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**Boulder, CO**

**Genetic Dissection of Hormesis:  
Ponce d'elegans**

# OVERVIEW

**How can an invertebrate be used to study hormesis?**

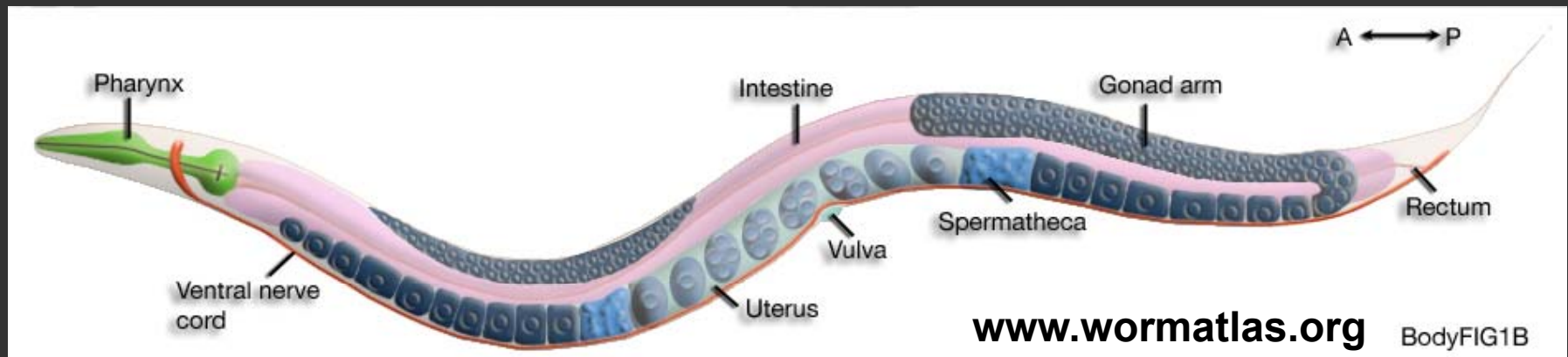
**Introduction to hormesis in the worm *C. elegans***

- **Genetic dissection of heat hormesis**
- **Individual worm response to heat as a predictor of longevity**
- **Heritability of worms' response to heat hormesis**

# •THE NEMATODE WORM MODEL SYSTEM: *C. elegans*

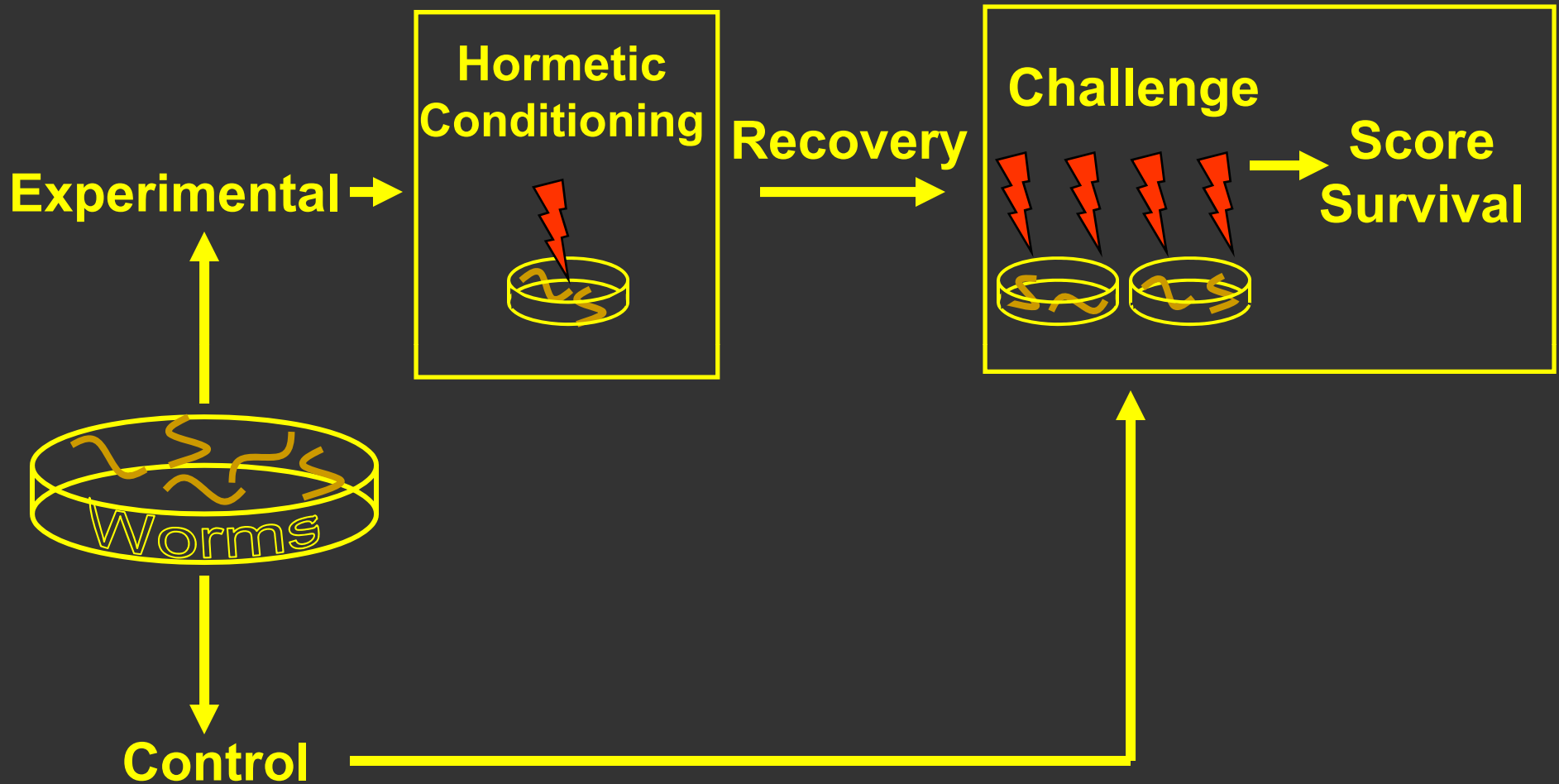
## Anatomy:

- Only 959 somatic cells in hermaphrodites (self-fertilizing!)
- Major tissues: muscle, hypodermis, neurons, gonads, gut
- Can visualize changes in tissues after stress / aging

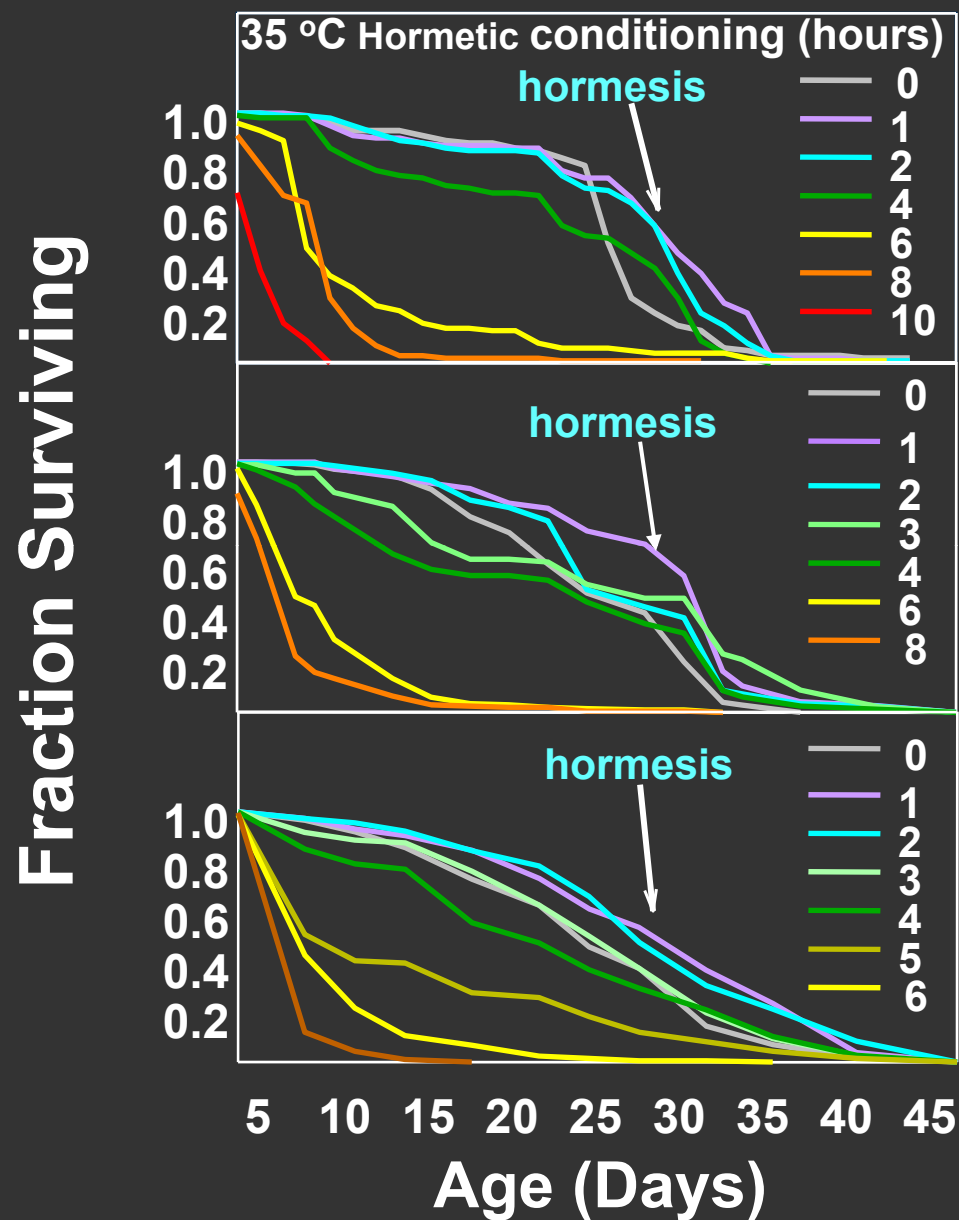




# HOW TO TEST FOR HORMESIS IN THE WORM



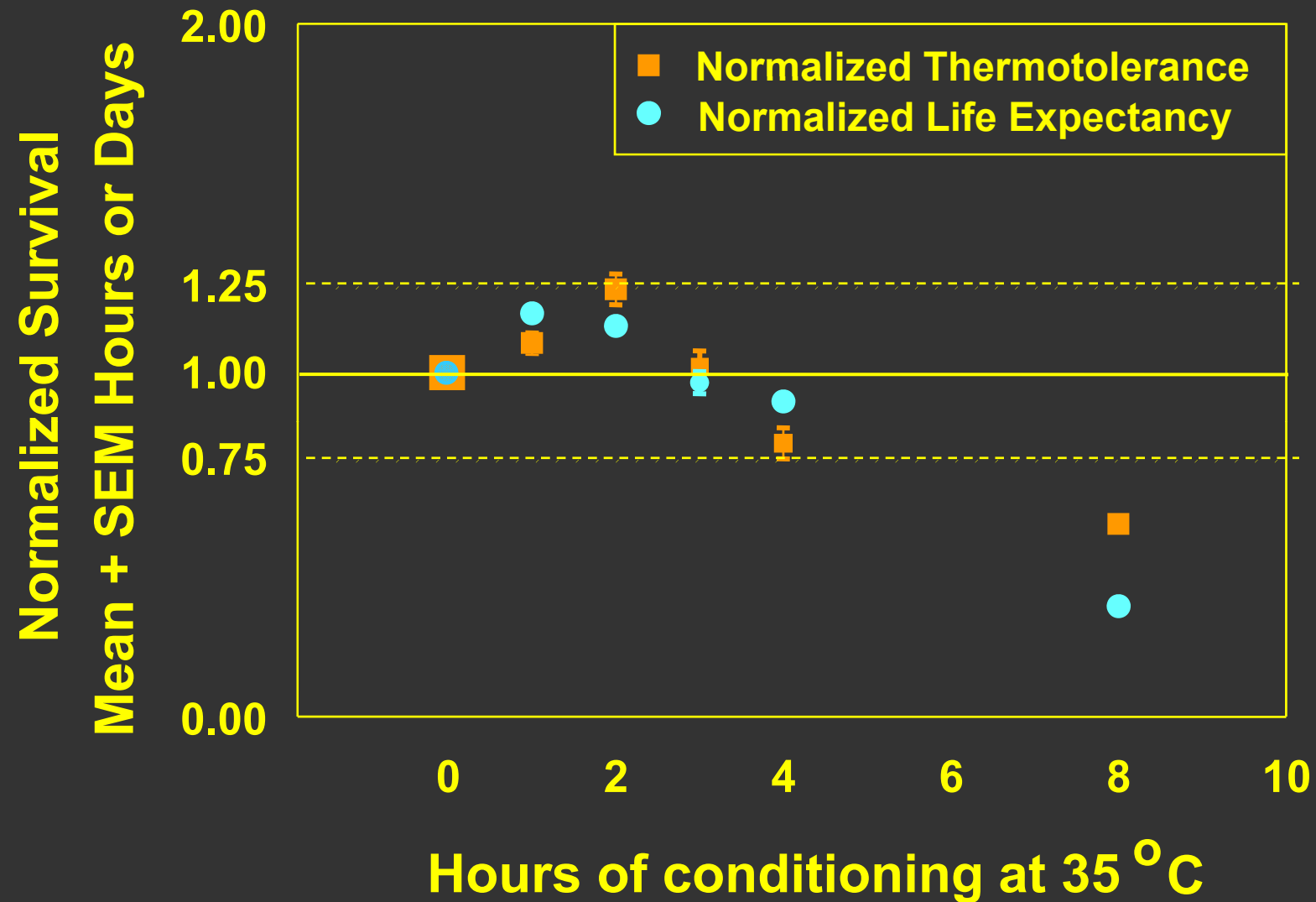
# SURVIVAL OF WORMS AFTER MILD HEAT STRESS



**HORMESIS CAN BE OBSERVED REPRODUCIBLY IN THE WORM**

*Michalski et al, Biogerontol. 2001*

# DOSE-RESPONSE CURVES OF HEAT HORMESIS IN THE WORM

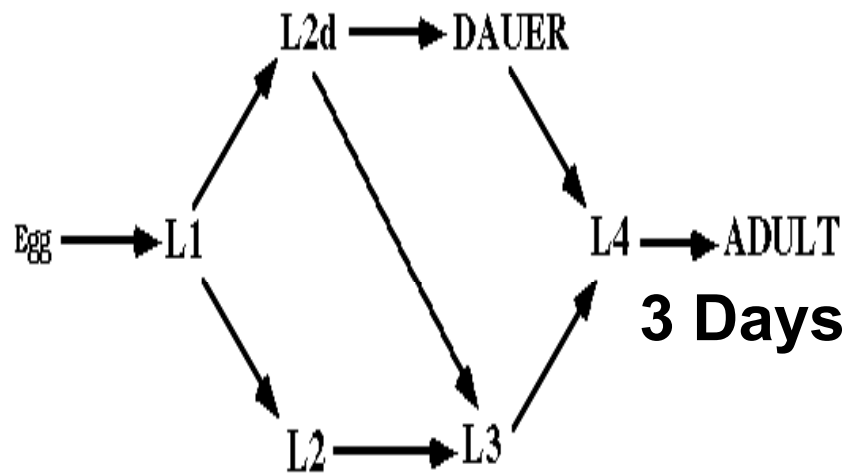


**THE WORM DISPLAYS PARALLEL  
HORMETIC THERMOTOLERANCE AND HORMETIC SLOWED AGING**

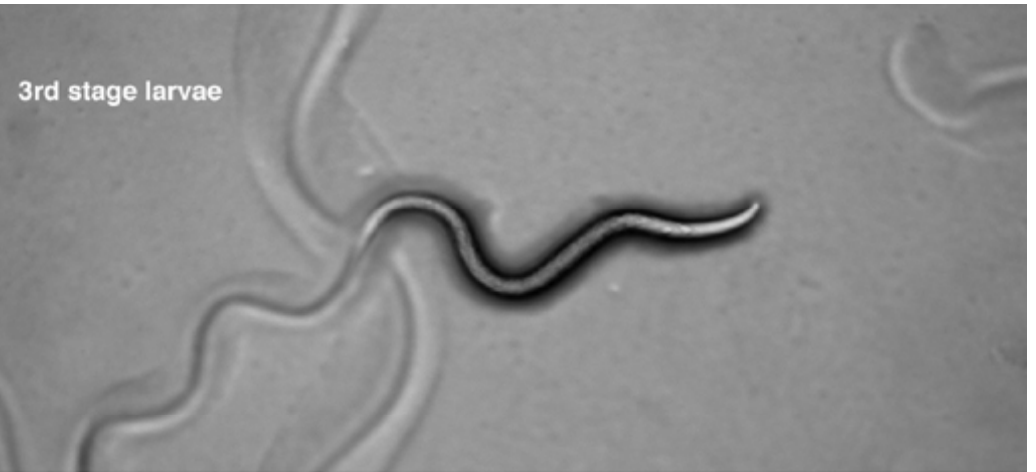
*Cypser et al, J. Gerontol. 2002*

# *Caenorhabditis elegans*: life cycle

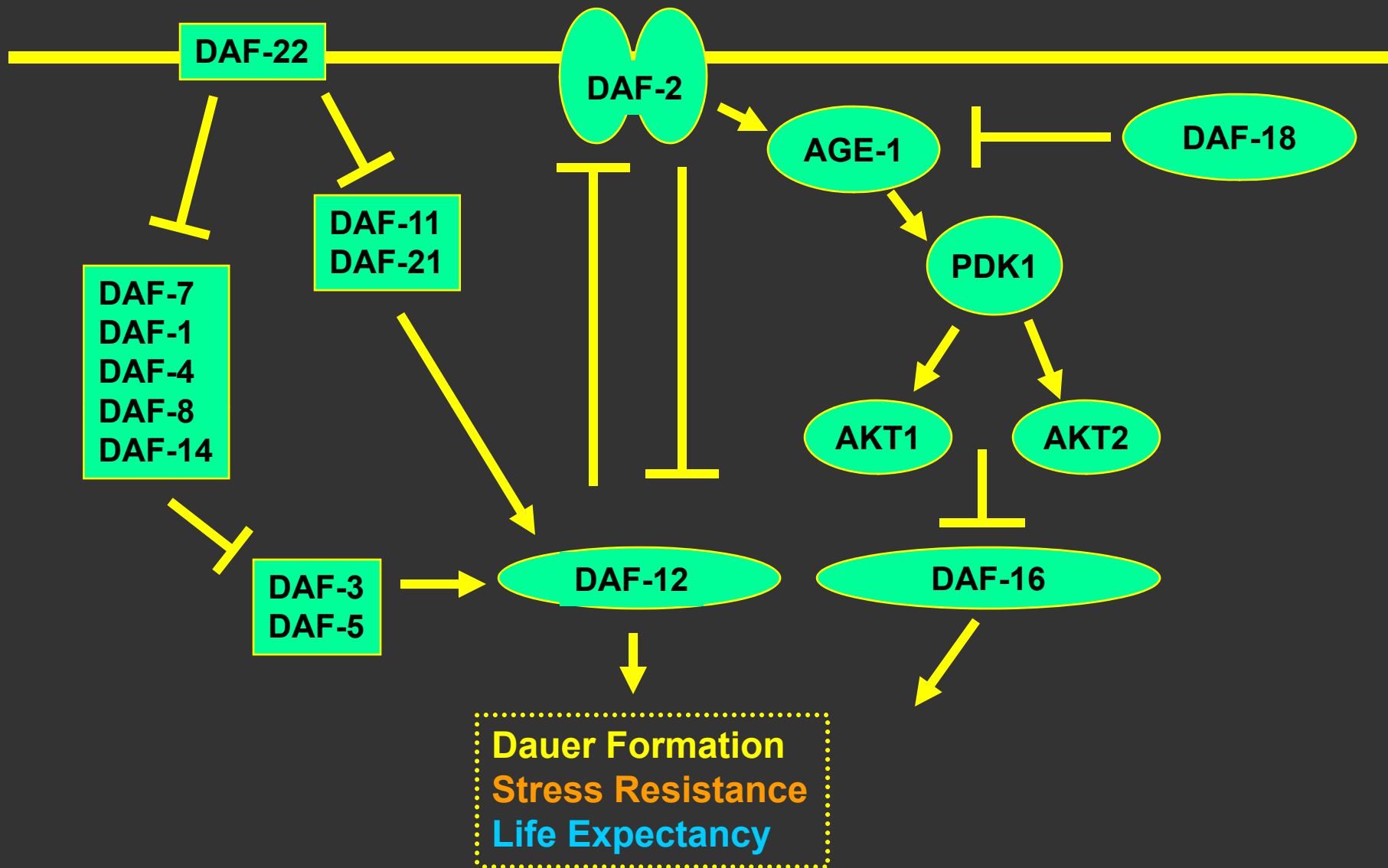
## Migratory (dauer) phase



## Reproductive phase



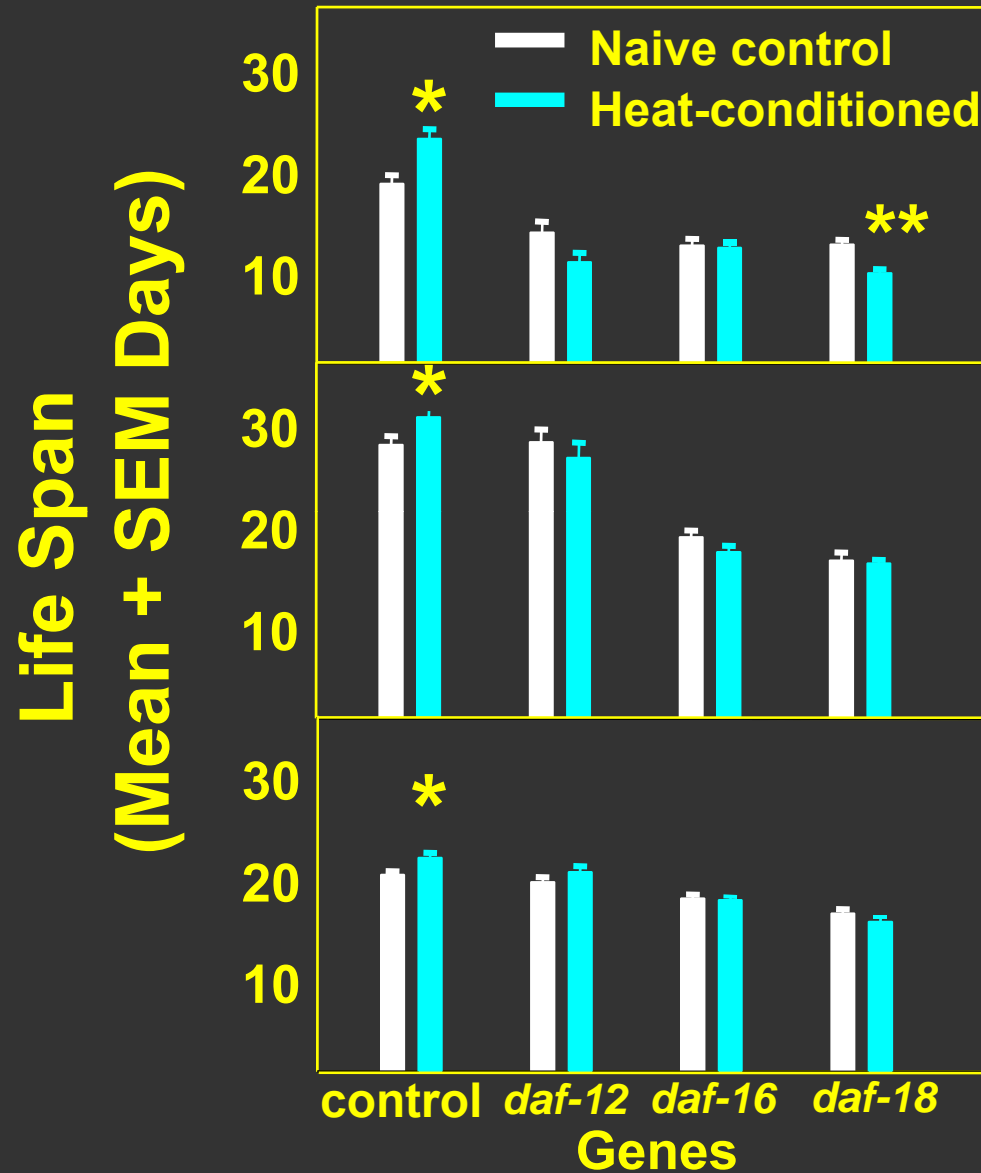
# DAUER FORMATION REQUIRES THE INSULIN-LIKE PATHWAY



THE DAUER / INSULIN PATHWAY ALSO MEDIATES STRESS RESISTANCE AND LIFE EXPECTANCY IN WORMS

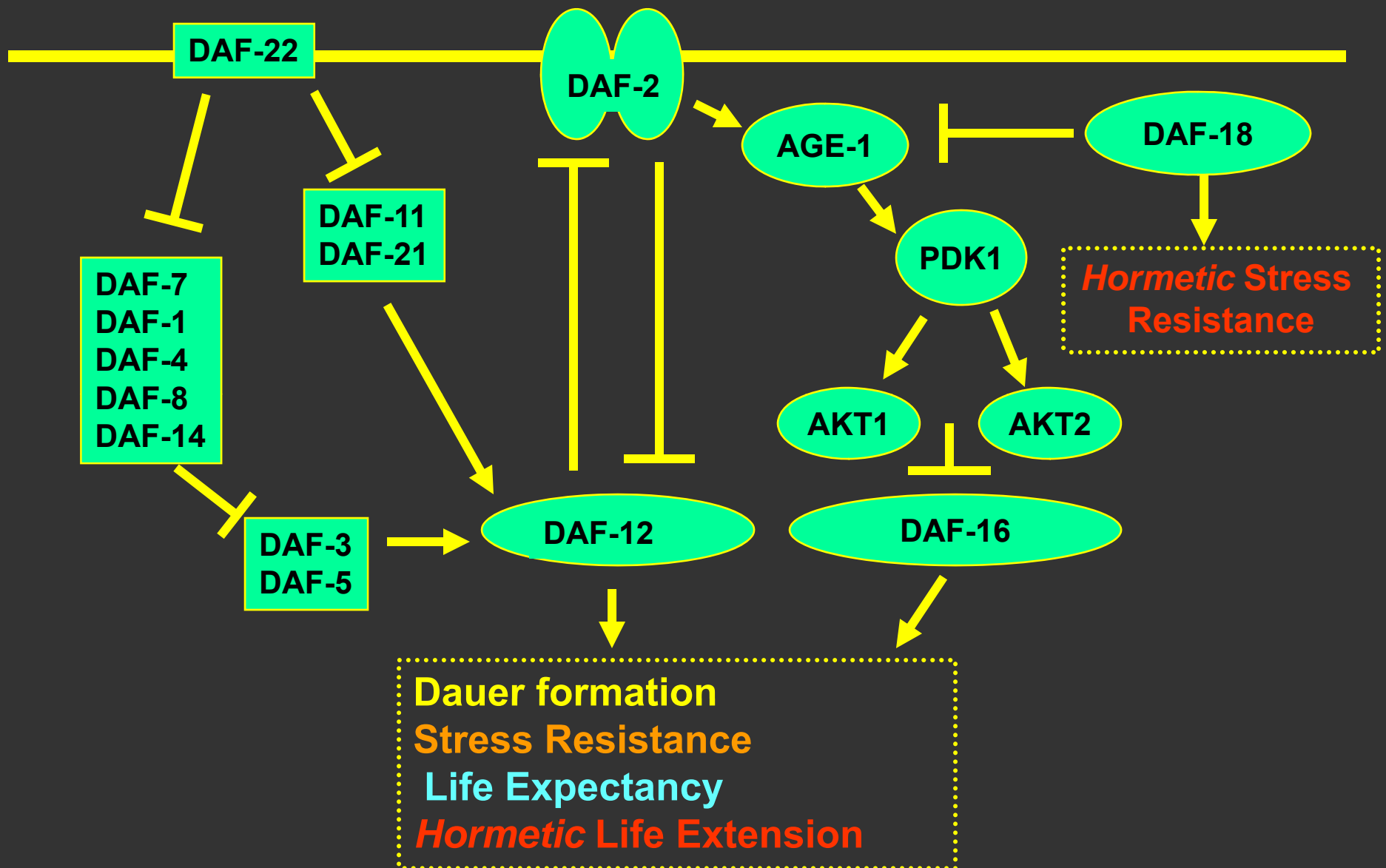


## *daf-12*, *daf-16* & *daf-18* REQUIRED FOR HORMETIC LIFE EXTENSION



**WORMS CARRYING MUTATIONS OF THE INSULIN PATHWAY DISPLAY DEFECTS IN HORMETIC LIFE EXTENSION** Cypser et al, *Biogerontol.* 2003

# DAUER FORMATION / INSULIN-LIKE PATHWAY OF THE WORM



**GENES OF THE INSULIN PATHWAY ARE REQUIRED FOR  
HORMESIS IN THE WORM**

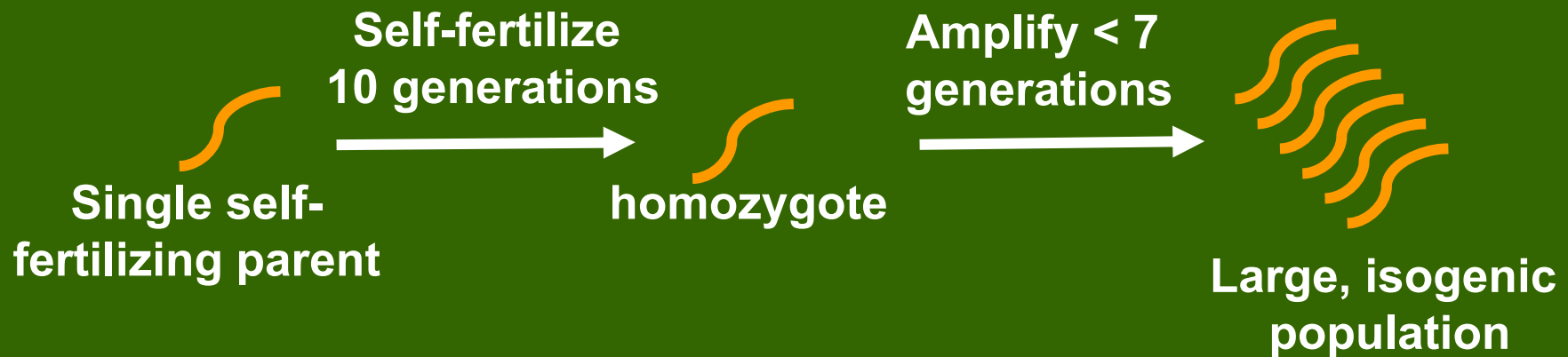
*Cypser et al, Biogerontol, 2003*

# SUMMARY OF WORM GENETICS AND HORMESIS

Genes (*daf-16*, *daf-12*, *daf-18*) of the insulin response pathway are required for extended life span after hormetic conditioning in the worm

The *daf-18* gene is required for a full hormetic response of (thermotolerance) in the worm

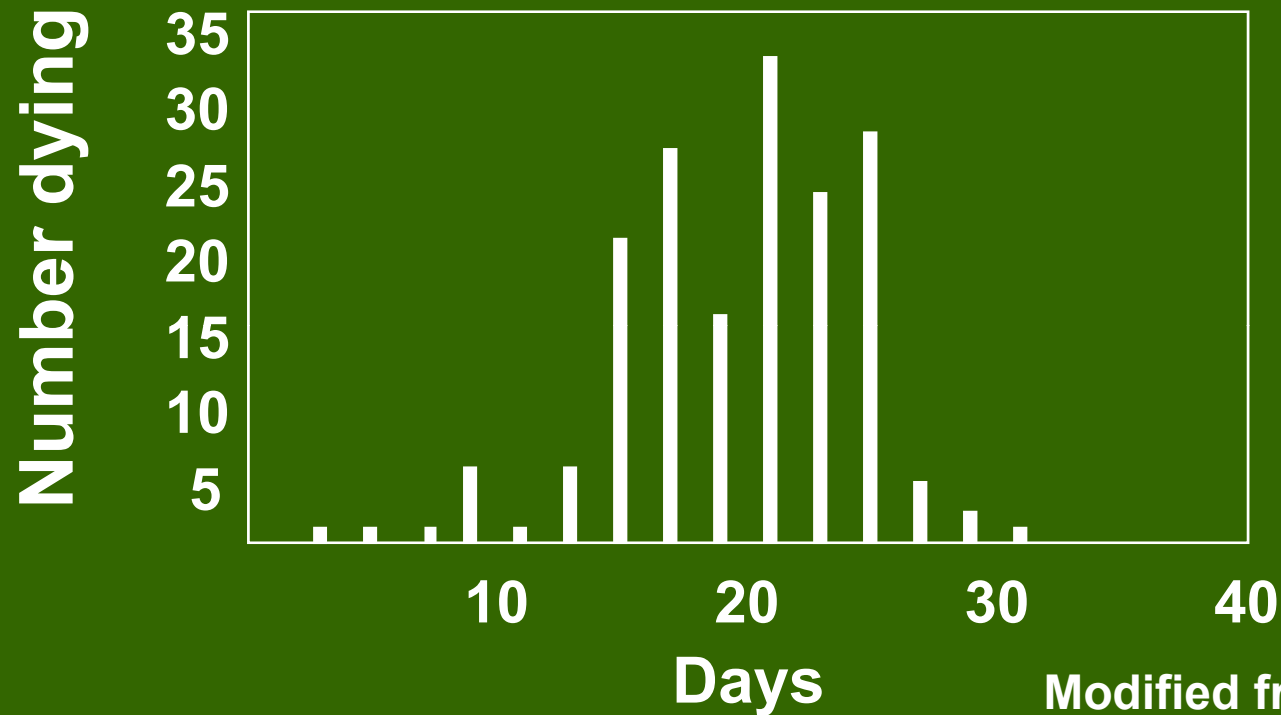
## CAN WE STUDY HORMESIS IN ISOGENIC POPULATIONS OF WORMS?



EASY TO MAKE

LARGE, SAME-AGED, ISOGENIC POPULATIONS OF WORMS

# Ages at death of wild-type, isogenic worms



Modified from Kirkwood,  
*Nature* 2002

**Variation in life span exists  
even in isogenic populations of worms:  
Variation in gene expression should also occur.**

# Heat Shock Protein-16 genes (*hsp-16*)

Specifically, *hsp-16.2* (family of four genes)

- Encodes a 16-kD heat shock protein
- Expression mediated by DAF-16 & Heat Shock Factor 1
- Induced in response to heat shock, hypoxia, other environmental stresses
- Up regulated in long-lived mutants
- Over expression has modest life extension

Can create a transgene

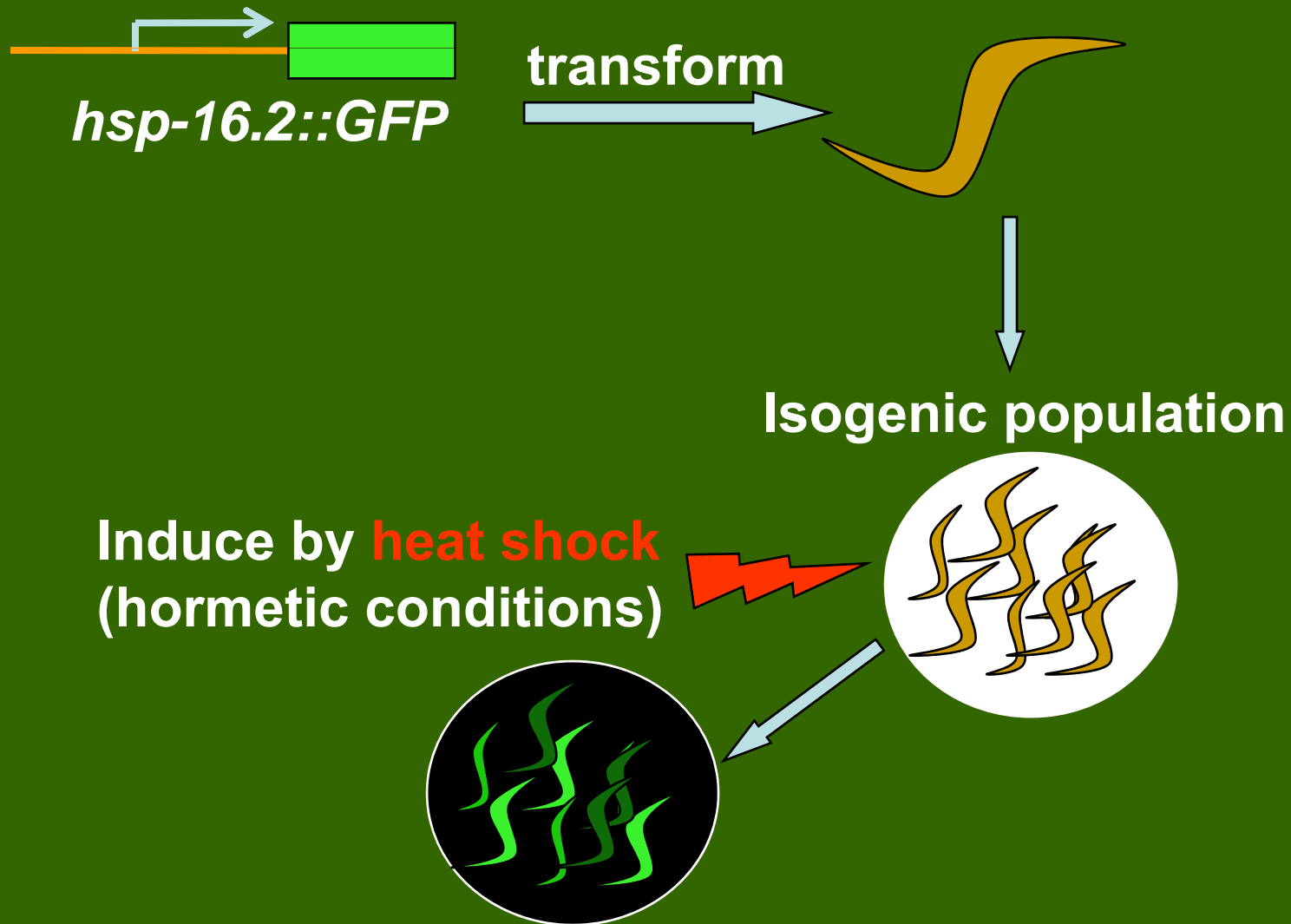
- **Promoter**: worm *hsp-16.2* gene
- **Coding**: jellyfish Green Fluorescent Protein (*GFP*) gene



*hsp-16.2::GFP*



# CREATING A TRANSGENIC WORM STRAIN



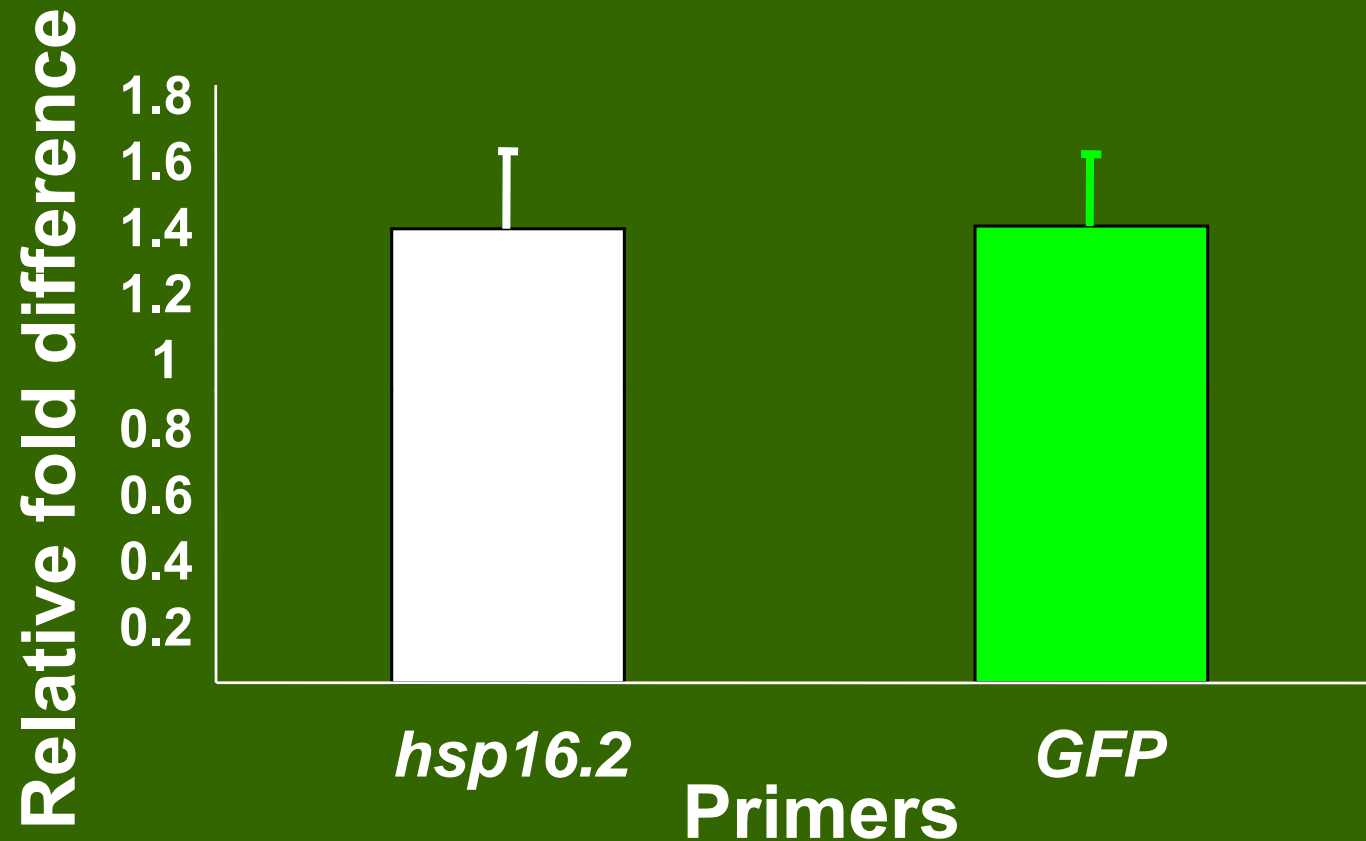
**Green Fluorescent Protein (GFP)** is a “reporter”  
for a single, stress-sensitive gene: *hsp-16.2*

## EXPRESSION OF *hsp-16.2::GFP*



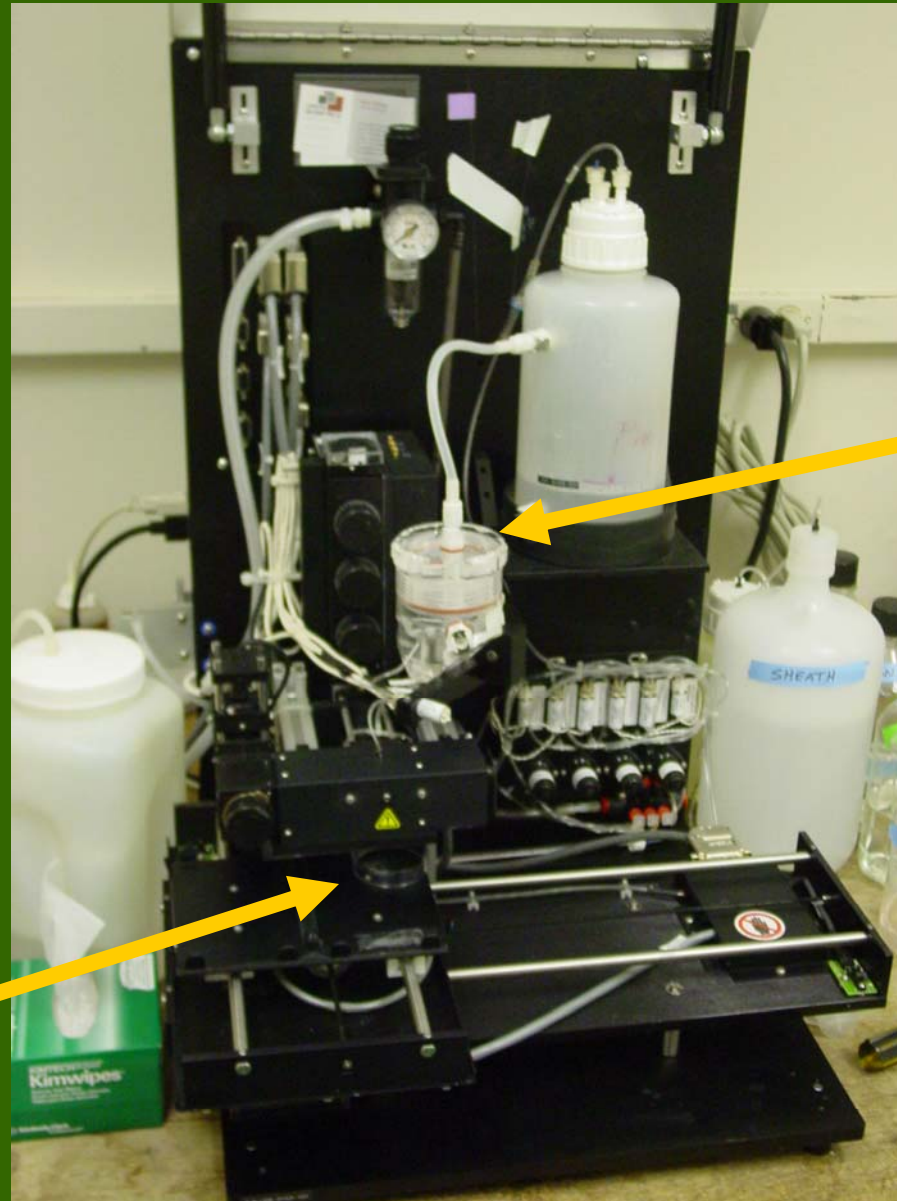
**Worms respond differently to heat shock.  
Due to differences in transgene copy number?**

# QPCR on genomic DNA of *hsp16.2::GFP*-expressing animals



Individual variation NOT due to differences  
in copy number of transgene or endogenous  
*hsp-16.2* gene (P. Tedesco, unpublished)

# GFP brightness is measured using a worm sorter



Sorted by  
GFP expression  
(bright v. dim)  
at  $\sim 10^4$  / hr

Worms go  
in here

Union Biometrica, Holliston, MA

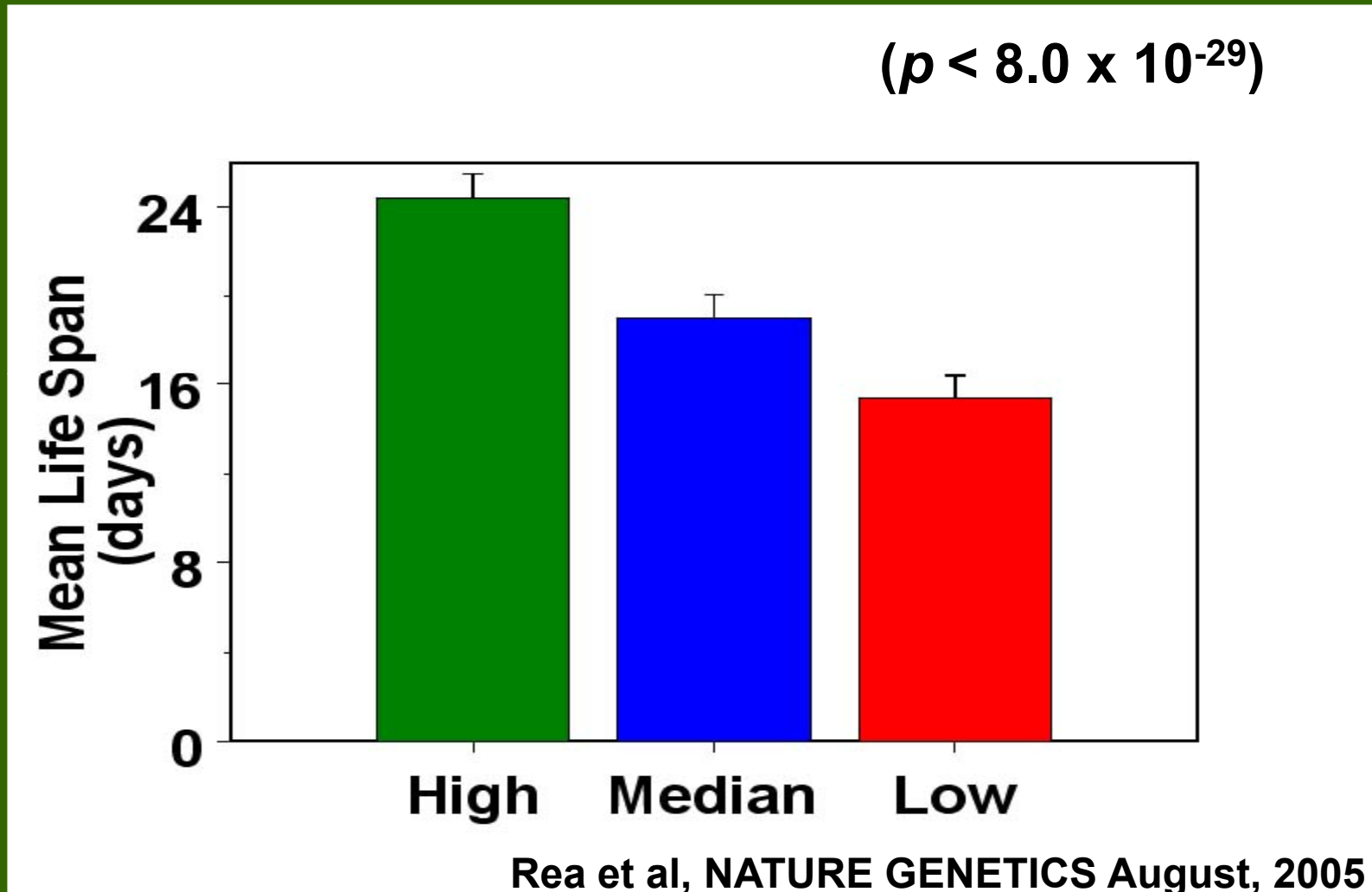
# Questions about individual variation and hormesis

Does *hsp-16.2* expression after hormetic conditioning correlate with:

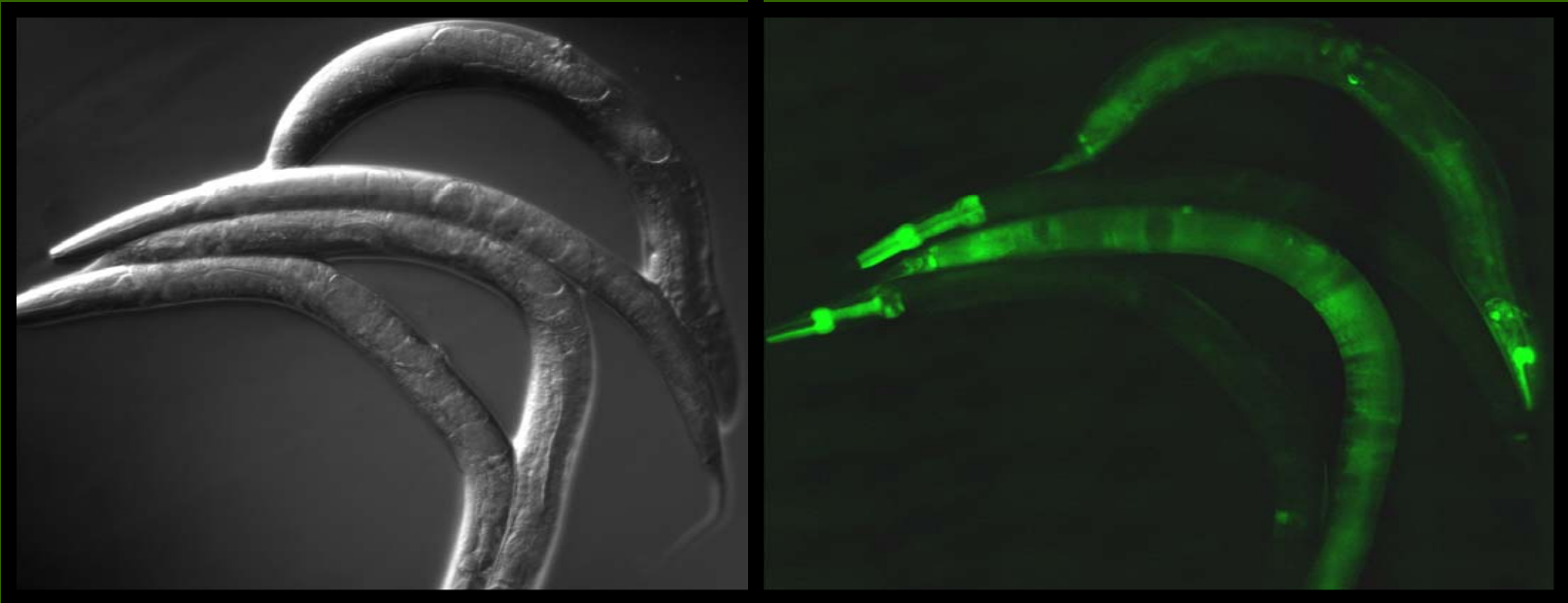
- Differential thermotolerance?
- With differential lifespan?



# COMBINED LONGEVITY EXPERIMENTS



Expression of *hsp-16.2::GFP* predicts longevity of genetically identical individuals



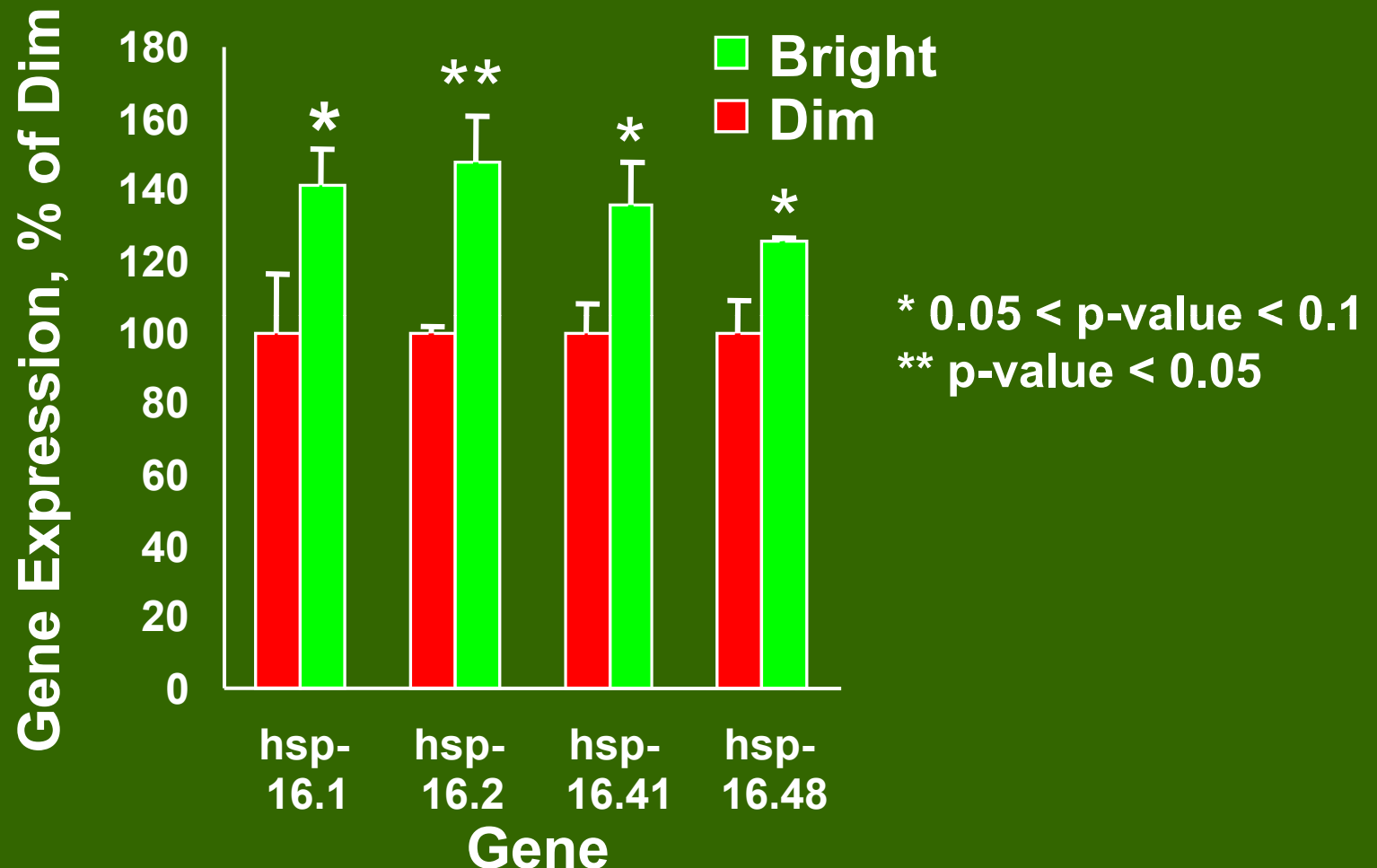
## CONCLUSIONS:

- Isogenic worms display variance in life span and thermotolerance
- Response to hormetic conditioning predicts life span
- Not due to genetic differences
- “Physiologic state” on first day of adulthood predicts life expectancy after hormetic conditioning

# QUESTIONS ABOUT PHYSIOLOGIC STATE AFTER HORMETIC CONDITIONING

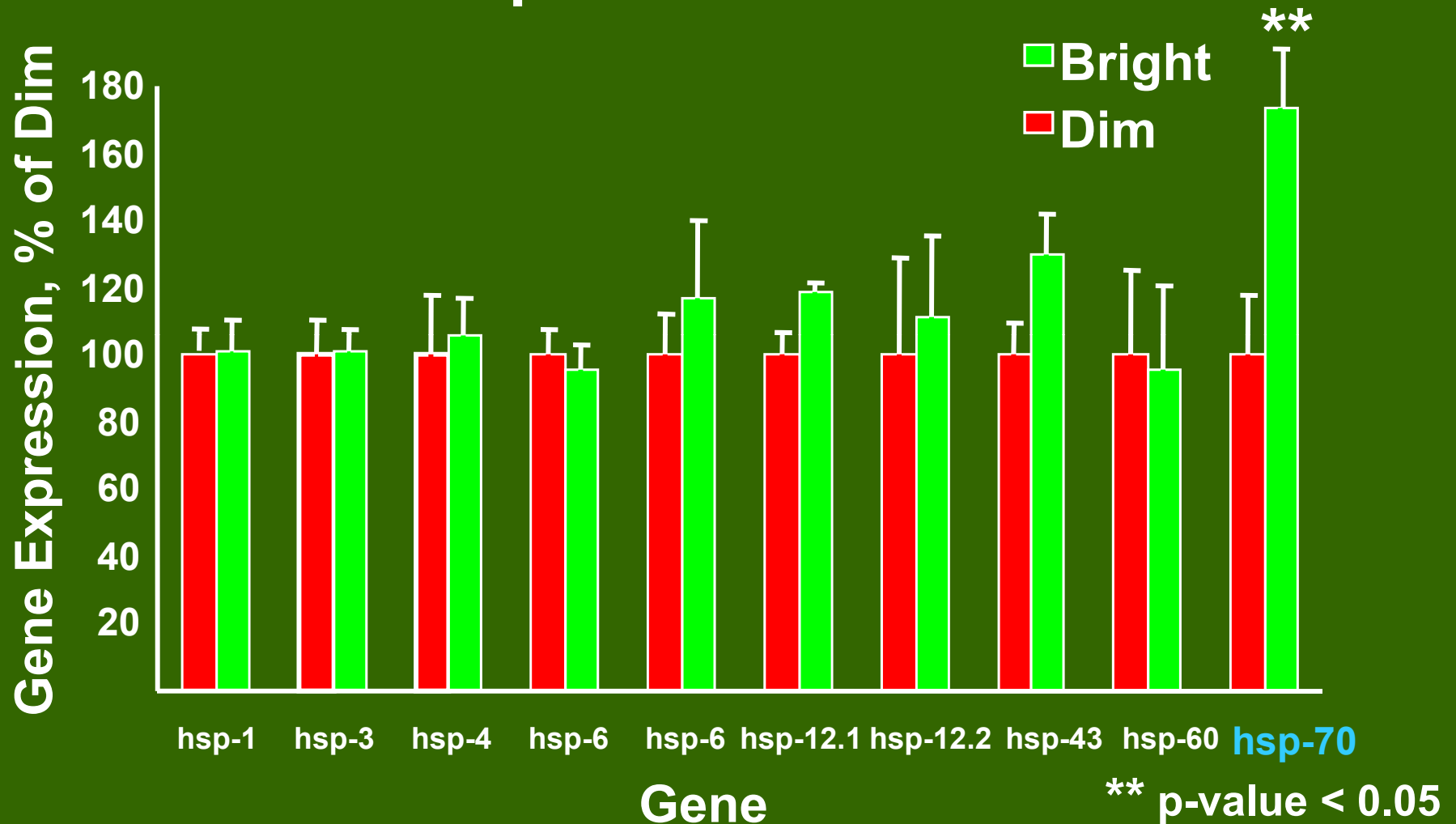
- Are genes besides *hsp-16.2* differentially expressed?
- Is physiologic state heritable?
- Are there epigenetic influences?

# MICRO-ARRAY ANALYSIS: INDUCTION OF *HSP-16* CLASS IN BRIGHT WORMS



**MULTIPLE GENES OF THE HEAT-SHOCK PROTEIN-16 CLASS  
ARE UPREGULATED IN BRIGHT WORMS (S.-K. Park, unpublished)**

# Micro array analysis: Other heat shock proteins in conditioned worms

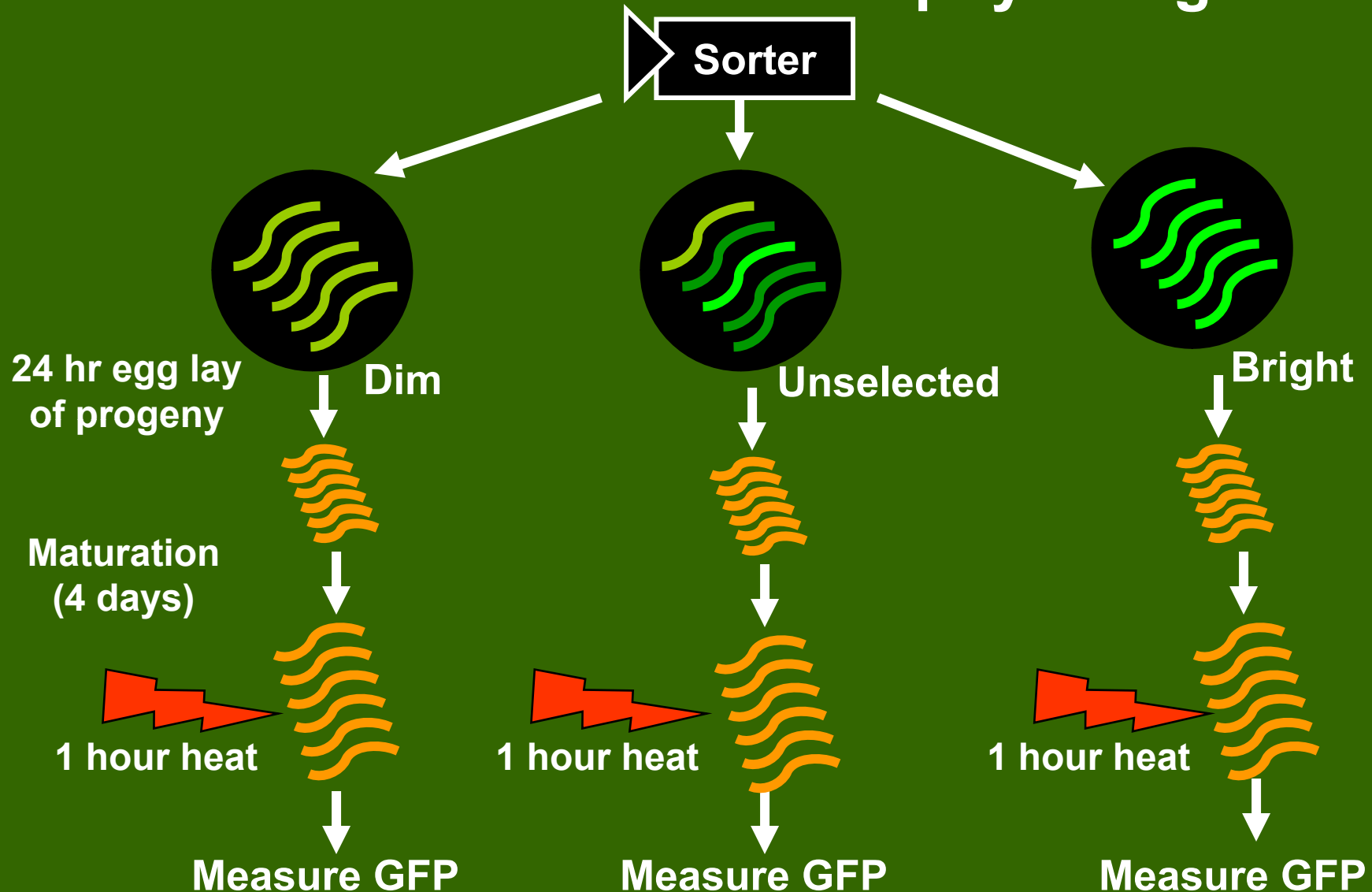


***hsp-70* is up-regulated in bright, longer-lived worms**

(S.-K. Park, unpublished)

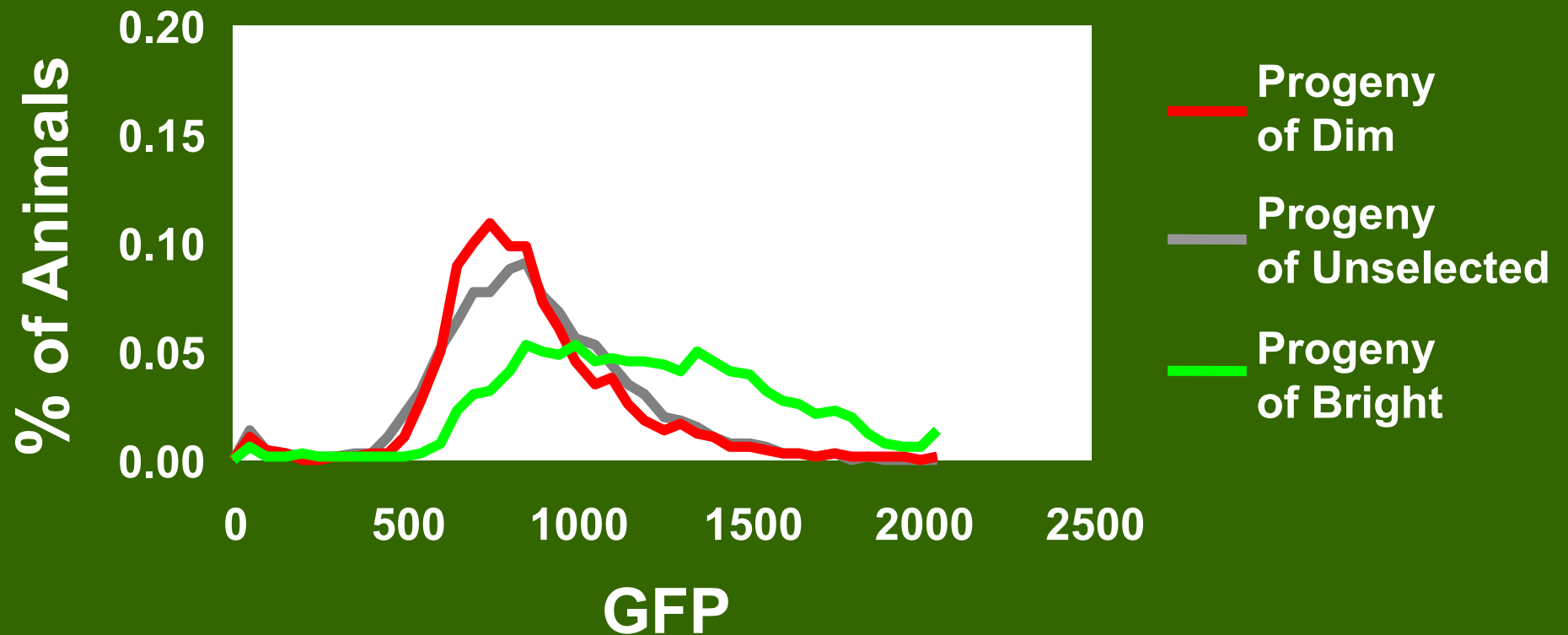


# How to test for inheritance of physiologic state



**Expectation: All groups of progeny  
*should* display similar GFP expression**

## *hsp-16.2::GFP* expression of progeny



n > 3000 / group  
Dim vs. Bright:  
 $p < 10^{-17}$  (chi-square test)

**PROGENY RESPONSE TO CONDITIONING**  
**SIMILAR TO PARENTS EVEN IN ISOGENIC POPULATIONS**  
(J.R. Cypser, unpublished)

# OVERALL SUMMARY

Insulin pathway genes required for hormetic heat conditioning in the worm

- For life expectancy: *daf-12*, *daf-16*, *daf-18*
- For thermotolerance: *daf-18*

Individual variation in individual gene expression after hormetic heat conditioning predicts subsequent thermotolerance and life expectancy.

- For life expectancy OR thermotolerance: *hsp-16.2* predictive

Physiologic state of worms after heat conditioning may be heritable even in an isogenic population

- hsp-16.2::GFP* expression after heat conditioning of progeny reflects expression displayed by parents.
- Epigenetic effects may act in response to heat conditioning; *sir-2.1* required for sustained conditioning

## Future Directions

Test inheritance of *hsp-16.2::GFP* response further  
Variation in promoter region of transgene?  
Inheritance of thermotolerance / life span?  
Difference observable at protein level?

Are there epigenetic influences on hormesis?  
Test histone-modifying genes  
Pharmacologic interventions

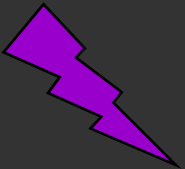
Does expression of *hsp-16.2::GFP* in specific tissues predict hormetic survival?

**Thanks to:**

**Deqing Wu  
Pat Tedesco  
Sang-Kyu Park  
Shane Rea**

**Tom Johnson**

**International Dose-Response Society**





# Heat Shock Protein-16 gene (*hsp-16*)

- *hsp-16.2* encodes a 16-kD heat shock protein (HSP)
- Member of the *hsp16/hsp20/alphaB-crystallin* (HSP16) family of heat shock proteins; 6 near identical homologs
- *hsp-16.2* expression, strongest in intestine and pharynx
- Under Daf-16 & HSF-1
- Induced in response to heat shock, hypoxia, or other environmental stresses
- Mount 36; other HSPs
- Up regulated in Age mutants
- Over expression has modest life extension
- Interacts with intracellular human beta amyloid peptide, a primary component of plaques in Alzheimer's disease
- HSP-16.2 is likely to function as a passive ligand temporarily preventing unfolded proteins from aggregating

# Summary:

## Physiologic state after conditioning

Physiologic state associated with induction of:

- multiple heat shock protein-16 class genes
- heat shock protein-70

Degree of expression of GFP driven by *hsp-16.2* is **heritable even in populations of isogenic animals.**

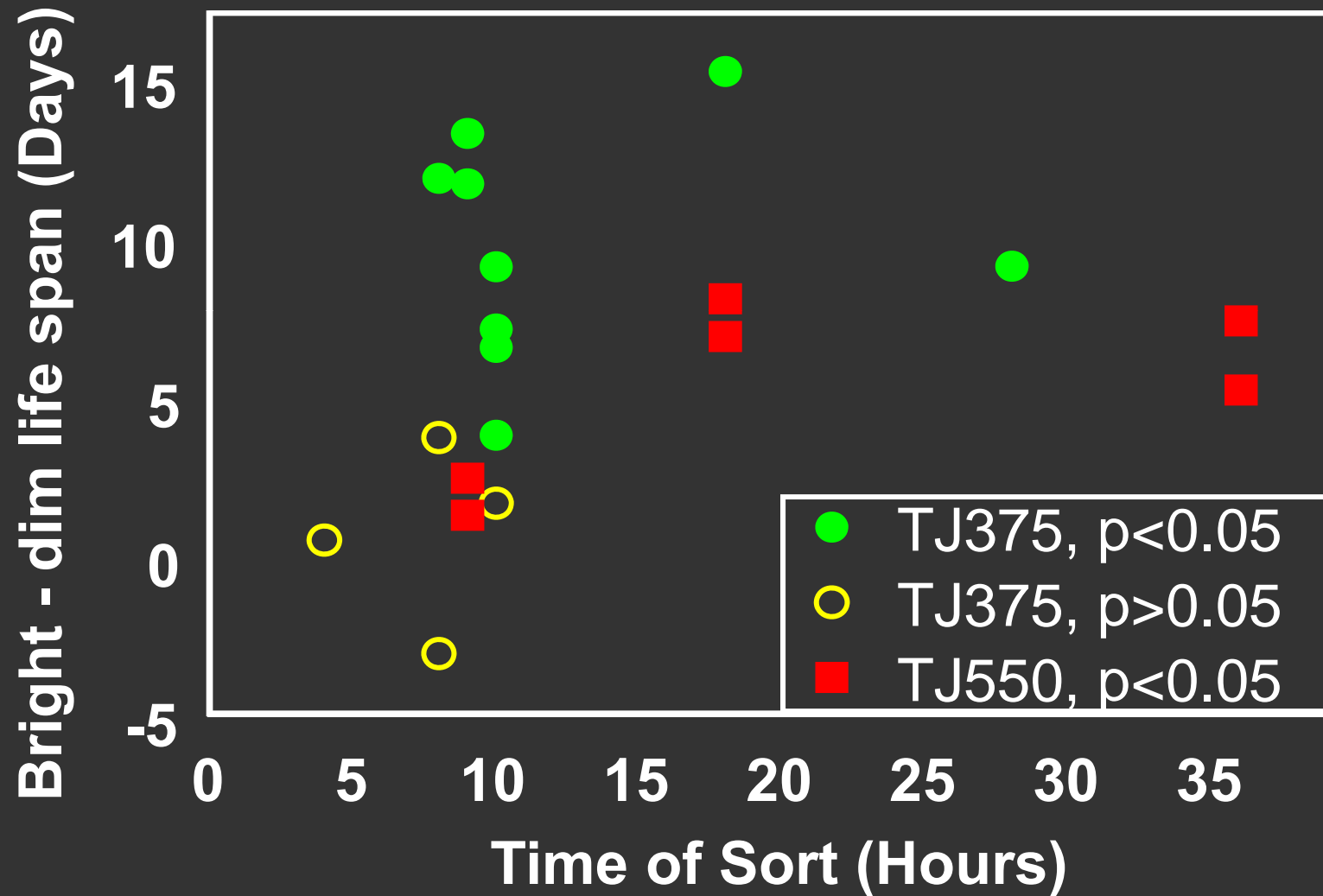
Models:

Duplications / deletions of the *hsp-16.2::GFP* transgene occur rapidly, creating subpopulations with different copy numbers: **NO**

Differences in promoter sequence? **NOT TESTED**

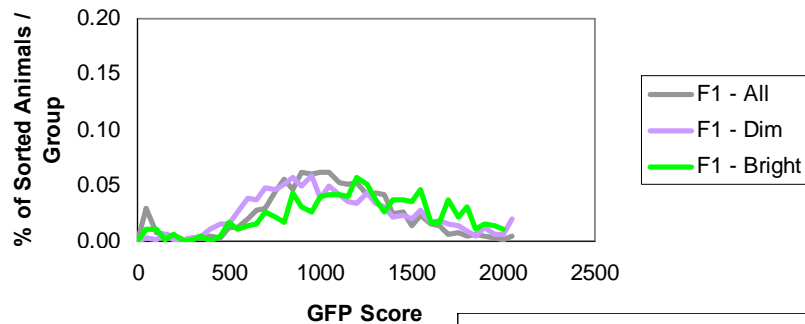
Differences in epigenetic state? **MAYBE; *sir-2.1* role in heat conditioning**

# Differential Longevity: All Individual Experiments

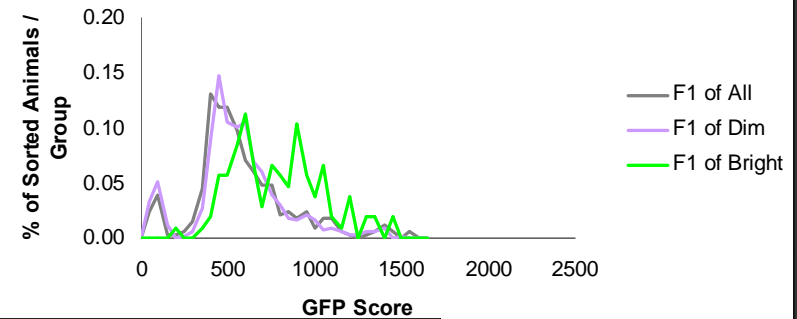


# Expression of *hsp-16.2::GFP* in progeny of dim vs. bright worms

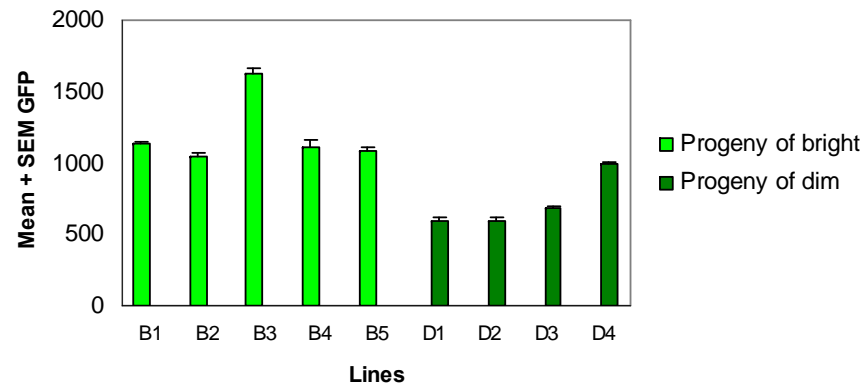
Experiment S8 Summary  
Acq, Sort, Rej, Coin



Experiment S12 Summary  
(Sorted, Rejected, Coincident)

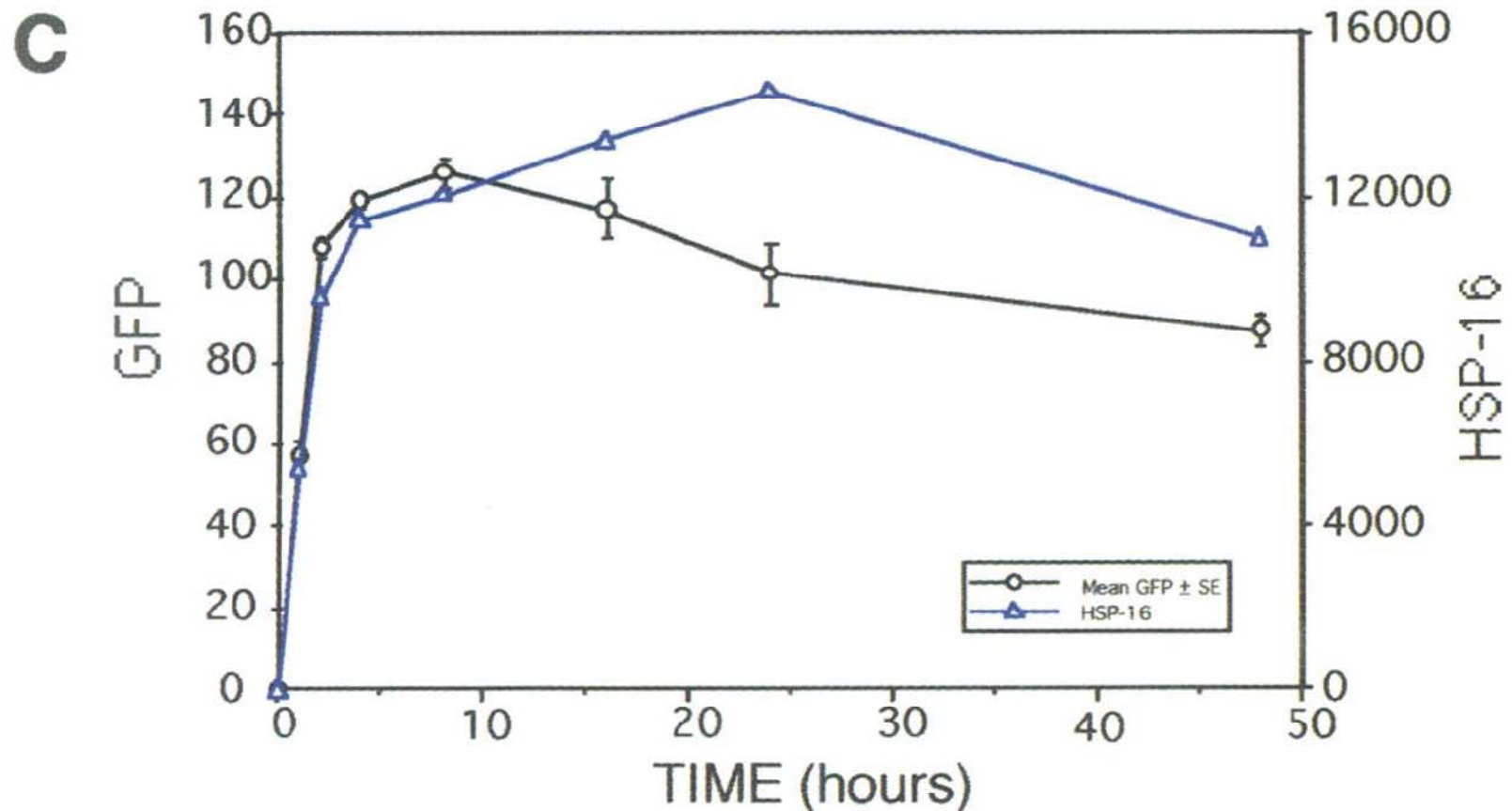


Inheritance of *hsp-16.2::GFP* expression in isogenic population



**Progeny reliably reflect expression of GFP even in isogenic populations** J. R. Cypser, unpublished

## Endogenous HSP-16.2 expression tracks HSP-16.2::GFP expression



**hsp-16.2::GFP accurately reports expression of the native HSP-16.2 protein**

Link et al, *Cell Stress & Chaperones*, 1999

**Direct Observation of Stress Response in *Caenorhabditis elegans* Using a Reporter Transgene.** Christopher D. Link, James R. Cypser, Carolyn J. Johnson and Thomas E. Johnson  
*Cell Stress & Chaperones*, Vol. 4, No. 4 (Dec., 1999), pp. 235-242

*Michalski et al, Biogerontol. 2001*

*Cypser et al, J. Gerontol. 2002*

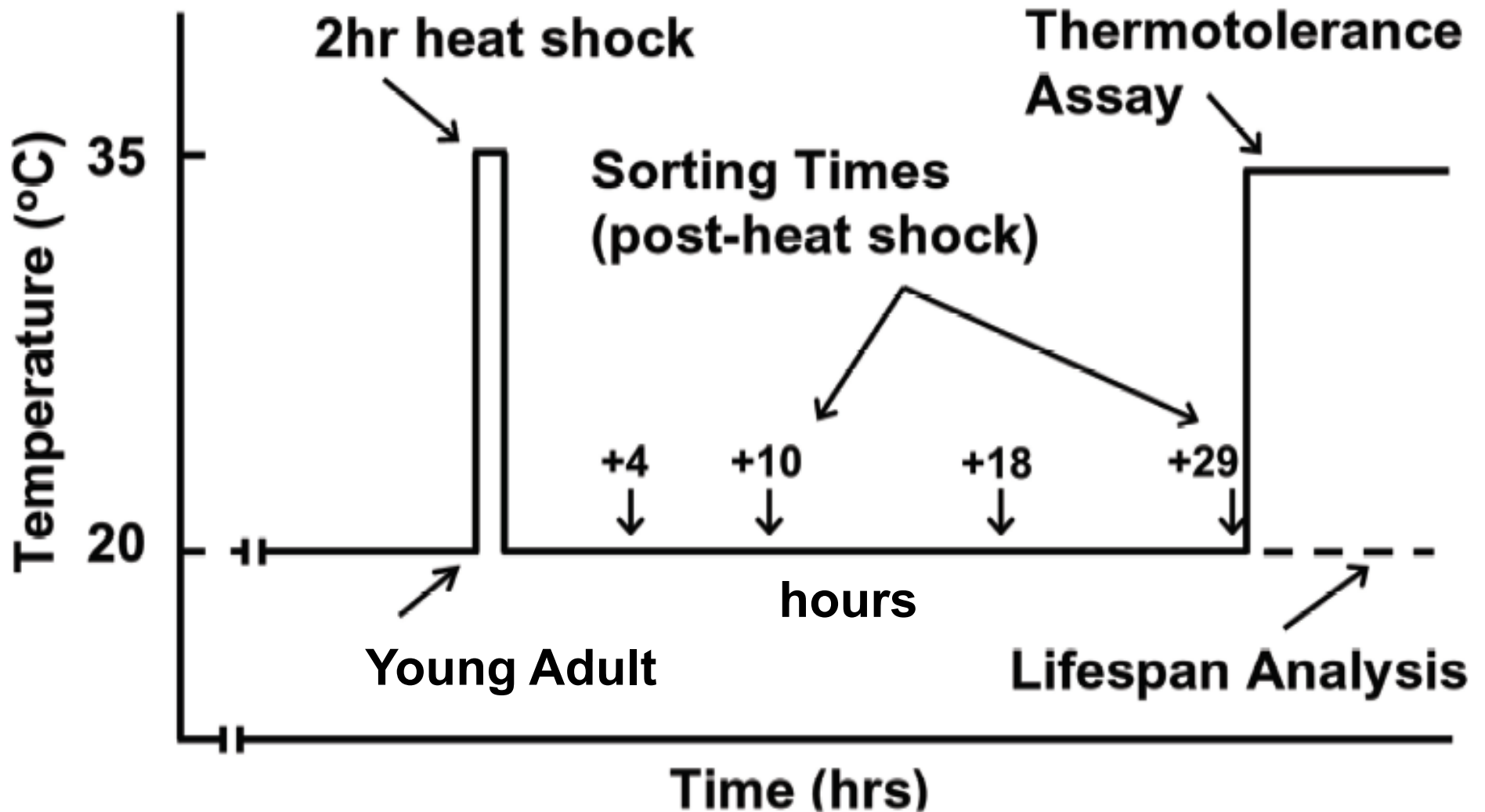
*Cypser et al, Biogerontol, 2003*

Modified from Kirkwood,  
*Nature 2002*

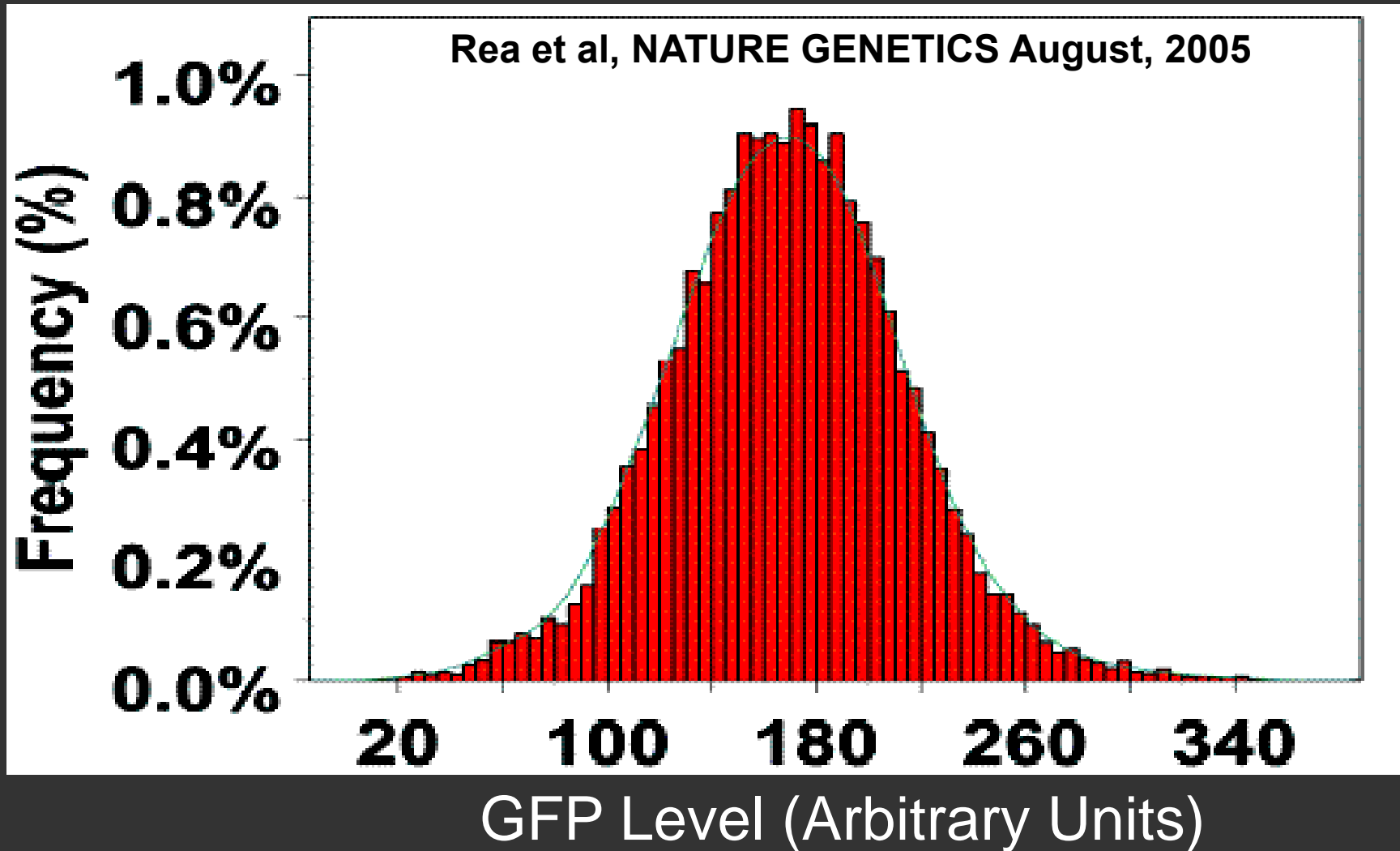
Rea et al, NATURE GENETICS August, 2005

# Induction Methods

Rea et al, NATURE GENETICS August, 2005



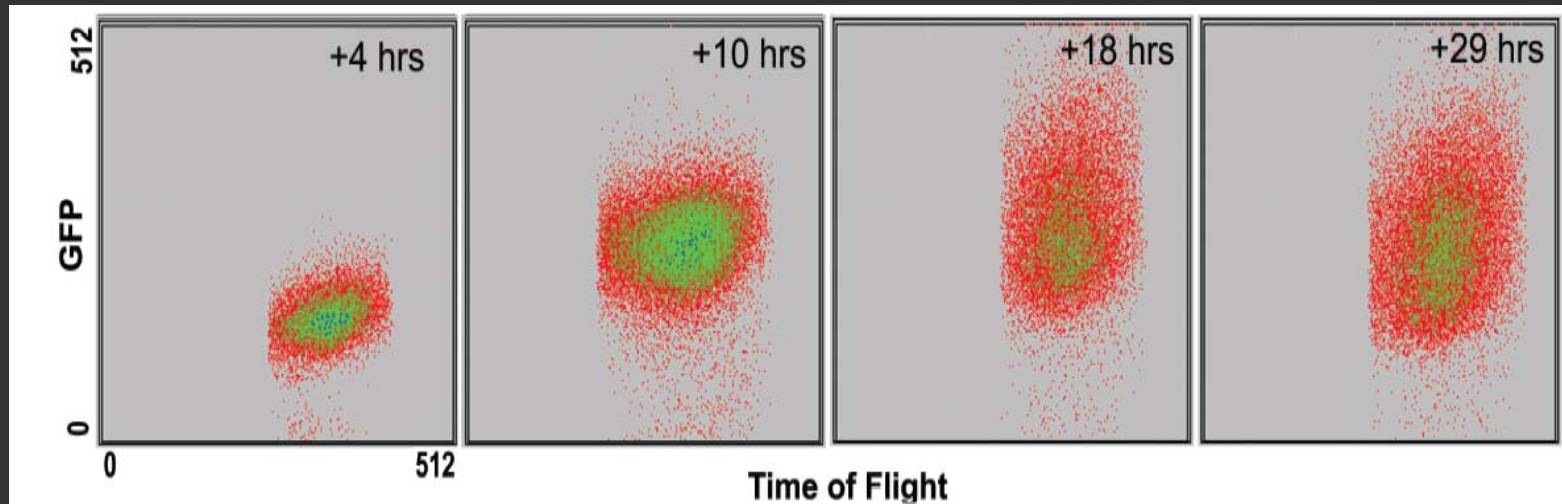
# GFP Is Normally Distributed



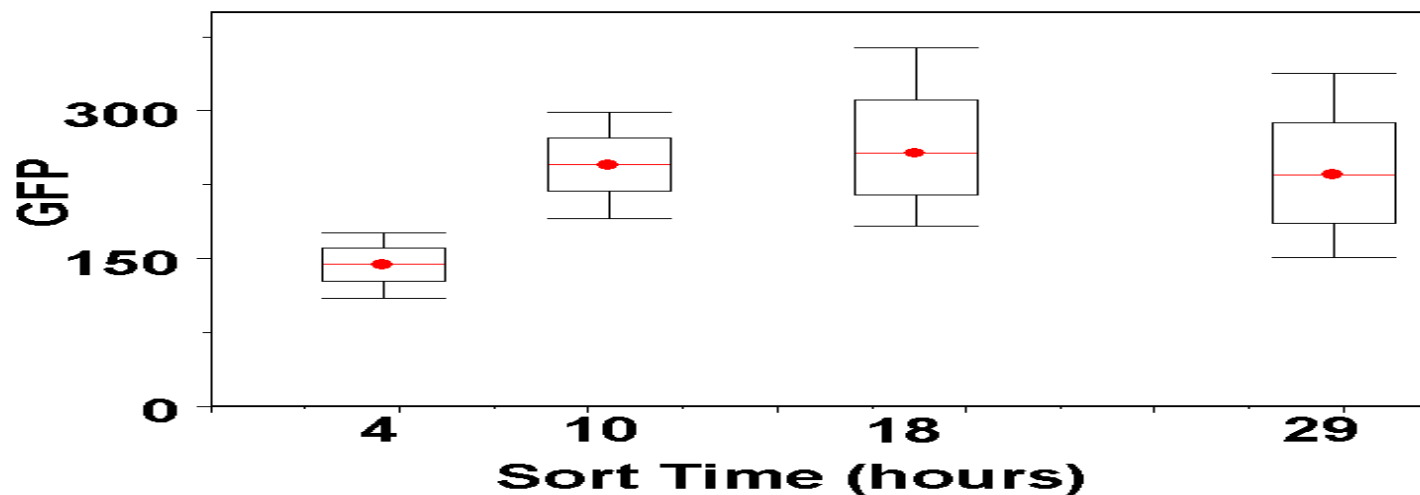
2 hour conditioning, 19 hour sort, 60,000 worms



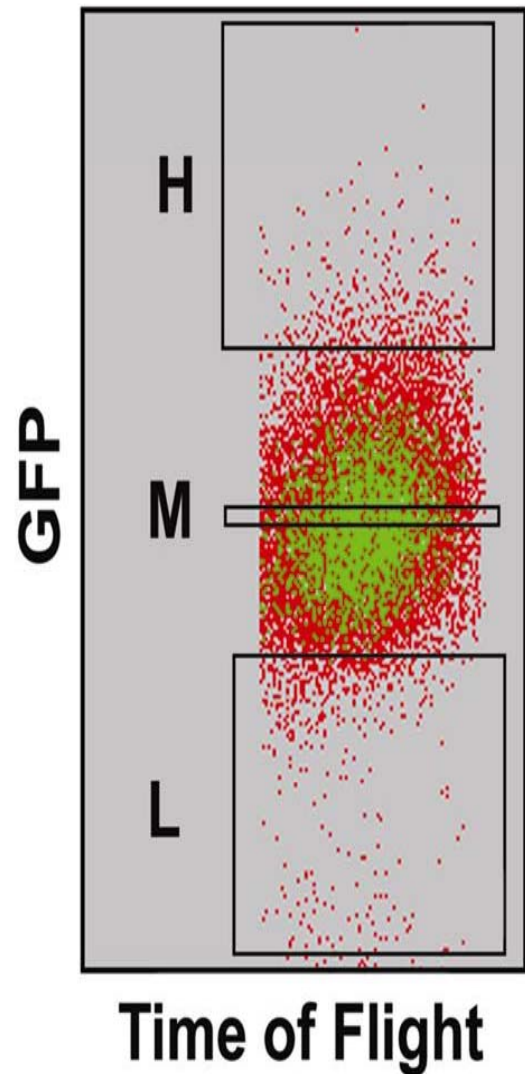
# Variance Increases with Time after Sort



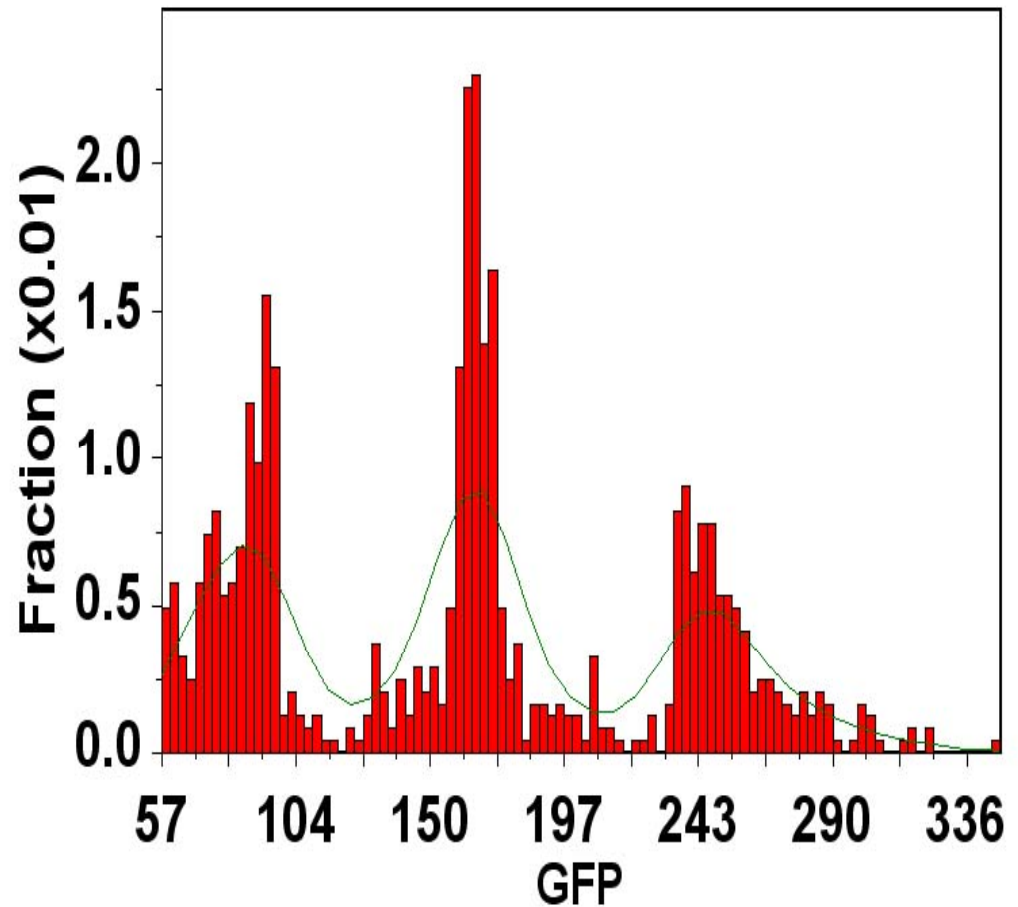
Rea et al, NATURE GENETICS August, 2005



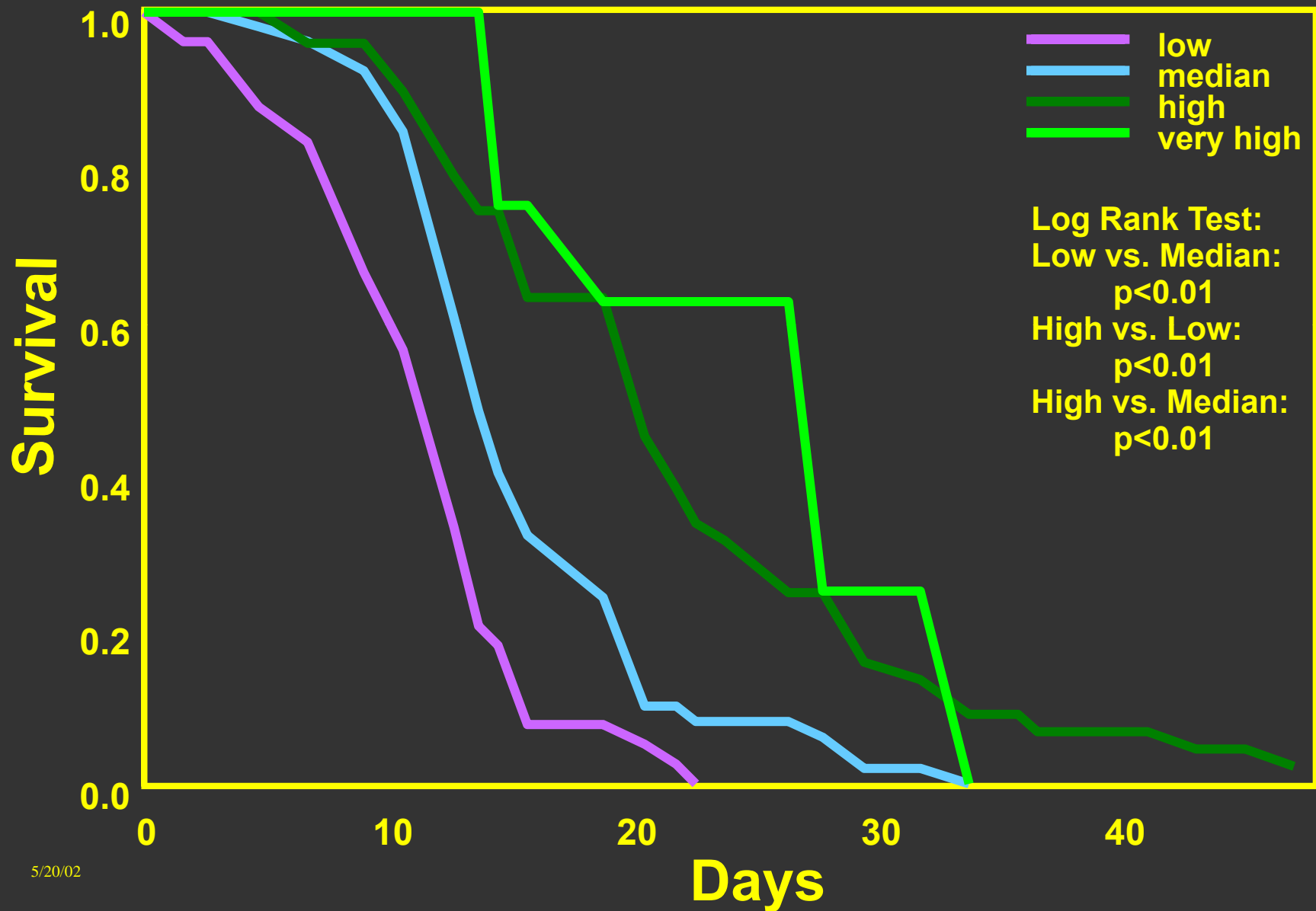
# Sorting for Differential Brightness



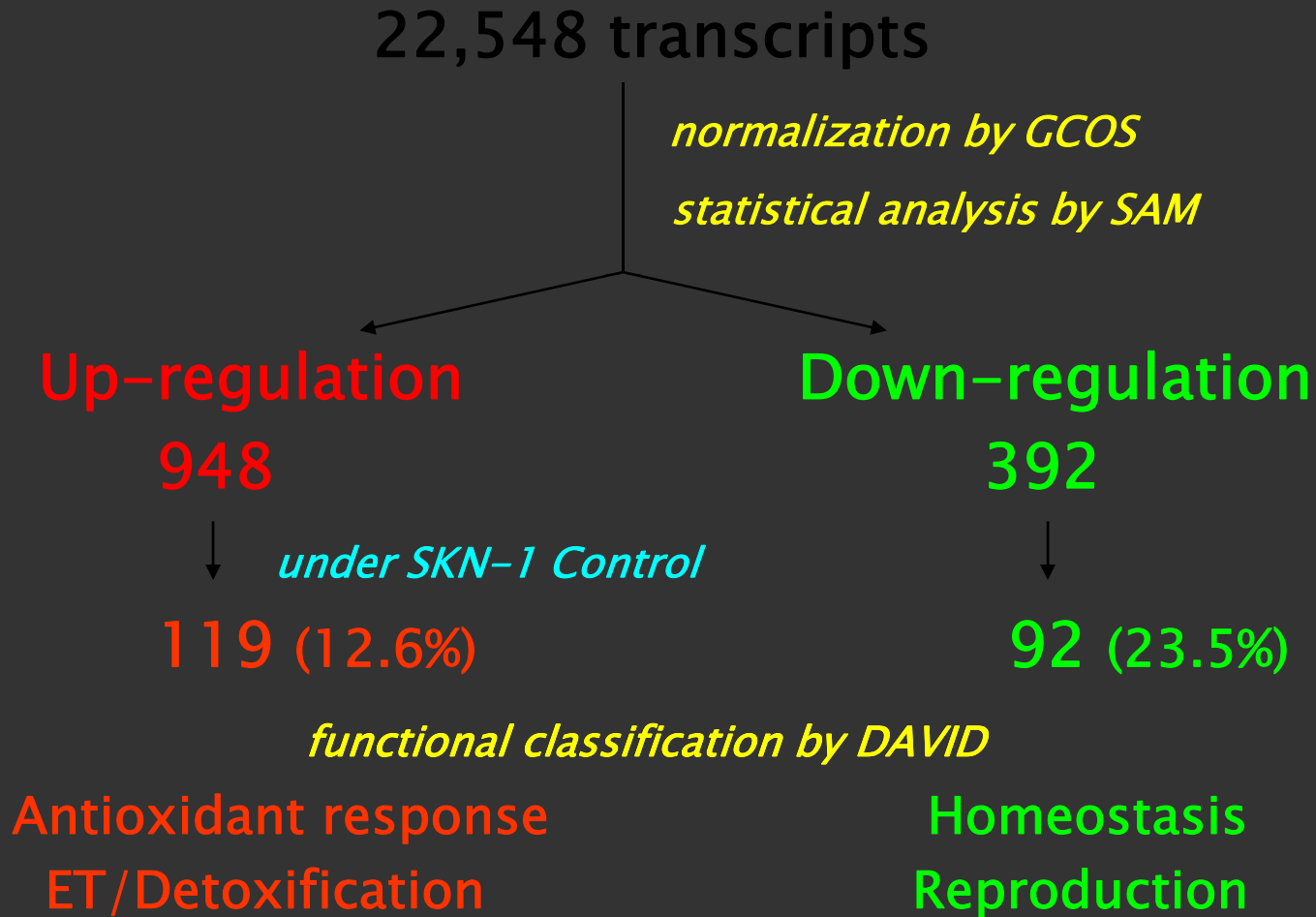
Rea et al, NATURE GENETICS August, 2005



# Life Expectancy



# Microarray Data: control vs. hyperbaric O<sub>2</sub> (effect of reduced skn-1)



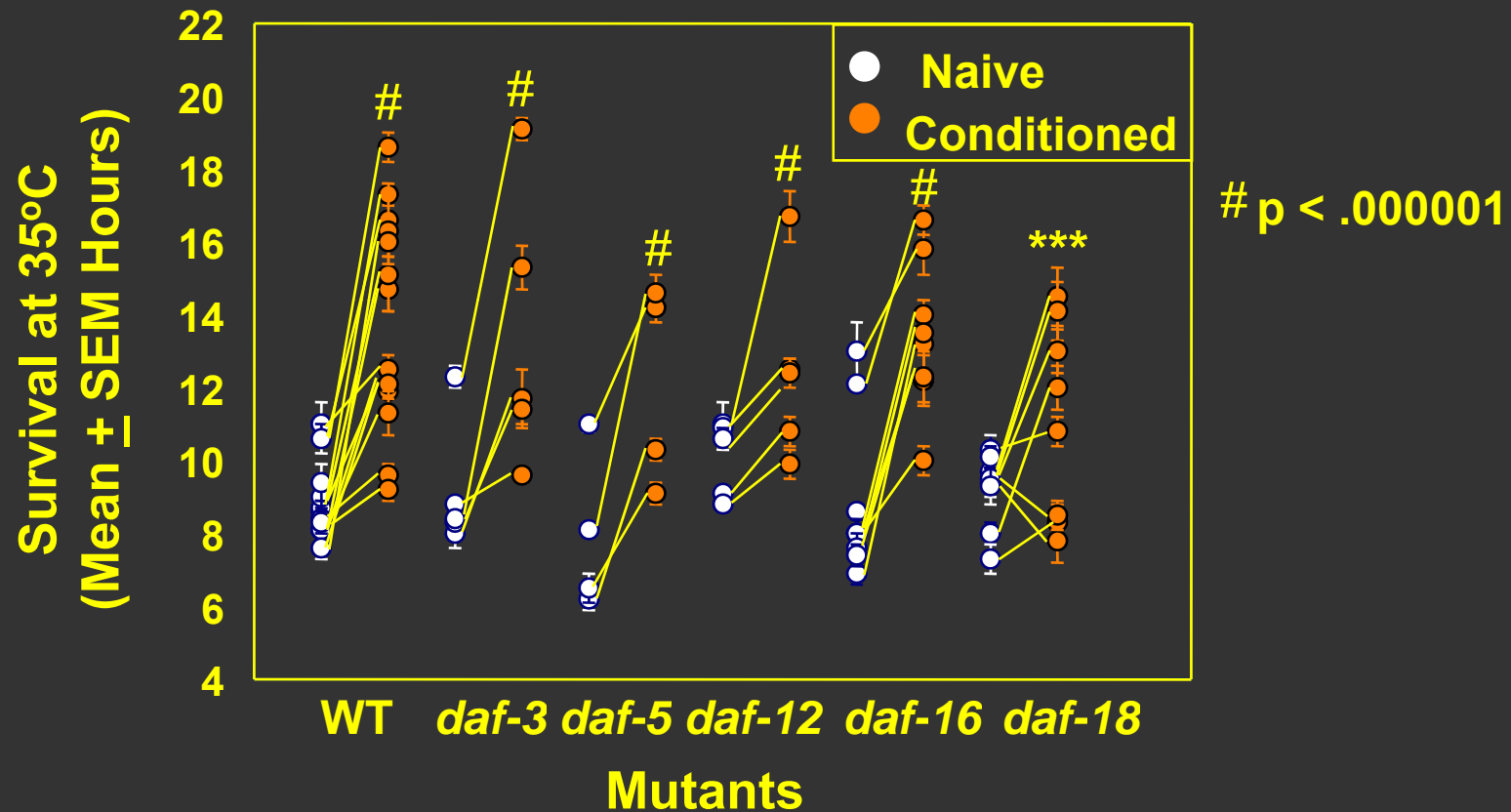
Analysis by S.-K. Park



**Juan Ponce de León** (ca. 1471 – July 1521) was a Spanish conquistador. He was born in Santervás de Campos (Valladolid). Ponce de León accompanied Christopher Columbus on the latter's second voyage to the New World. He became the first Governor of Puerto Rico by appointment of the Spanish Crown. He is also notable for his voyage to Florida, the first known European excursion there, as well as for being associated with the legend of the Fountain of Youth, which was said to be in Florida.

- Wikipedia

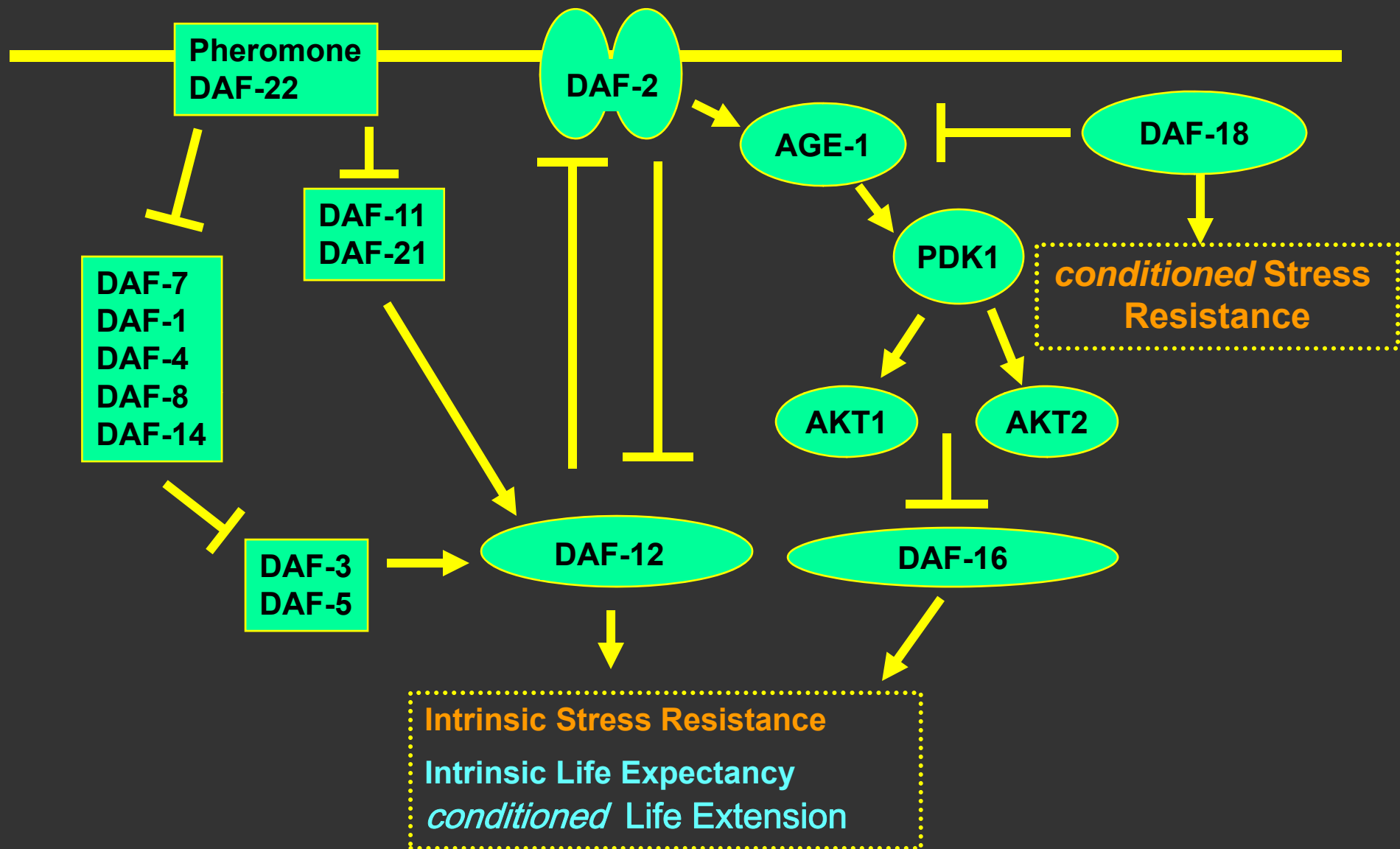
# GENETIC DISSECTION OF HEAT HORMESIS



**MUTANT WORMS THAT CANNOT FORM DAUERS  
STILL DISPLAY HORMETIC THERMOTOLERANCE**

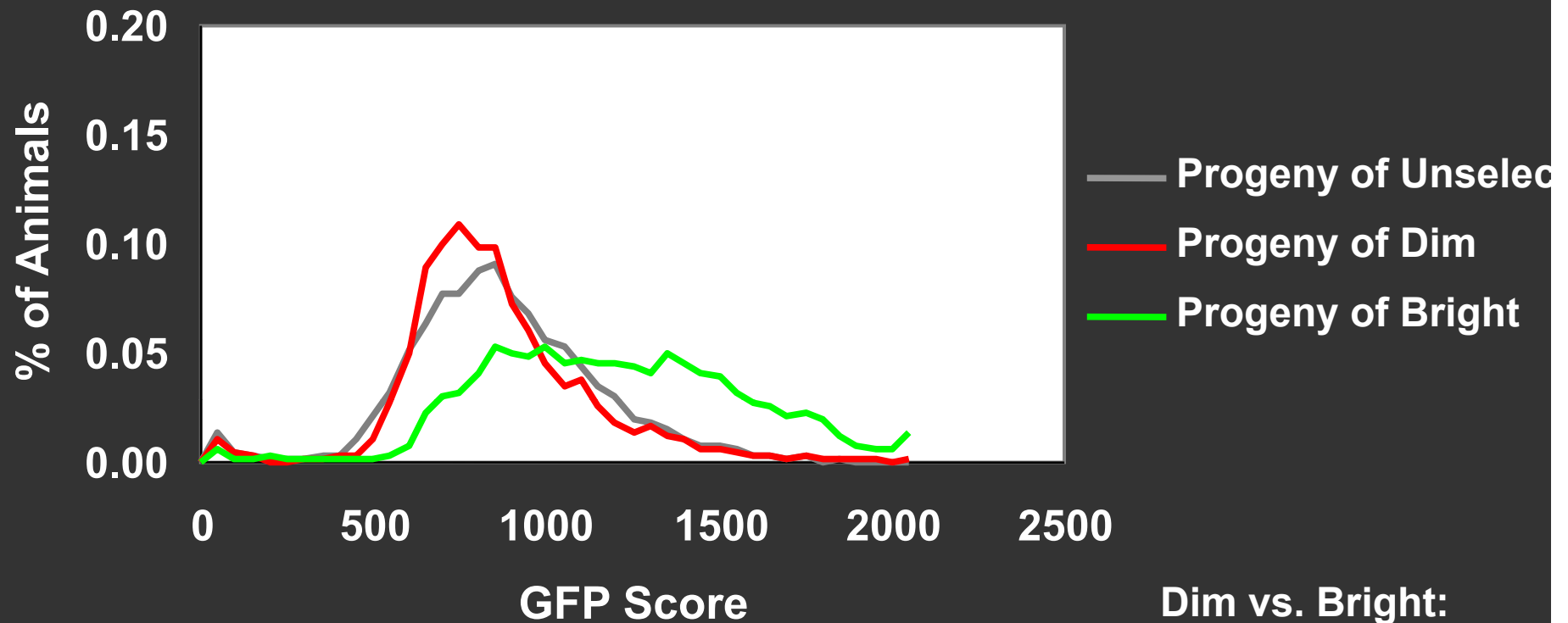
*Cypser et al, Biogerontol. 2003*

## Dauer formation requires the insulin-like response pathway



Elements of this pathway are required for hormesis in the worm

# hsp-16::GFP expression of progeny



Dim vs. Bright:  
 $p \ll 10^{-17}$  ( $\chi^2$  test)

	Mean $\pm$ SD	$\sim n$
F1 of All	847 $\pm$ 273	6300
F1 of Dim	835 $\pm$ 271	3300
F1 of Bright	1167 $\pm$ 371	3100

Progeny tend to display same response to preconditioning as parents in isogenic populations (replicated five times; J.R. Cypser, unpublished)



Differentiate daf-16 from hsp-16  
Use notes features at bottom  
Fold diff in life span (red-blue-green bars) twin analogy  
As usual more ideas than funding  
Check no slides are hidden  
Emphasize HEAT?  
Printouts of slides?  
Other replicates of heritability  
Italics  
Reporter tracking endogenous HSP16 slide  
Slide with references