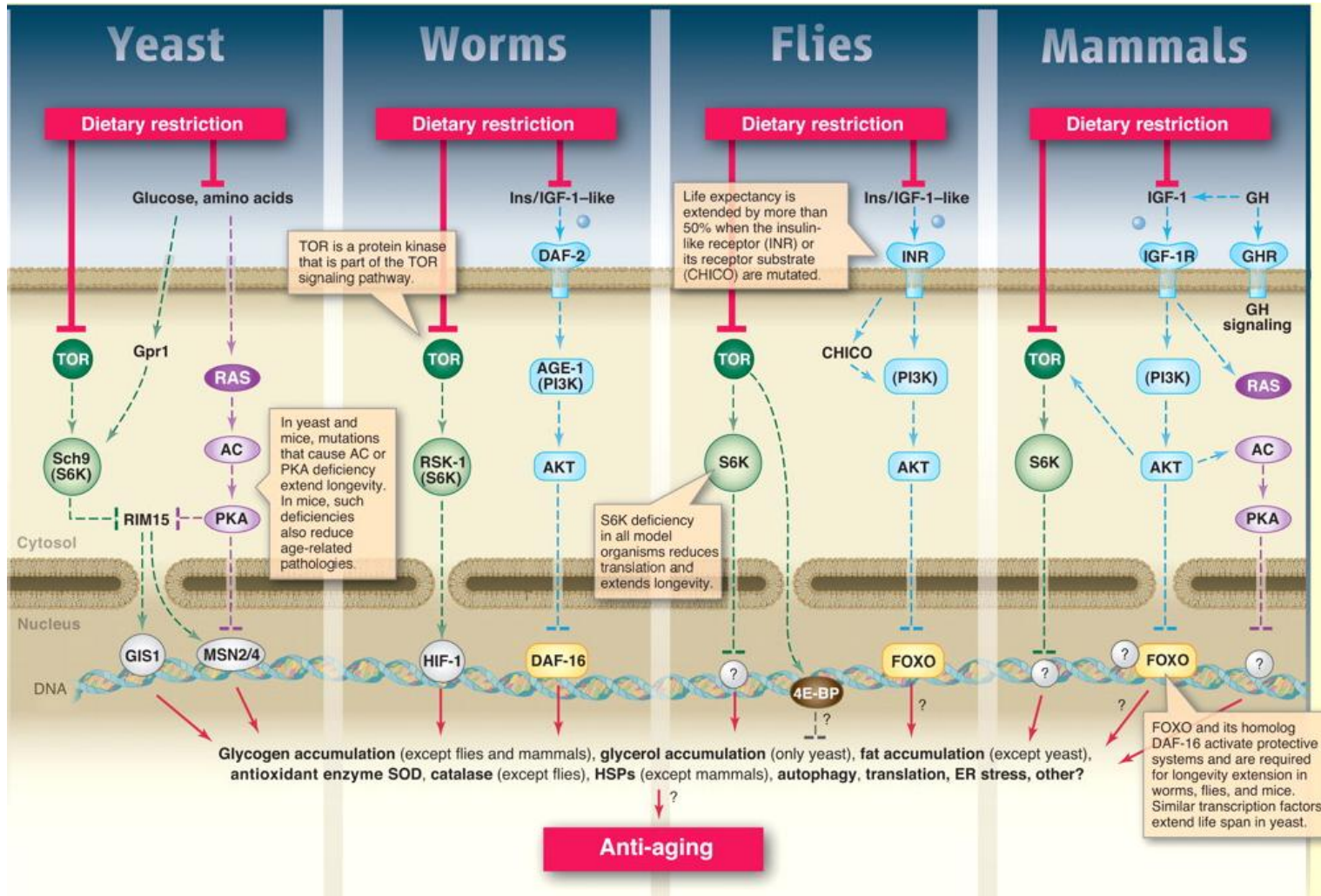
# Retrograde flow and selection



# The Hallmarks of Aging

Carlos López-Otín,<sup>1</sup> María A. Blasco,<sup>2</sup> Linda Partridge,<sup>3,4</sup> Manuel Serrano,<sup>5,\*</sup> and Guido Kroemer<sup>6,7,8,9,10</sup>





L Fontana et al. Science 2010;328:321-326



# Historical Conclusions

Based on evolutionary and molecular studies:

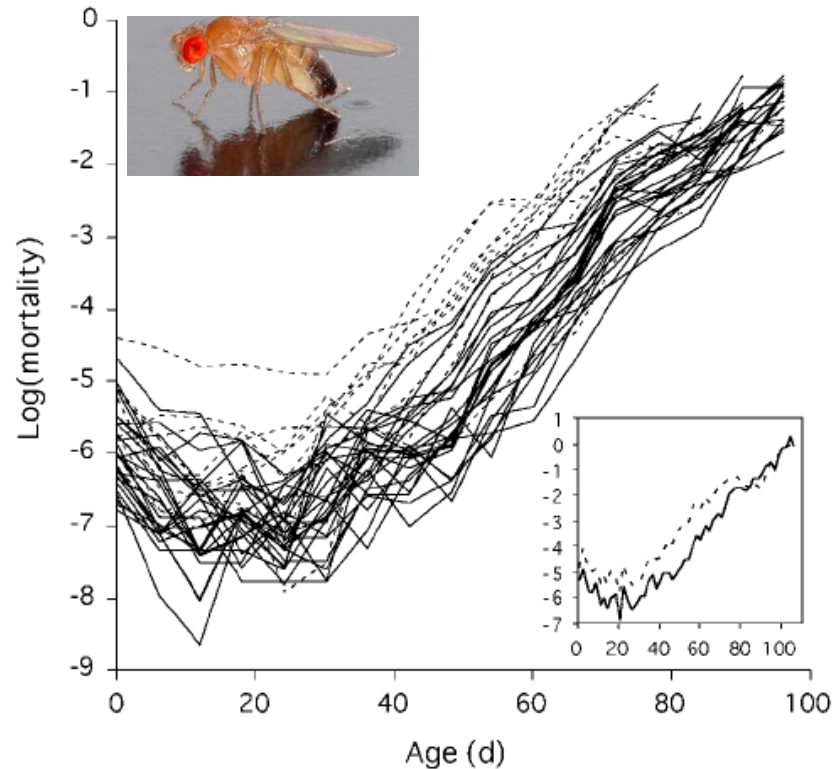
- Aging has a genetic basis
- Longevity influenced by genes with major effects
- Gene pathways and cellular targets of aging appear to be evolutionarily conserved

Challenge:

- Can we translate lab findings to the real world?

# Natural genetic variation

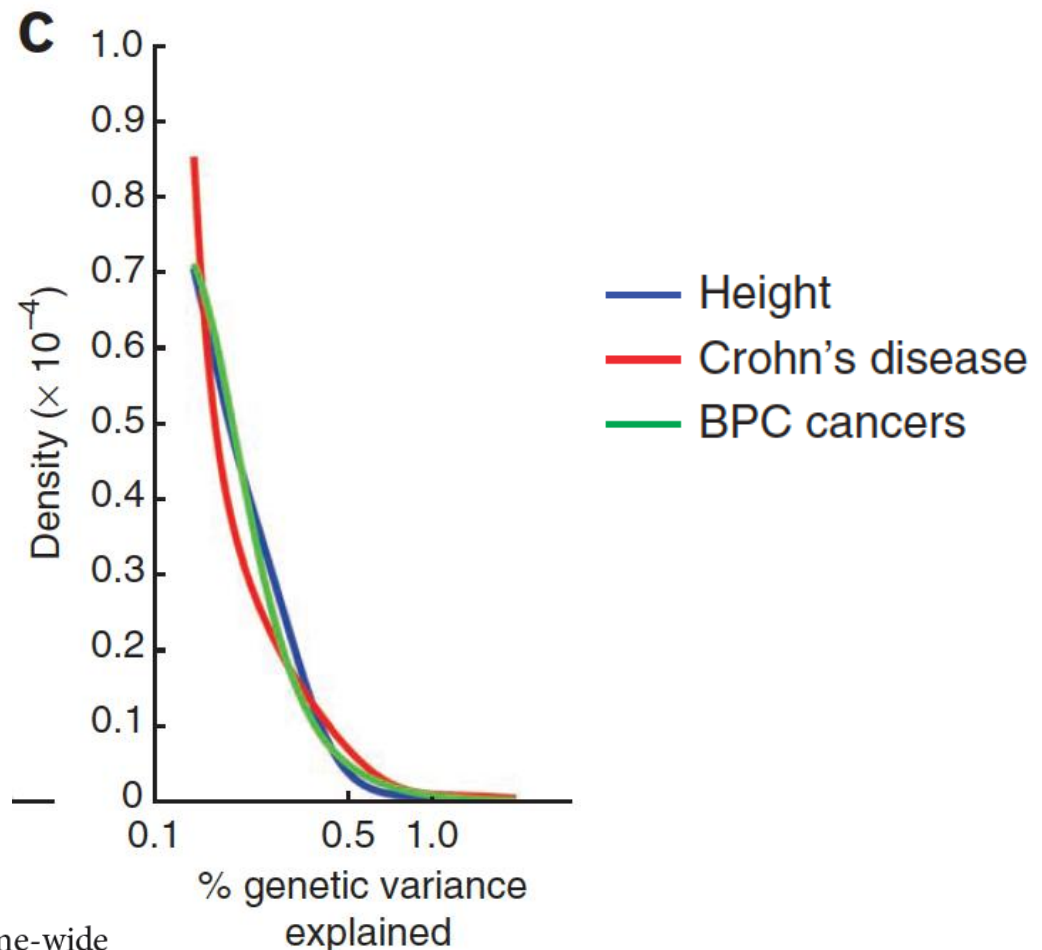
Lifespan is variable, heritable and polygenic



Snoke & Promislow, 2003



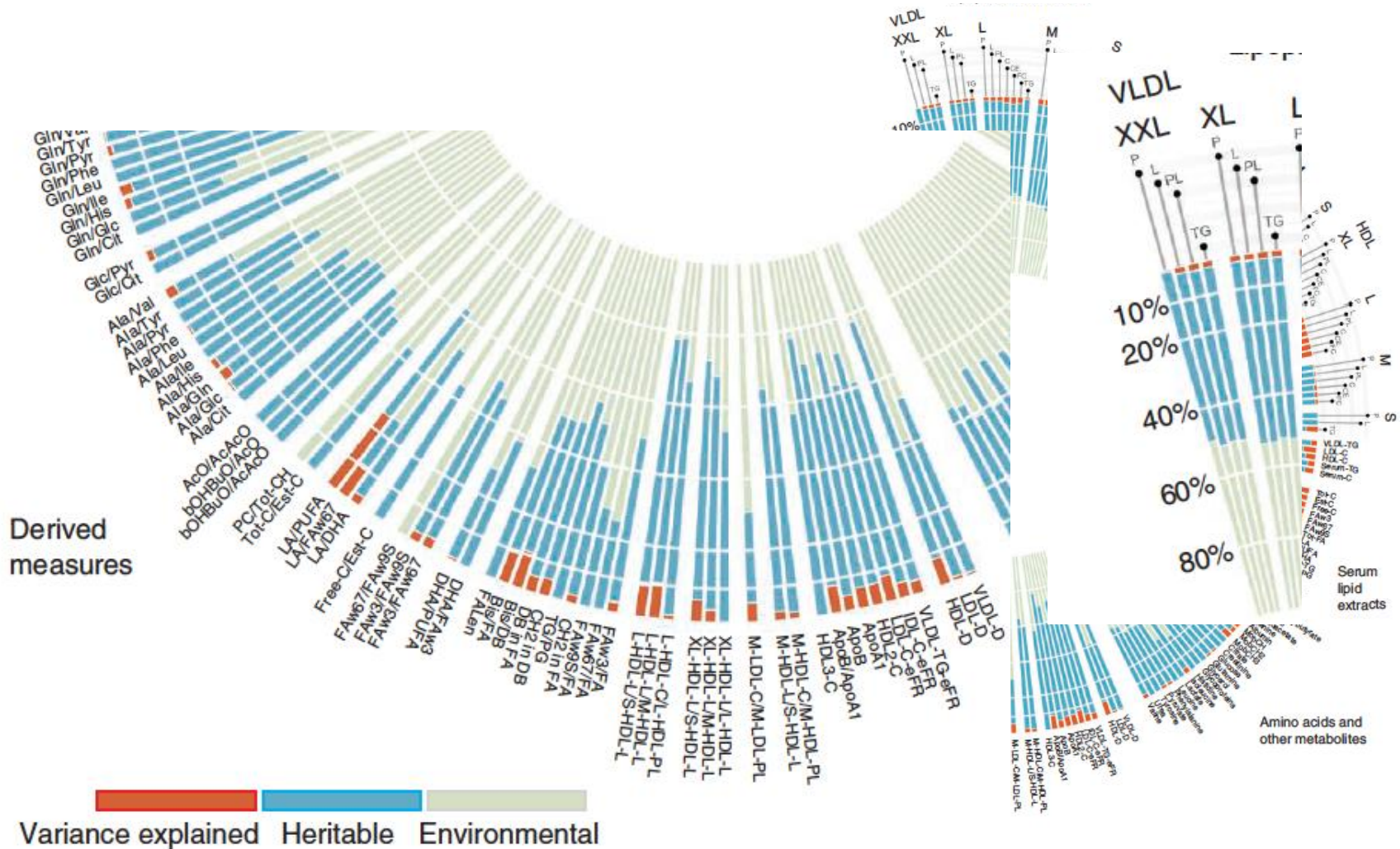
# Lab—genes of major effect Nature—genes of minor effect



Estimation of effect size distribution from genome-wide association studies and implications for future discoveries

VOLUME 42 | NUMBER 7 | JULY 2010 **NATURE GENETICS**

# Genome → Metabolome





# Human metabolic individuality in biomedical and pharmaceutical research

Karsten Suhre<sup>1,2,3</sup>, So-Youn Shin<sup>4\*</sup>, Ann-Kristin Petersen<sup>5\*</sup>, Robert P. Mohney<sup>6</sup>, David Meredith<sup>7</sup>, Brigitte Wägele<sup>1,8</sup>, Elisabeth Altmaier<sup>1</sup>, CARDIoGRAM†, Panos Deloukas<sup>4</sup>, Jeanette Erdmann<sup>9</sup>, Elin Grundberg<sup>4,10</sup>, Christopher J. Hammond<sup>10</sup>, Martin Hrabě de Angelis<sup>11,12</sup>, Gabi Kastenmüller<sup>1</sup>, Anna Köttgen<sup>13</sup>, Florian Kronenberg<sup>14</sup>, Massimo Mangino<sup>10</sup>, Christa Meisinger<sup>15</sup>, Thomas Meitinger<sup>16,17</sup>, Hans-Werner Mewes<sup>1,8</sup>, Michael V. Milburn<sup>6</sup>, Cornelia Prehn<sup>11</sup>, Johannes Raffler<sup>1,2</sup>, Janina S. Ried<sup>5</sup>, Werner Römisch-Margl<sup>1</sup>, Nilesh J. Samani<sup>18</sup>, Kerrin S. Small<sup>10</sup>, H.-Erich Wichmann<sup>19,20,21</sup>, Guangju Zhai<sup>10</sup>, Thomas Illig<sup>22</sup>, Tim D. Spector<sup>10</sup>, Jerzy Adamski<sup>11,12</sup>, Nicole Soranzo<sup>4\*</sup> & Christian Gieger<sup>5\*</sup>

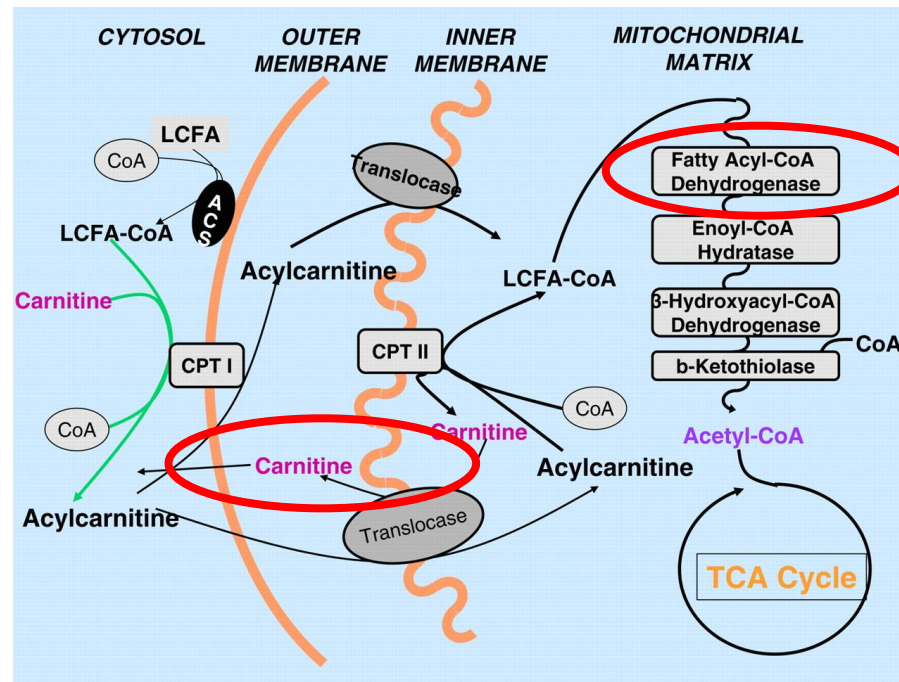
NATURE | VOL 477 | 1 SEPTEMBER 2011

for diabetes and cardiovascular disease. In all except three loci, the SNPs are common, with minor allele frequencies greater than 10%. In 25 cases, the effect size per allele copy is larger than 10%, and up to 60% in the case of the acyl-CoA dehydrogenase (*ACADS*) locus.

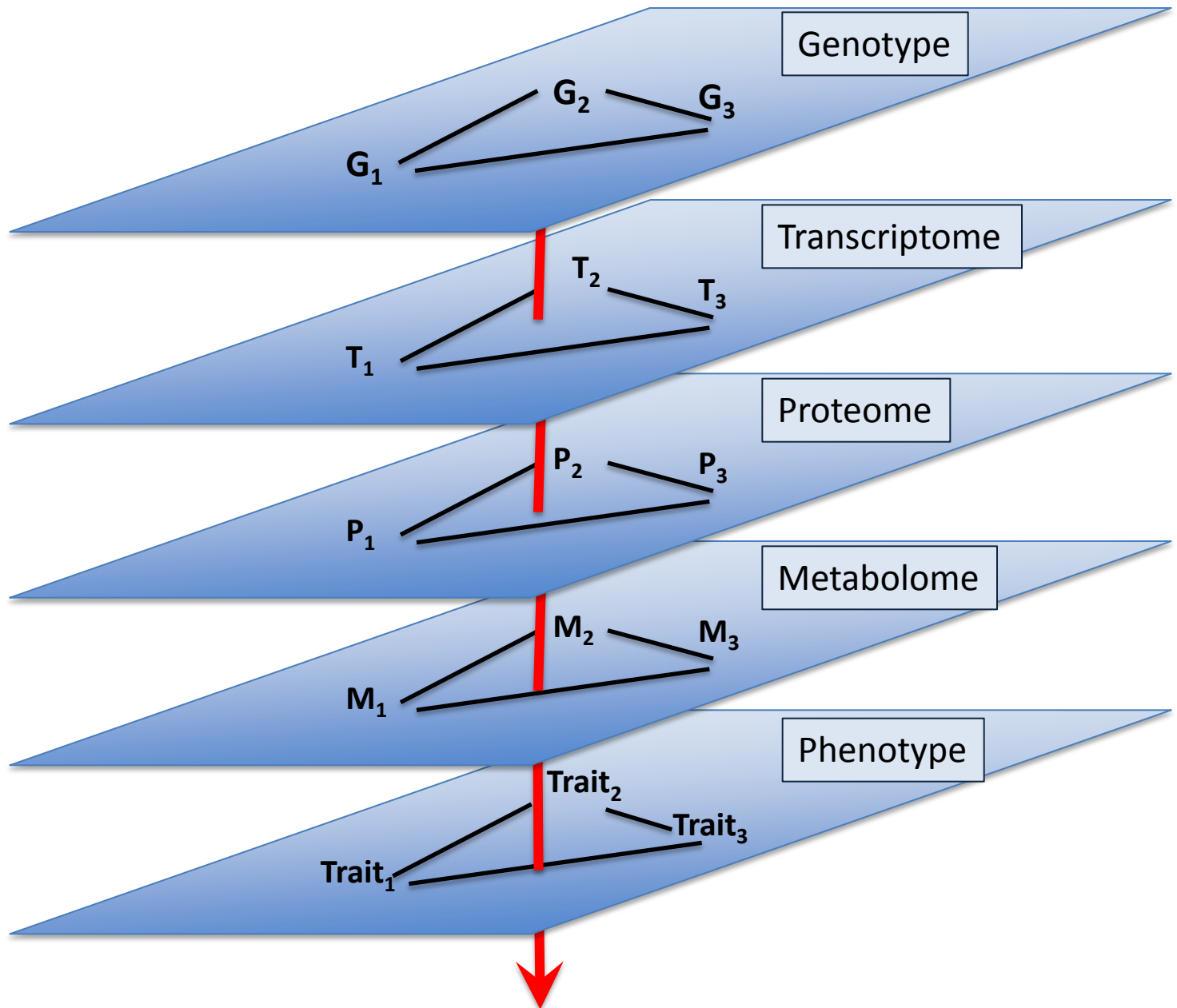
ACADS  
rs2066938

Butyrylcarnitine/propionylcarnitine

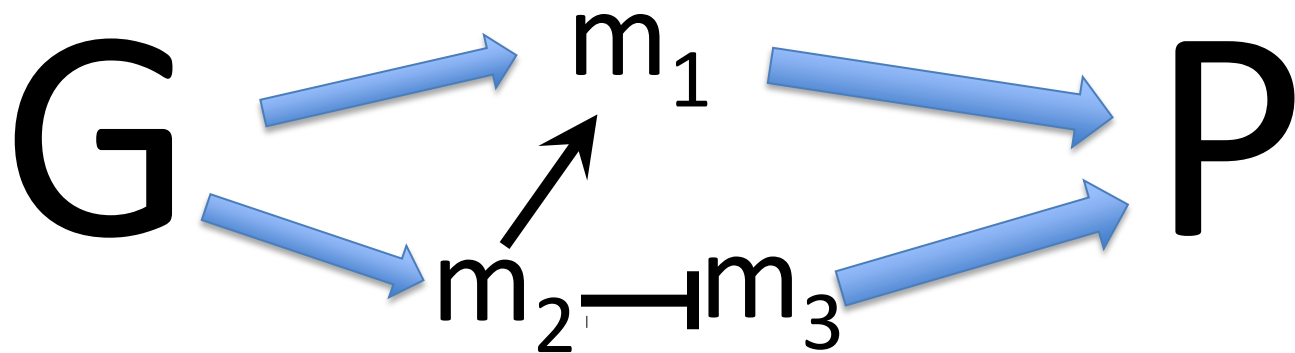
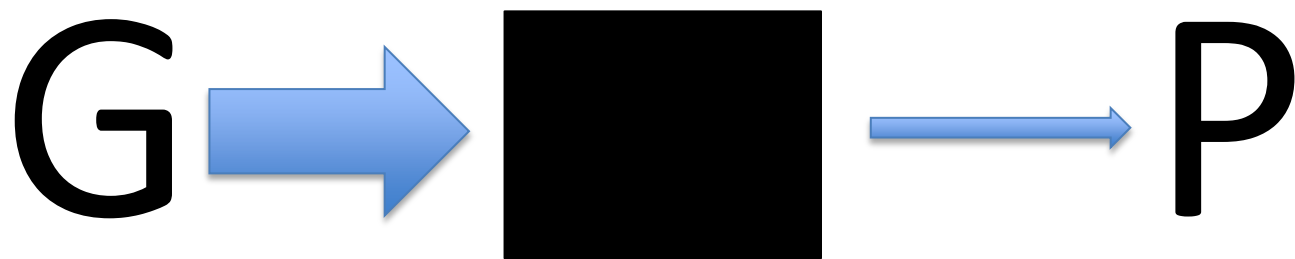
$< 4.4 \times 10^{-305}$



Environment



Life History Traits (survival and reproduction)



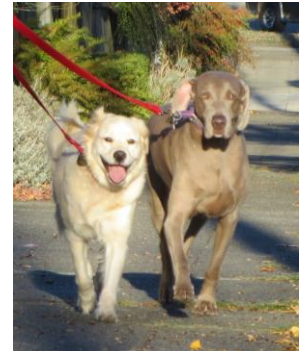
# Metabolomics and Aging

1. Phylogenetic variation
  - evolution of the metabolome
2. Genetic variation
  - Genetics, metabolomics, and aging
3. Environmental variation
  - Diet restriction and metabolomic networks

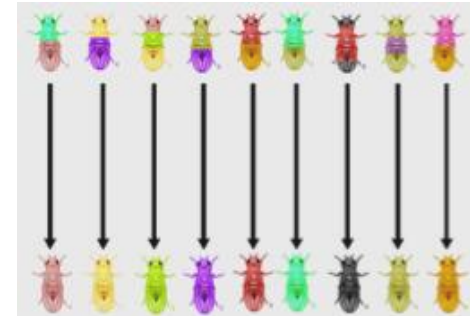
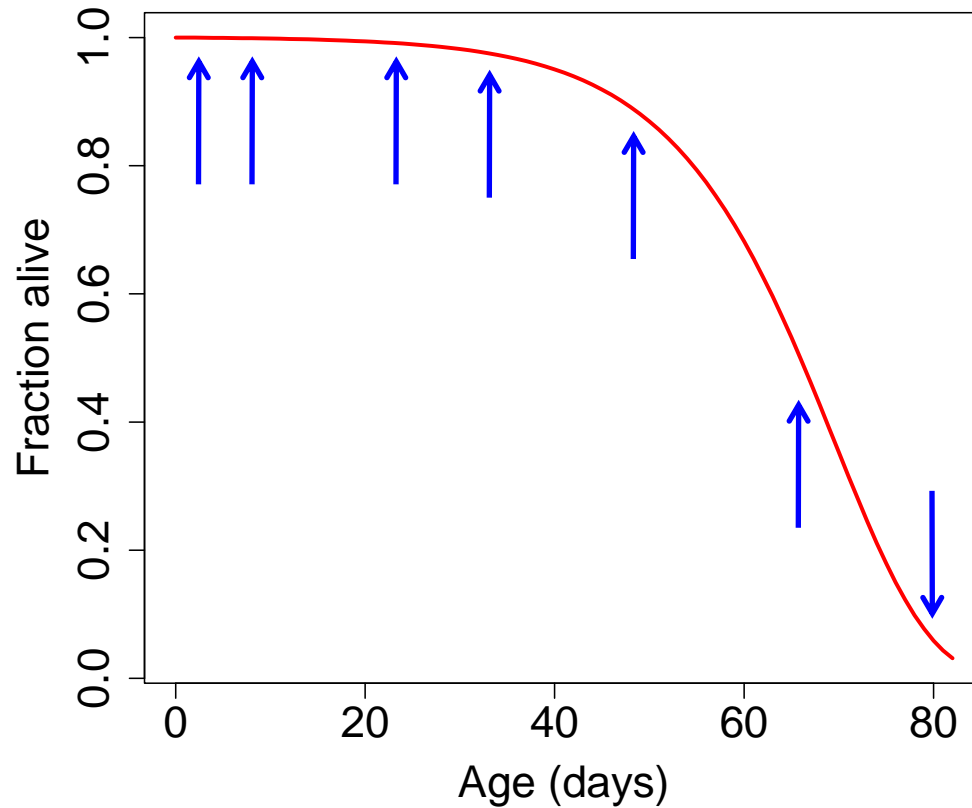


# Metabolomics and Aging

1. Evolution of the metabolome
2. Genetics, metabolomics, and biomarkers of aging
3. Diet restriction and metabolomic networks



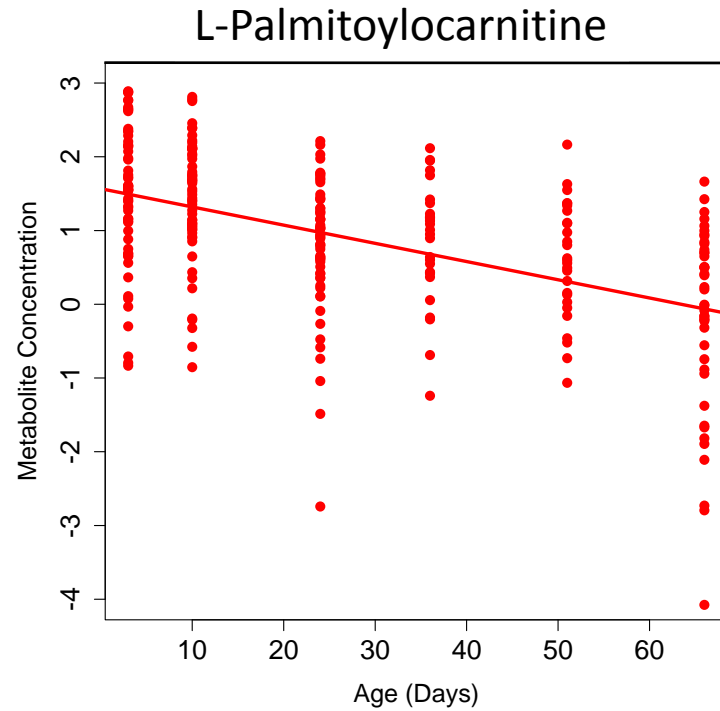
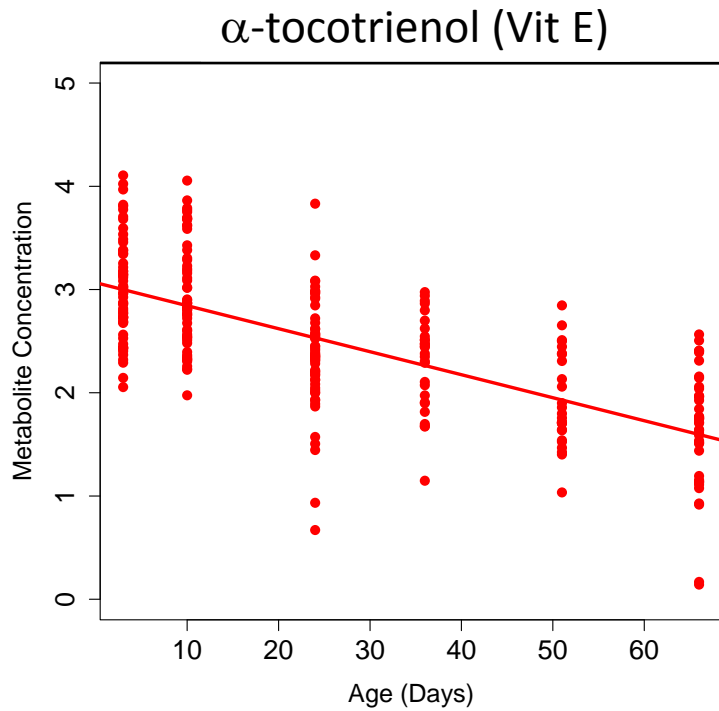
# Effect of aging on the metabolome in the *Drosophila* Genome Reference Panel?



Assay three key factors that affect survival:

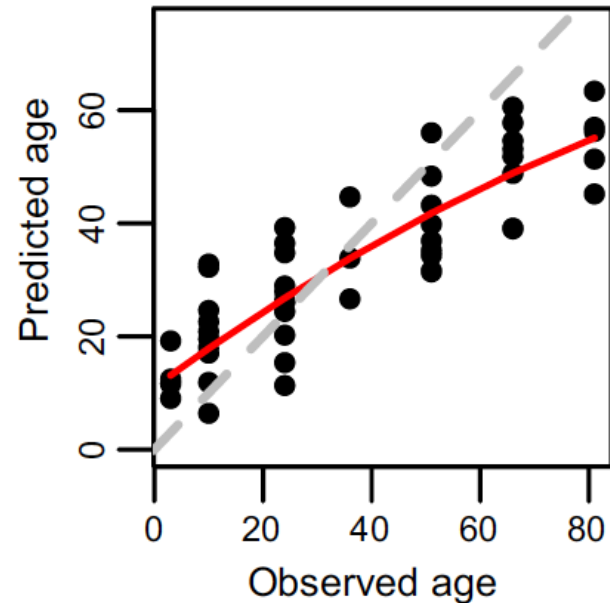
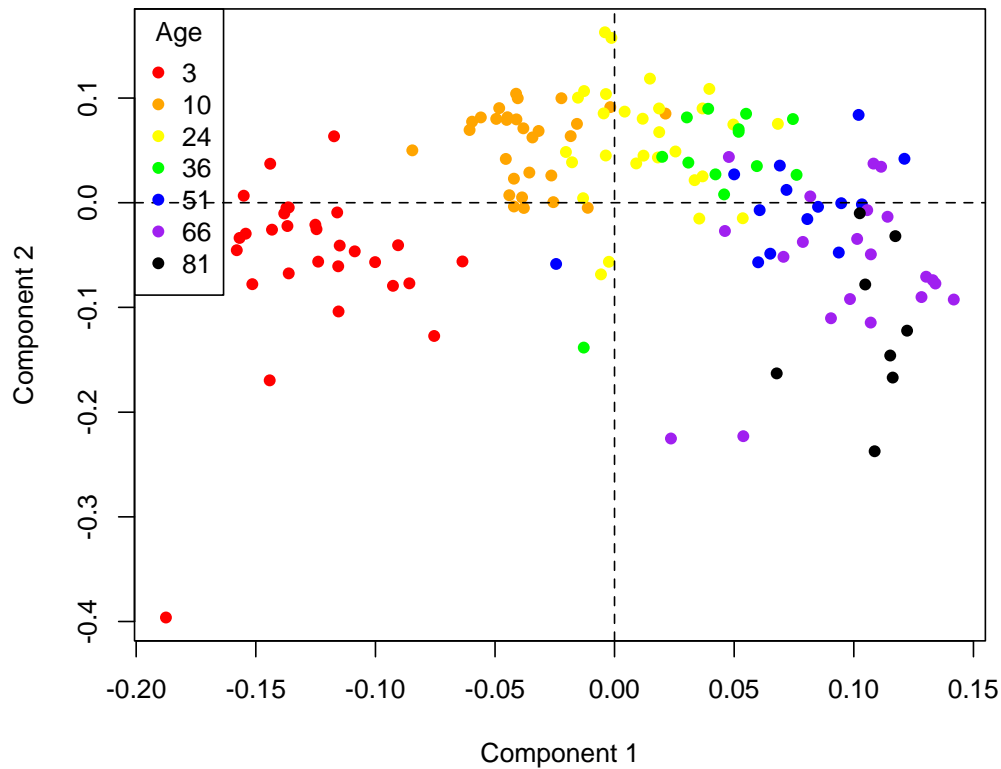
- Sex
- Age (7 ages)
- Genotype (15 inbred lines)
- Biological and technical replicates (~600 samples)

# Effects of age

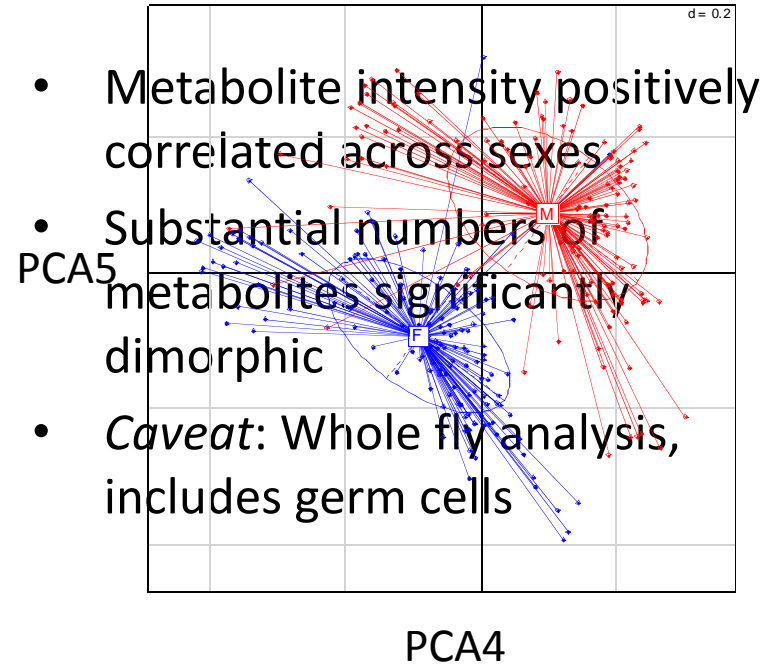
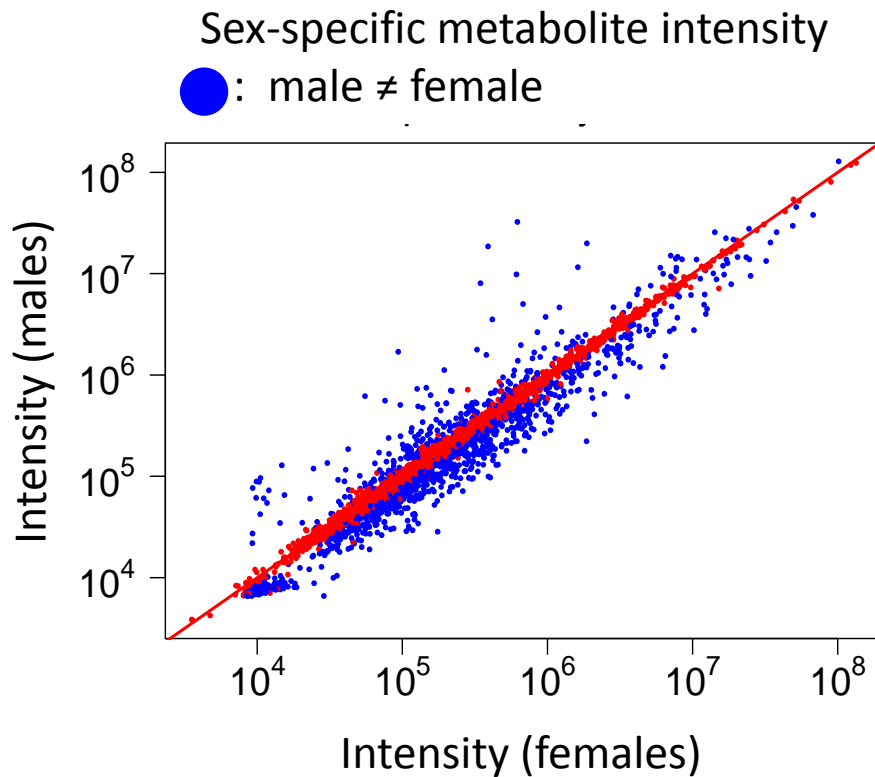




# The *Drosophila* metabolome is age-specific

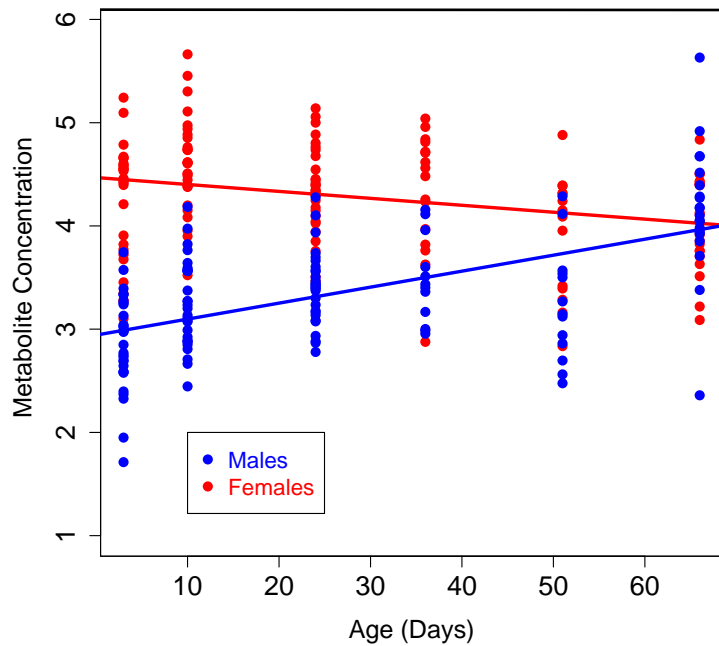


# The *Drosophila* metabolome is sexually dimorphic

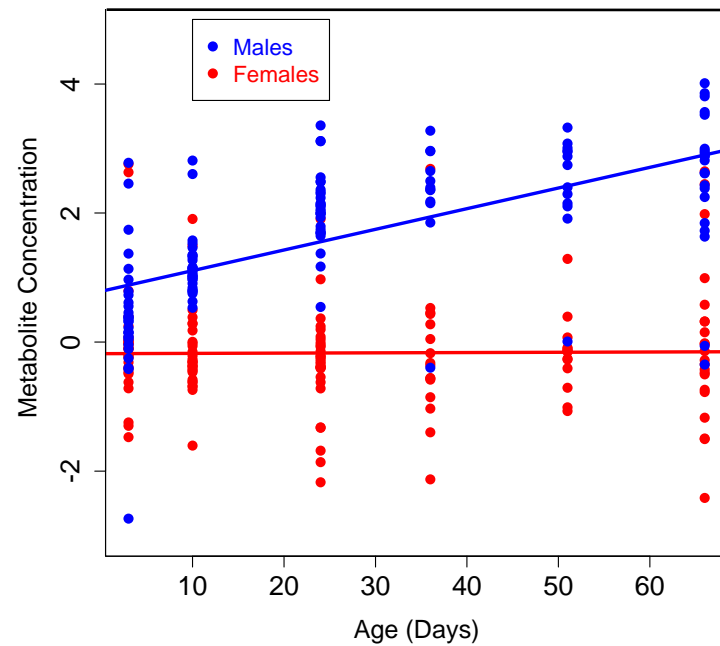


# Age-Sex interactions

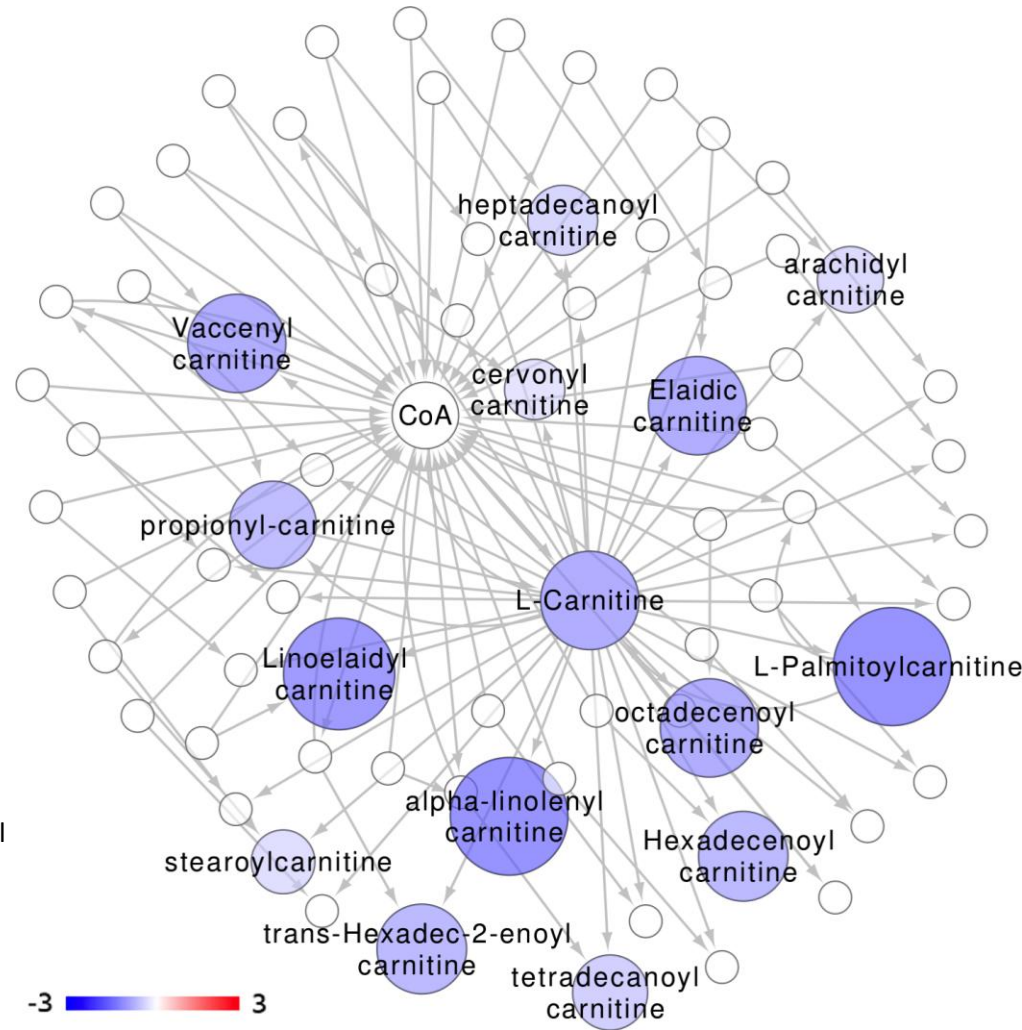
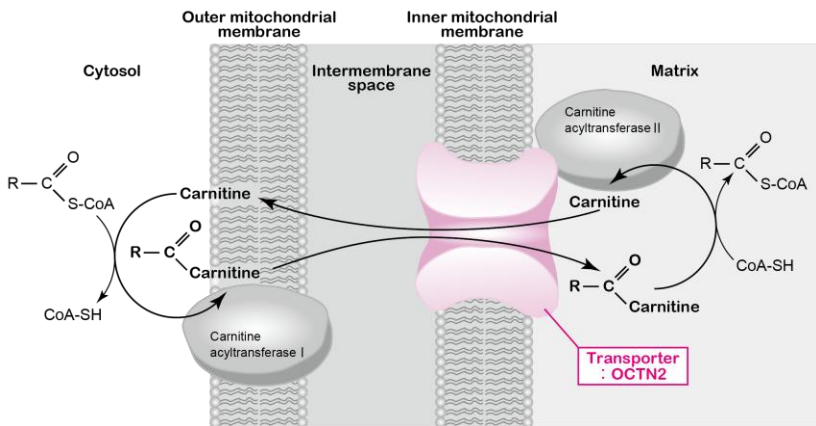
5-dehydro avenasterol (plant sterol)



Pantothenate (CoA biosynthesis)

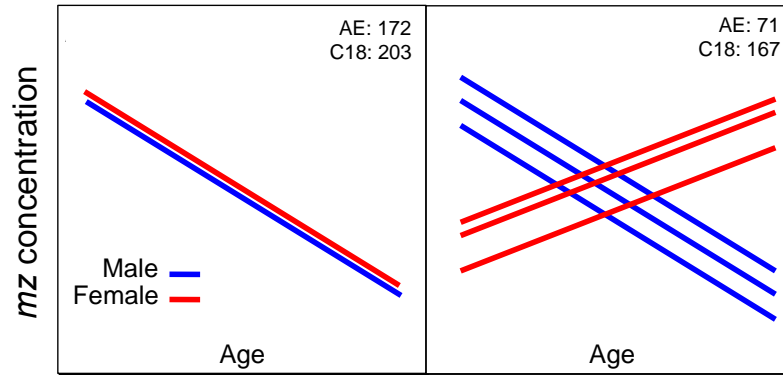


# Fatty acid metabolism

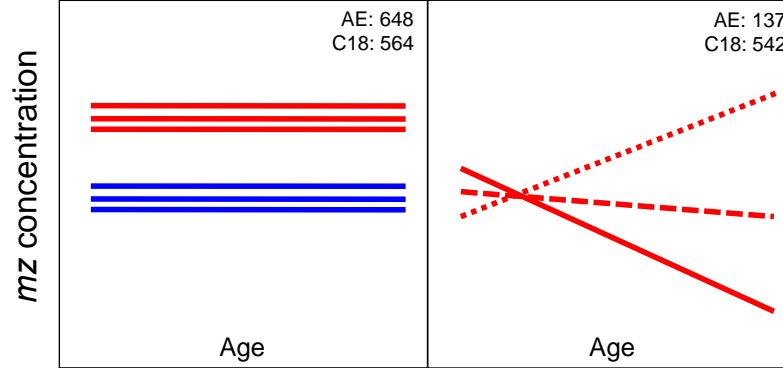


[http://www.kanazawa-u.ac.jp/research\\_bulletin/feature201309F003.html](http://www.kanazawa-u.ac.jp/research_bulletin/feature201309F003.html)

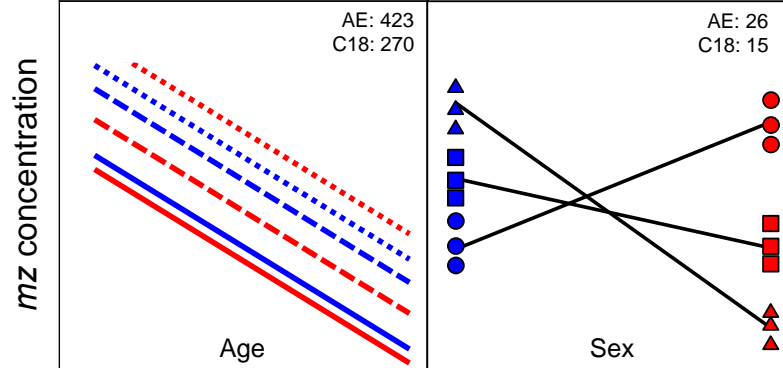
Age



Sex



Genotype

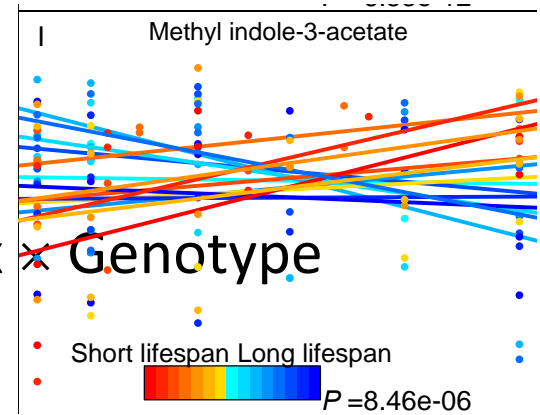


Age × Sex

- Fatty acid metabolism
- Neurotransmitters
- Amino acids
- Glycerophospholipids

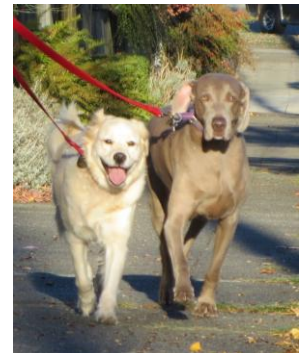
Age × Genotype

Sex × Genotype



# Metabolomics and Aging

1. Evolution of the metabolome
2. Genetics, metabolomics, and biomarkers of aging
3. Diet restriction and metabolomic networks



# Metabolomics and Diet Restriction



Available online at [www.sciencedirect.com](http://www.sciencedirect.com)

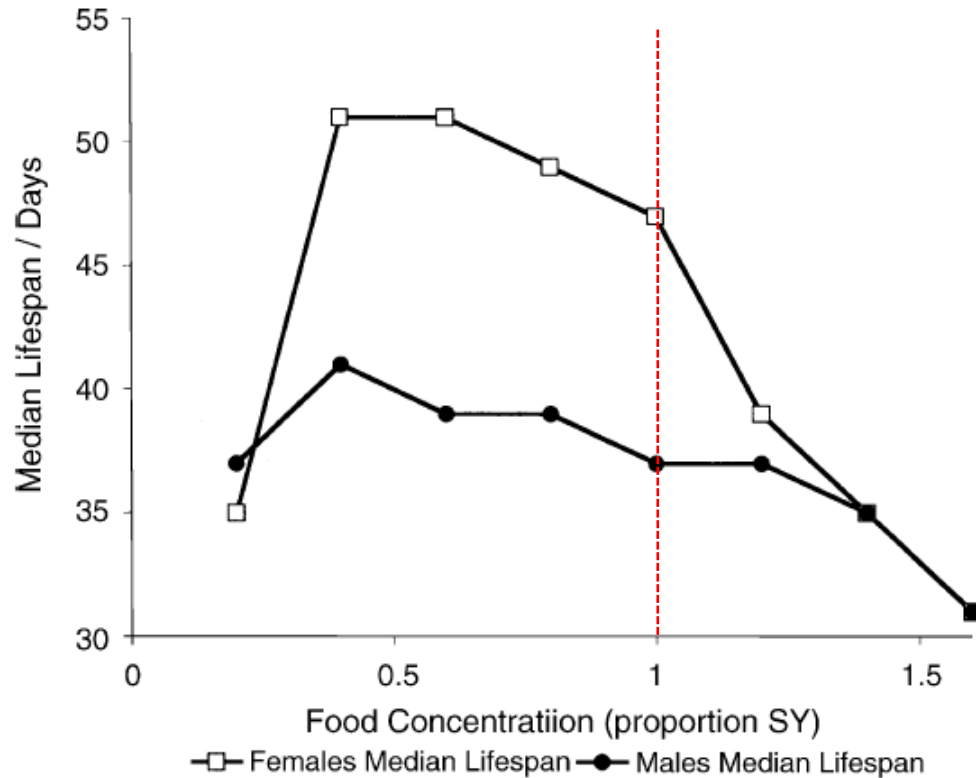
SCIENCE @ DIRECT®

mechanisms of ageing  
and development

Mechanisms of Ageing and Development 126 (2005) 938–950

## Dietary restriction in *Drosophila*

Linda Partridge\*, Matthew D.W. Piper, William Mair

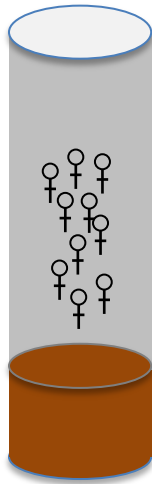




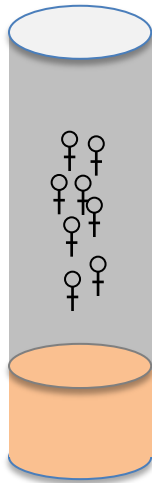
# Metabolomics & Diet Restriction

1. Does DR work by slowing aging of the metabolome?
2. Can metabolome networks point to mechanism?

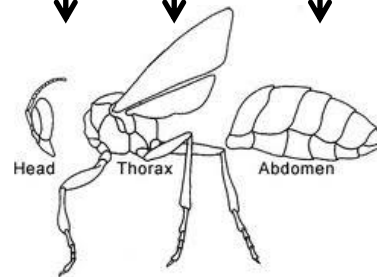
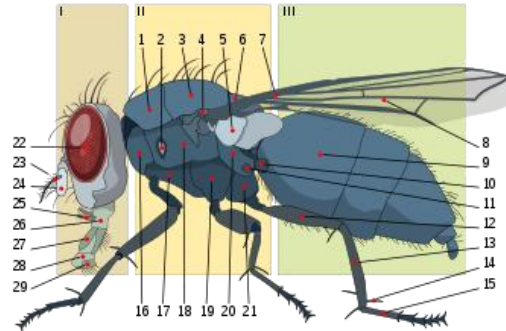
# Metabolomics and Diet Restriction



High yeast (AL) diet  
Collect age 10, 20, 40 d



Low yeast (DR) diet  
Collect age 10, 20, 40 d



X 6 samples/tissue

Buck Institute

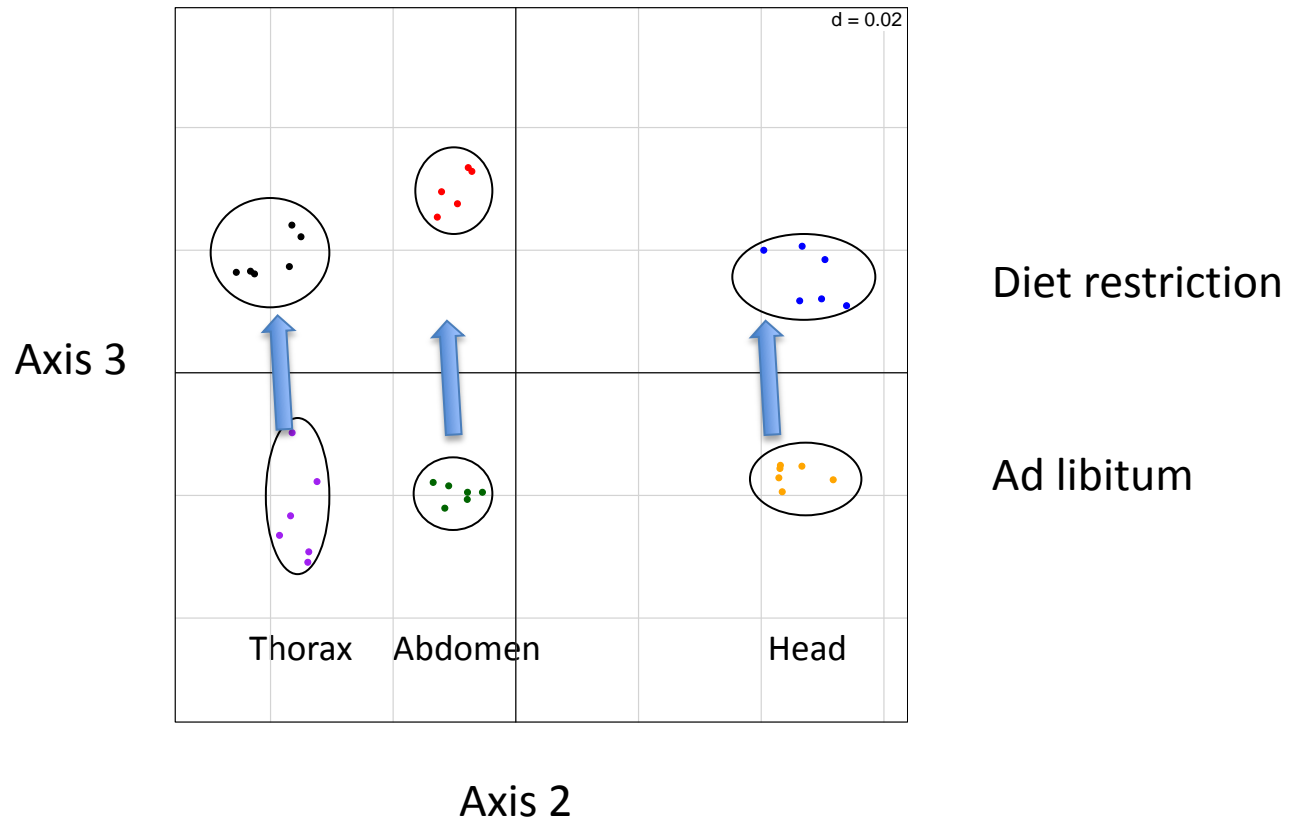


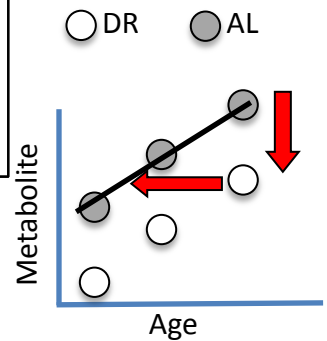
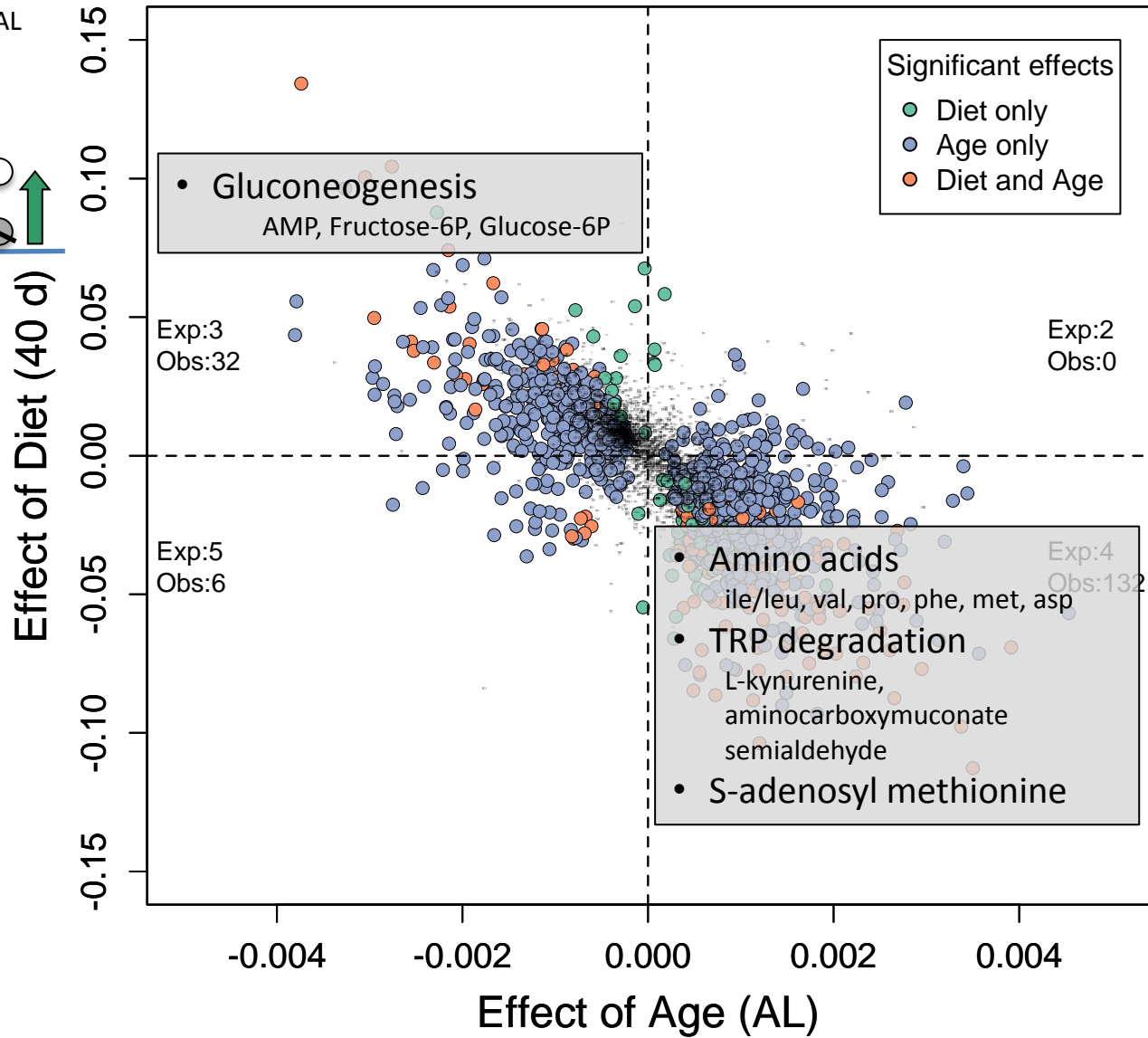
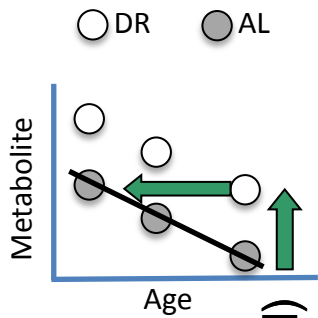
Pankaj  
Kapahi

Matt  
Laye

# Principle Component Analysis

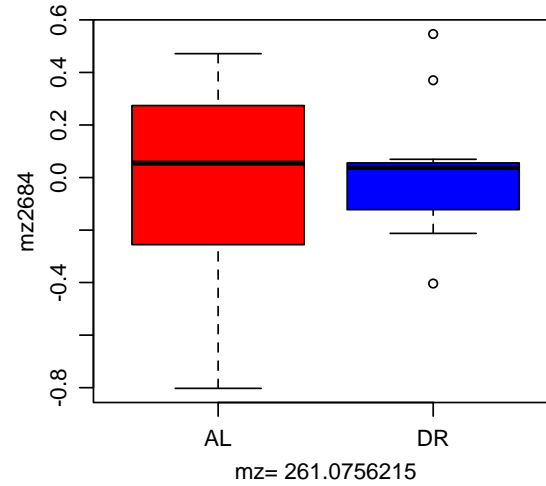
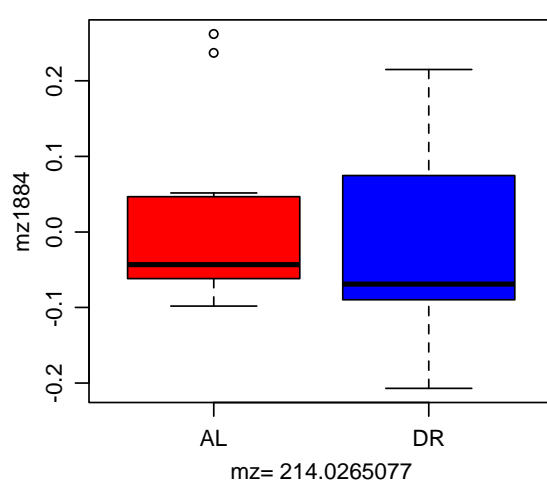
## DR and body part





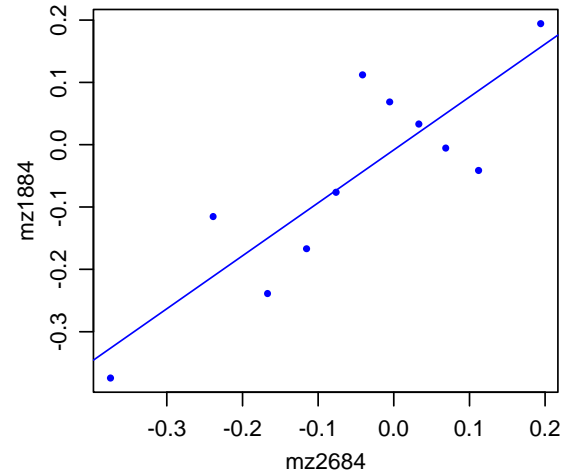
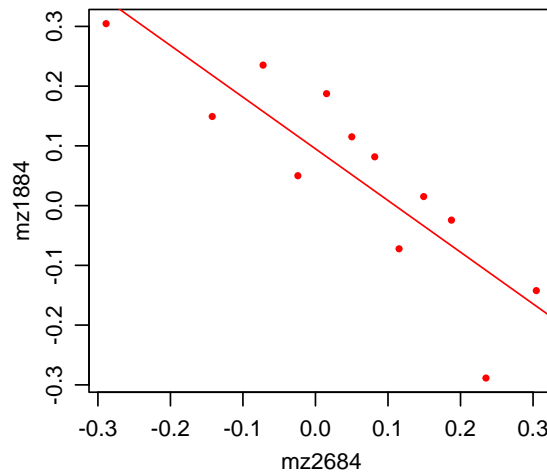
DR and  
metabolome network structure

# Correlations vs. Main Effects



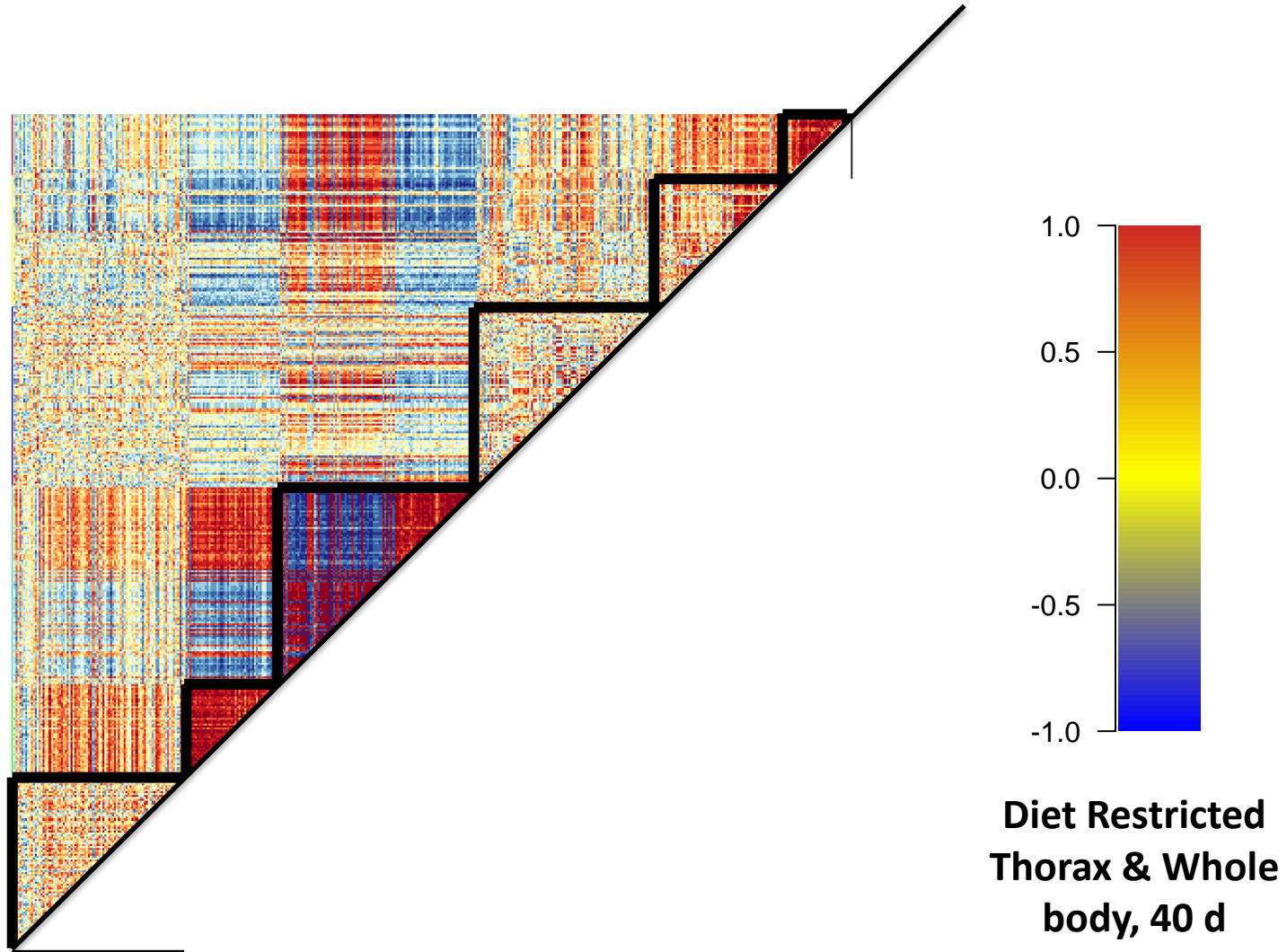
*Ad Lib*

*DR*

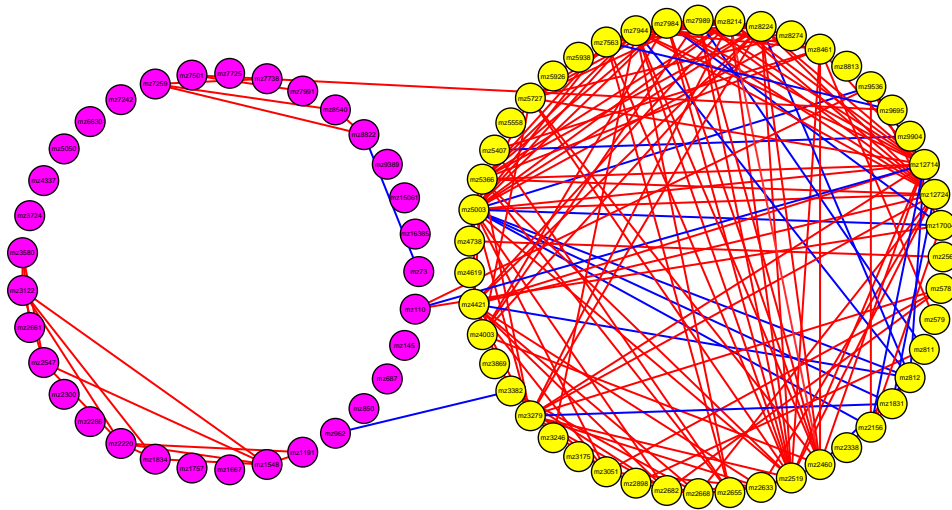


# WGCNA analysis of DR in flies

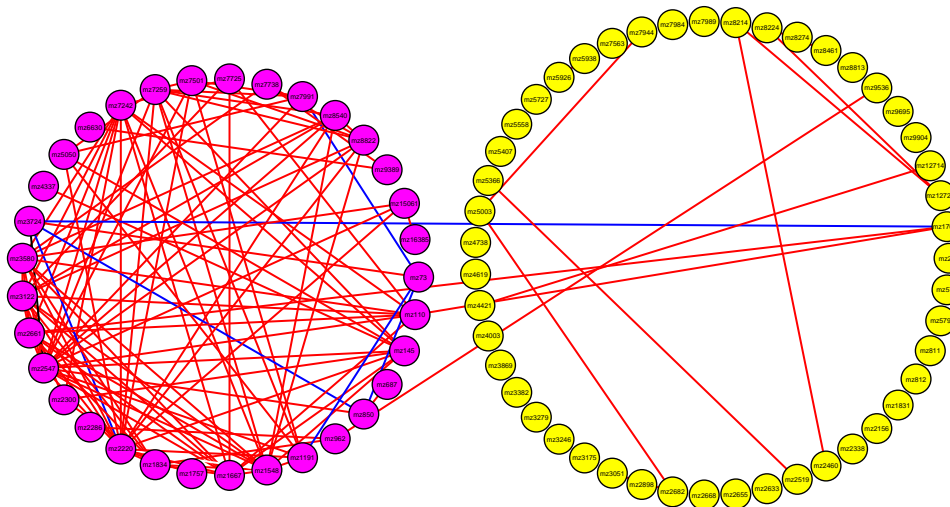
**Ad Lib Thorax &  
Whole body, 40 d**



# Some gain connections, some lose connections...



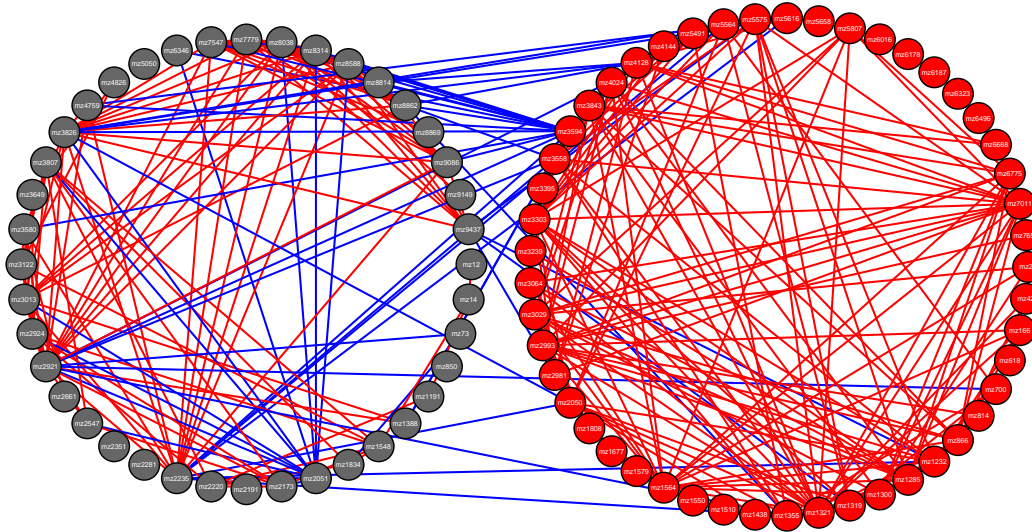
Thorax, Magenta/Yellow, AL



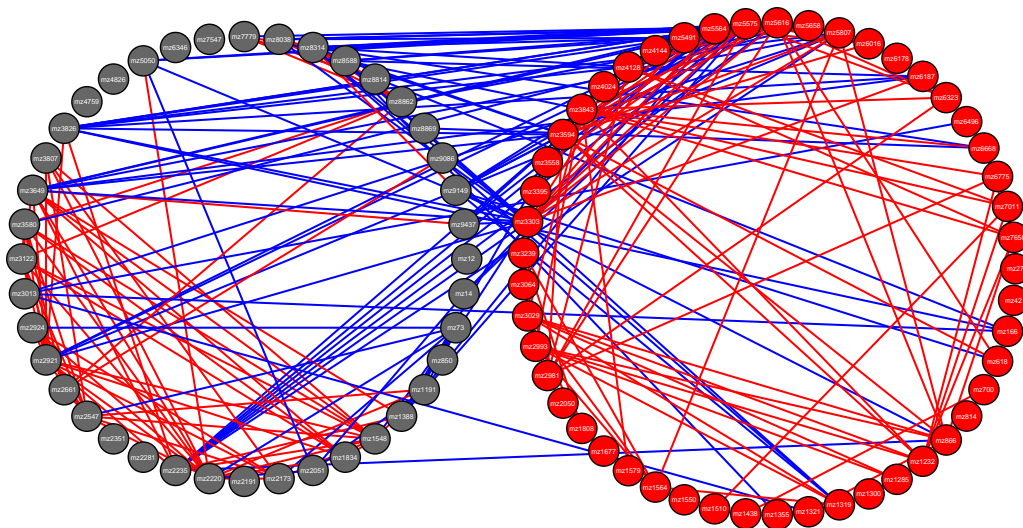
Thorax, Magenta/Yellow, DR



# ...and some stay the same.

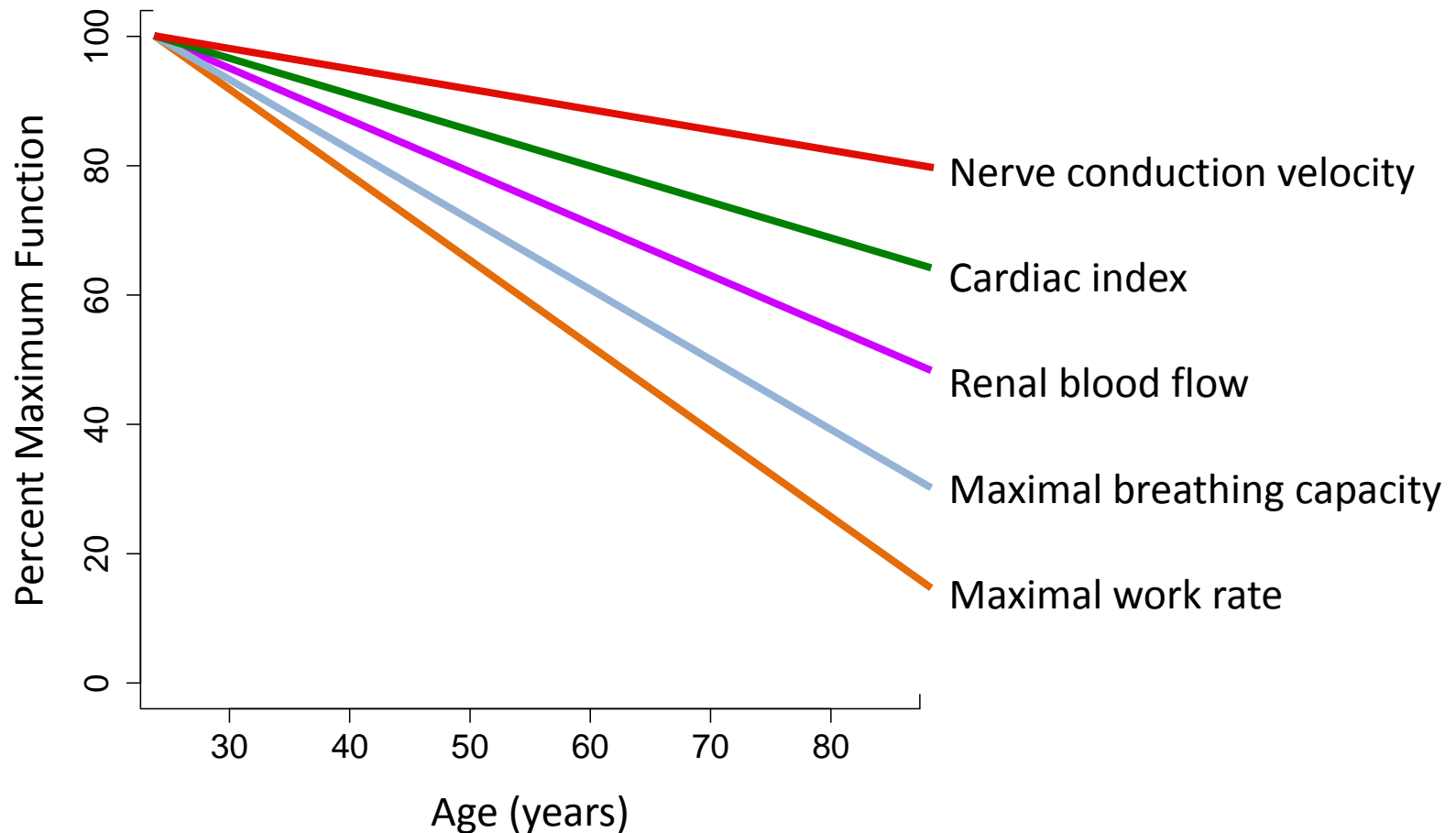


Thorax Red/Black, AL



Thorax Red/Black, DR

# Next step: Identify modules that age at different rates



After Baker & Martin in Geriatric Medicine (ed. Cassel *et al.*, 1997)



Emory University  
 Dean Jones  
 ViLinh Tran  
 Karan Uppal  
 Shuzhao Li

Buck Institute  
 Pankaj Kapahi  
 Matt Laye  
 Rachel Brem

Promislow lab

Devin Arbuthnott  
 William Gordon  
 Ram Hariharan  
 Jessica Hoffman  
 Kelly Jiang  
 Forrest Nussdorfer  
 Adrienne Wang

Nick Force  
 Erika Gajda  
 Jessica Jang  
 Jake Mouser  
 Sharon Ornelas  
 Romeo Quach  
 Jake Mouser  
 Eric Vanderbilt-  
 Matthews

Ariana Samuelson  
 Whitney Sharp  
 Katie Strehler  
 Quynh Tran  
 Cindy Tseng  
 Erin Tudor  
 Deborah Xi  
 Nicole Bergman  
 Ijay Okeke

UW  
 Peter Hoff  
 Matt Kaeberlein  
 Mike MacCoss  
 Leo Pallanck  
 Peter Rabinovitch

Aberdeen/Beijing  
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