



Marine invertebrate environmental memory

Implications, mechanisms, and opportunities



Steven Roberts - University of Washington
github.com/sr320/talk-duke-2024

Duke

NICHOLAS SCHOOL *of*
the ENVIRONMENT
Duke University Marine Lab













Early-life Priming

Hardening



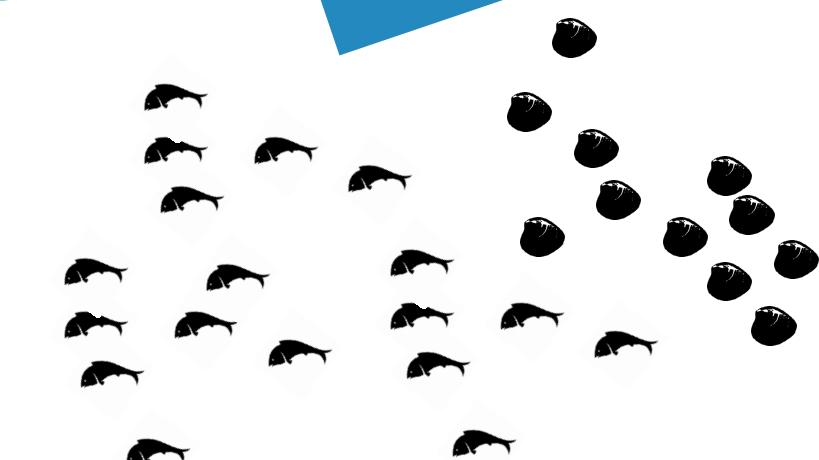
Adults

Influencing adult phenotype by altering early life environment



Transgenerational Plasticity

Carry-over effects



Larvae

Influencing offspring phenotype by altering environmental conditions of parents

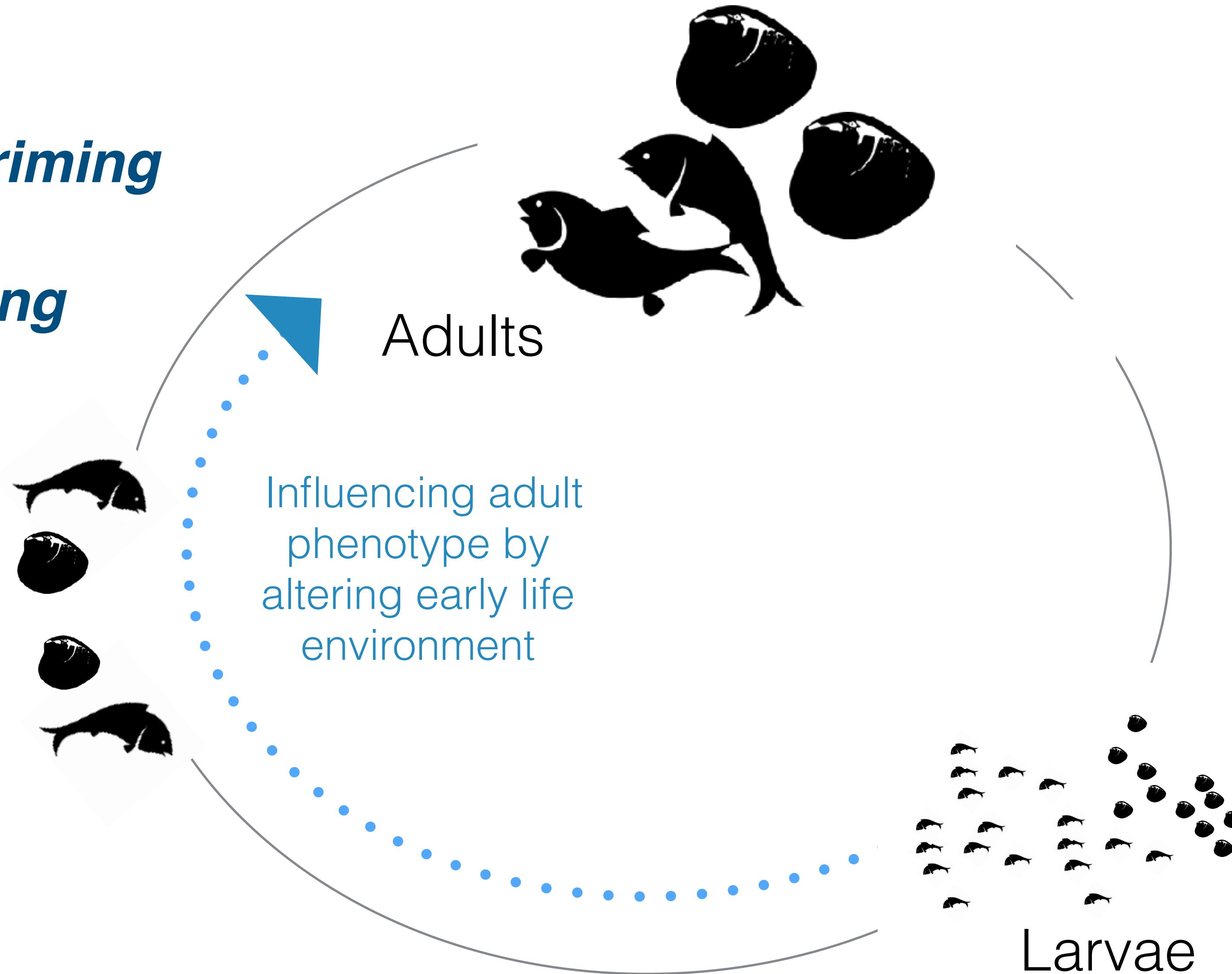
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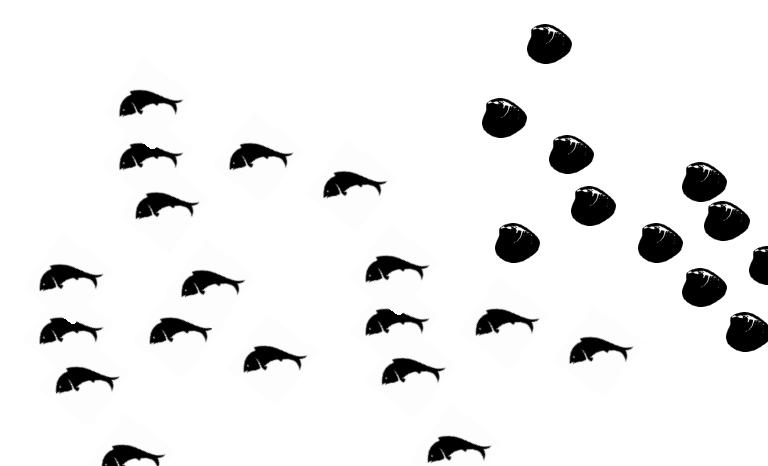
Trends in Plant Science

Early-life Priming

Hardening



Haipei Liu ,¹ Amanda J. Able ,¹ and Jason A. Able  ^{1,@,*}

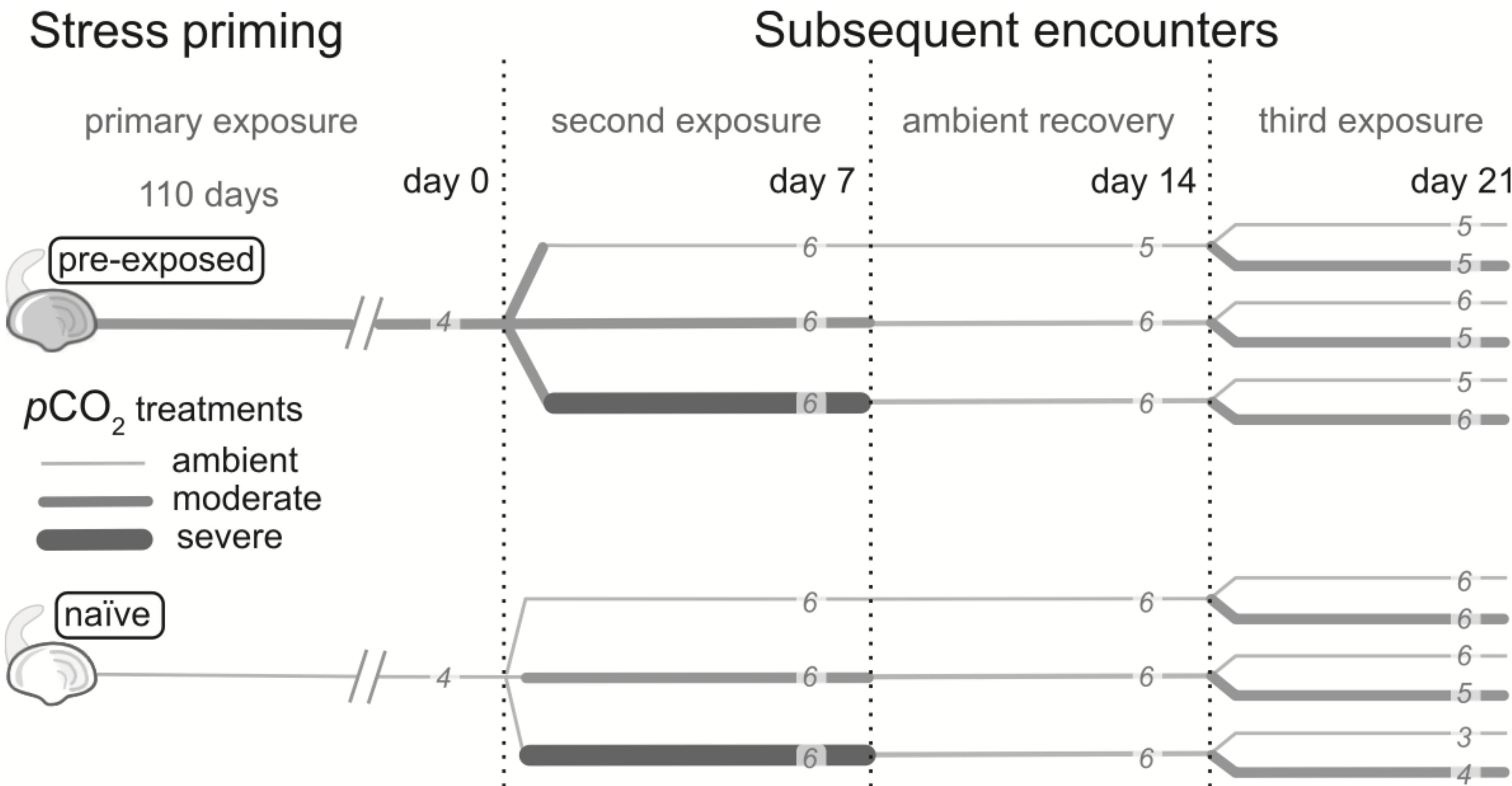


Cross-stress priming success relies on synergistic stress signalling pathways being shared across stresses varying in nature and intensity.

Geoduck Clams

Repeat exposure to hypercapnic seawater modifies growth and oxidative status in a tolerant burrowing clam

Samuel J. Gurr^{1,*}, Shelly A. Wanamaker², Brent Vadopalas³, Steven B. Roberts² and Hollie M. Putnam¹

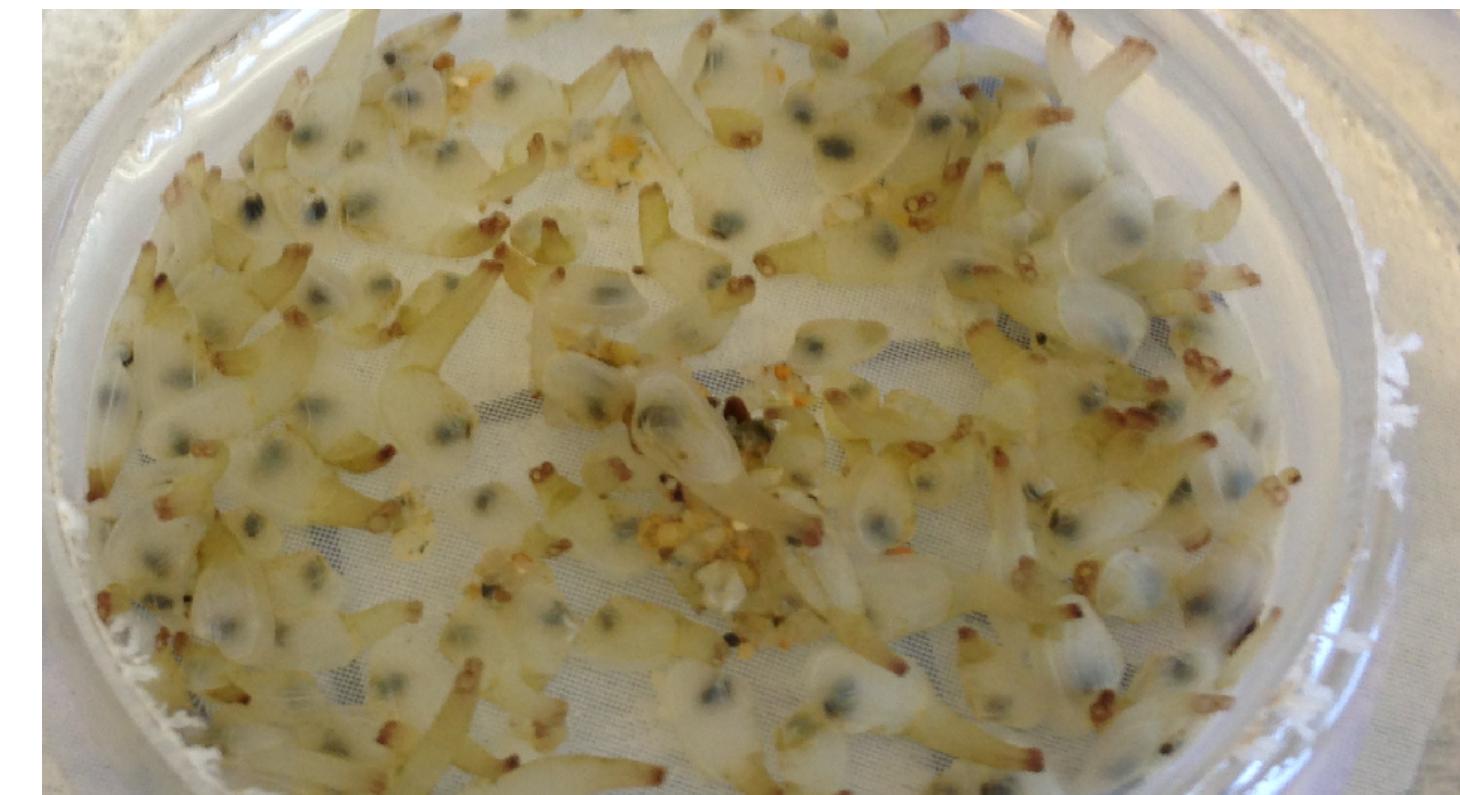


Geoduck Clams

Repeat exposure to hypercapnic seawater modifies growth and oxidative status in a tolerant burrowing clam

Samuel J. Gurr^{1,*}, Shelly A. Wanamaker², Brent Vadopalas³, Steven B. Roberts² and Hollie M. Putnam¹

- Initial conditioning followed by second and third exposure to severe and moderate PCO₂ stress increased respiration rate, organic biomass and shell size, suggesting a stress-intensity-dependent effect on energetics.
- Stress acclimated clams had lower antioxidant capacity compared with clams under ambient conditions, supporting the hypothesis that stress over postlarval-to-juvenile development affects oxidative status later in life.



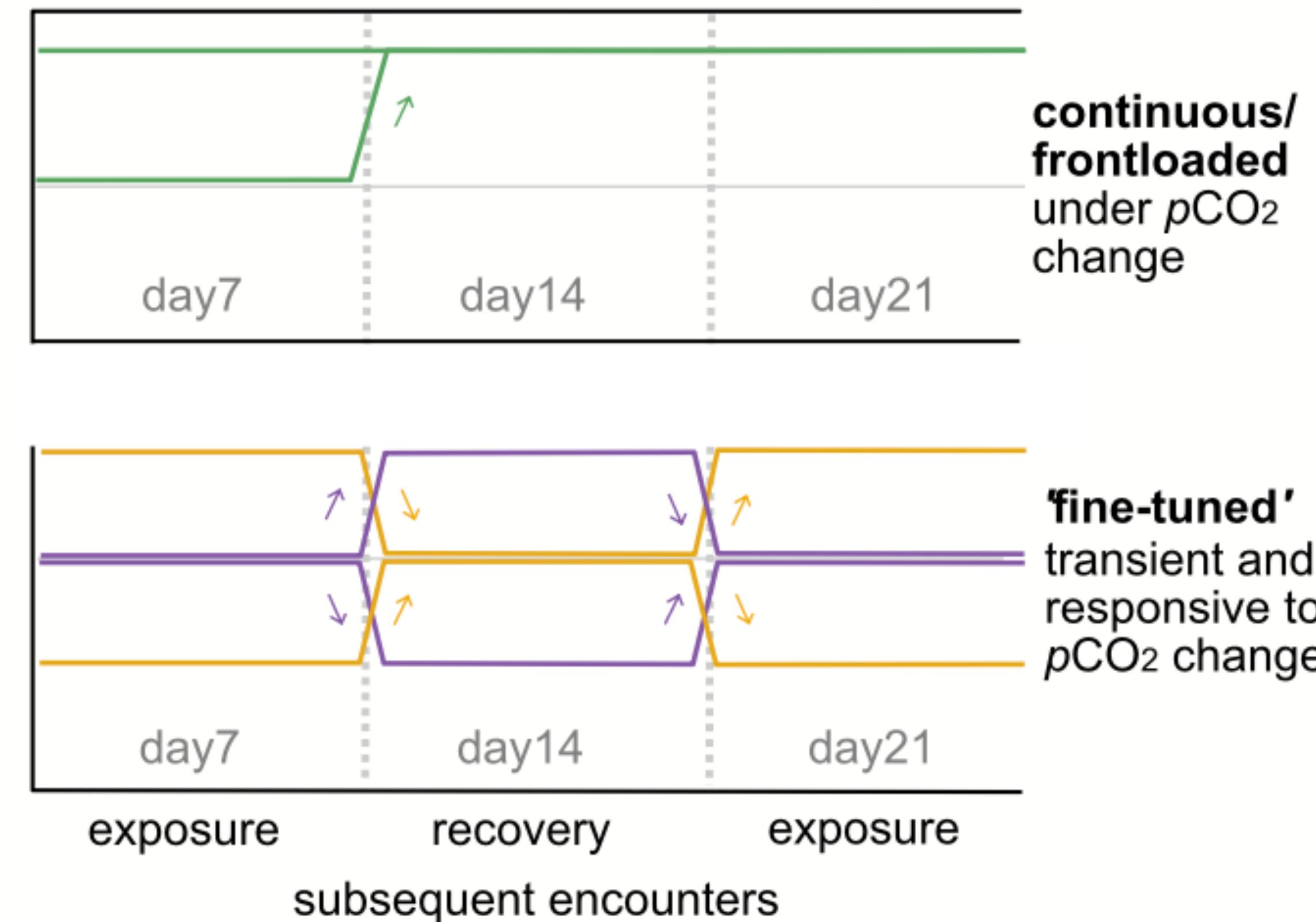
Geoduck Clams

Acclimatory gene expression of primed clams enhances robustness to elevated $p\text{CO}_2$ 

Samuel J. Gurr^{1,2}  | Shelly A. Trigg^{3,4}  | Brent Vadopalas⁵ | Steven B. Roberts³
Hollie M. Putnam¹ 

(b)

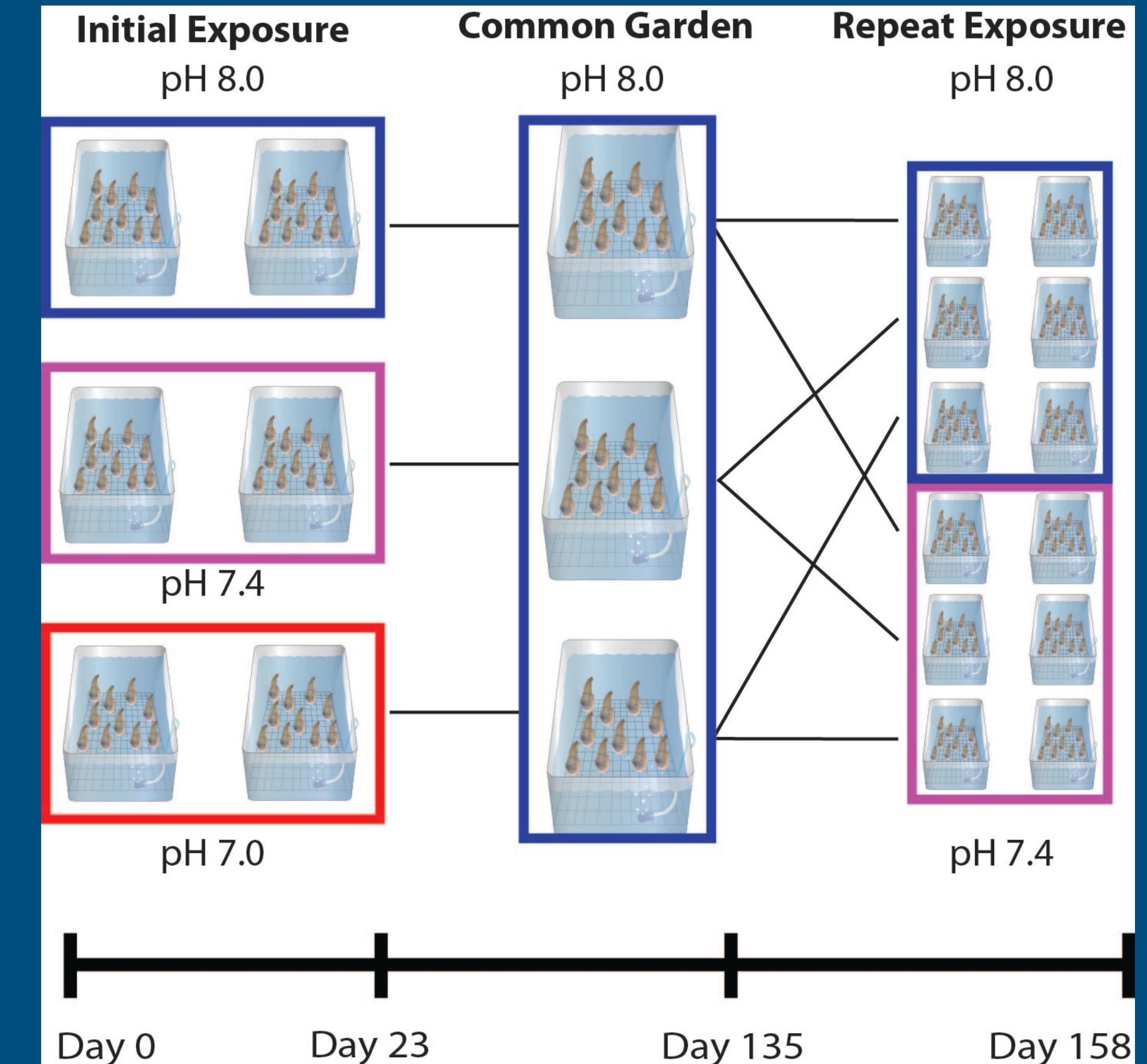
patterns in functional enrichment analysis



GEODUCKS CLAMS

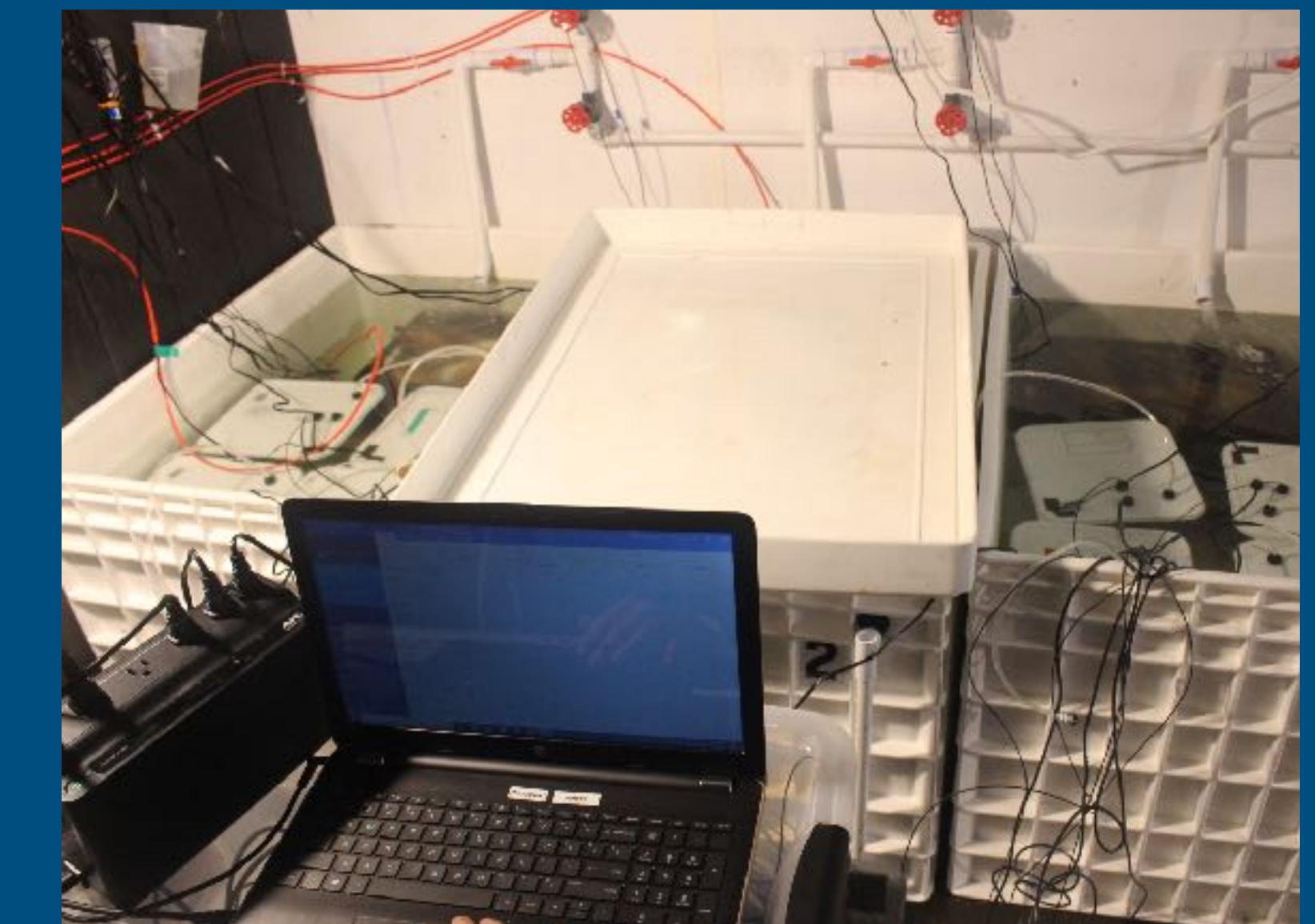


- ▶ Does conditioning to low pH confer tolerance within a generation?



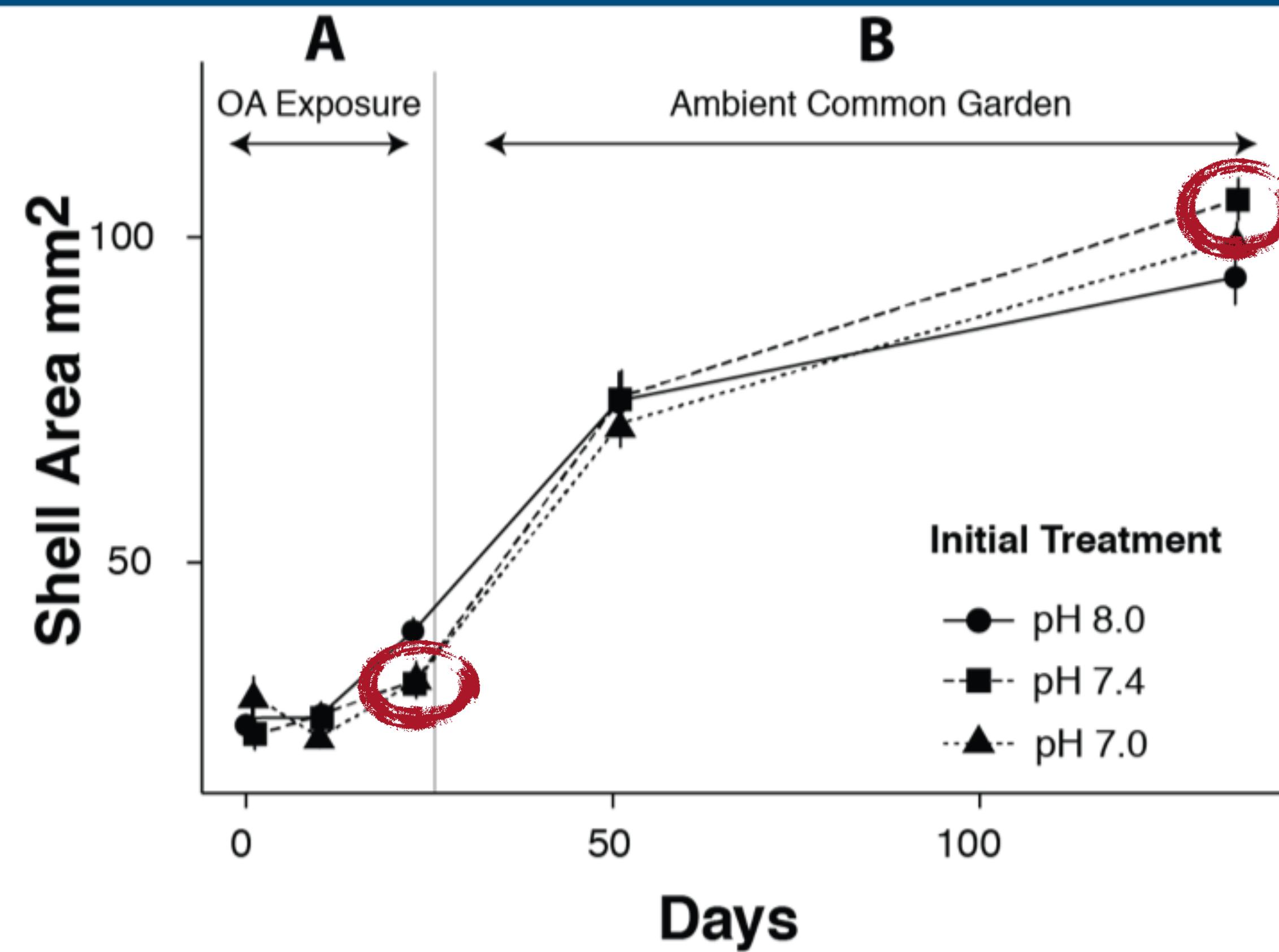
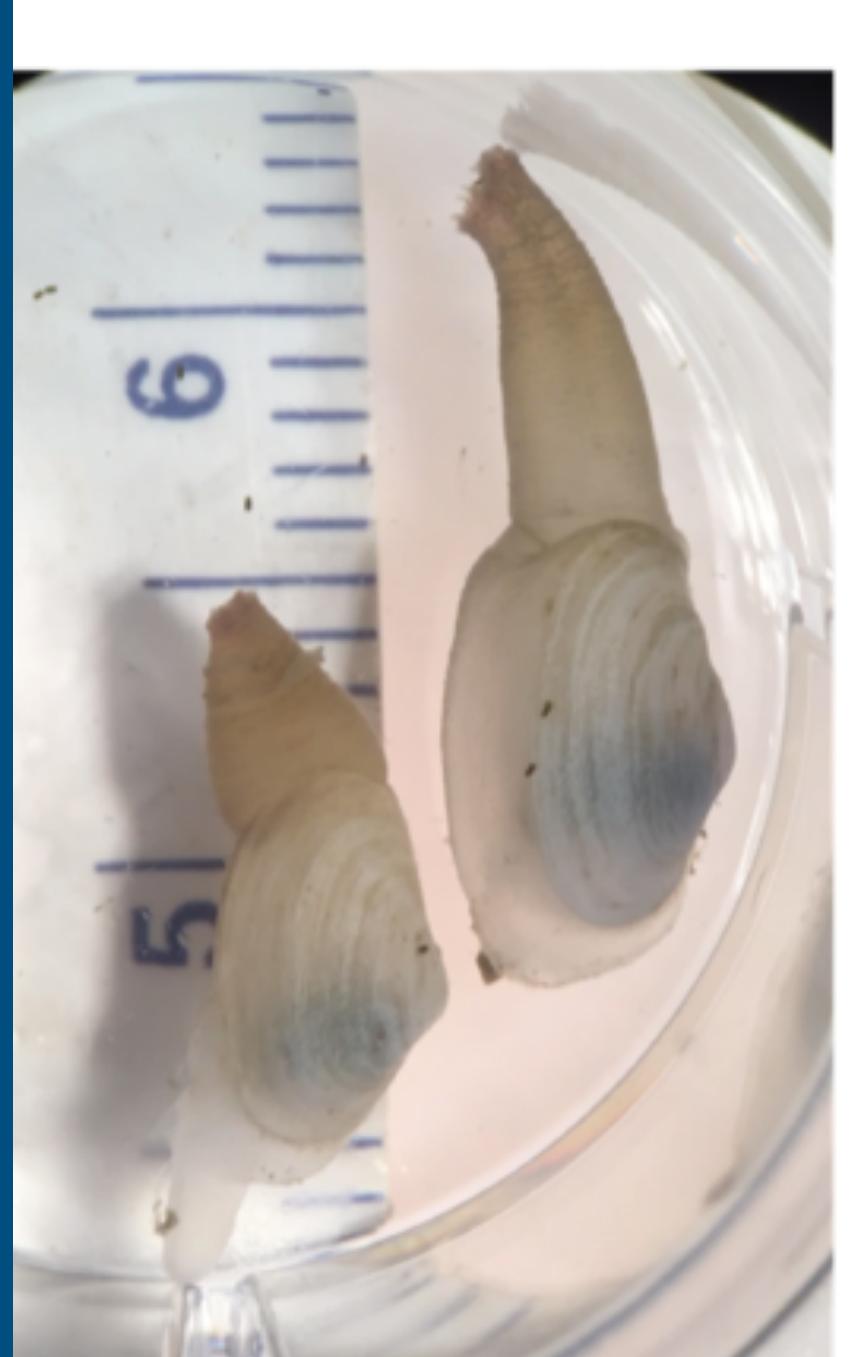
HOLLIE PUTNAM, SAM GURR, BRENT VADOPALAS, SHELLY TRIGG, JAMESTOWN S'KLALLAM TRIBE

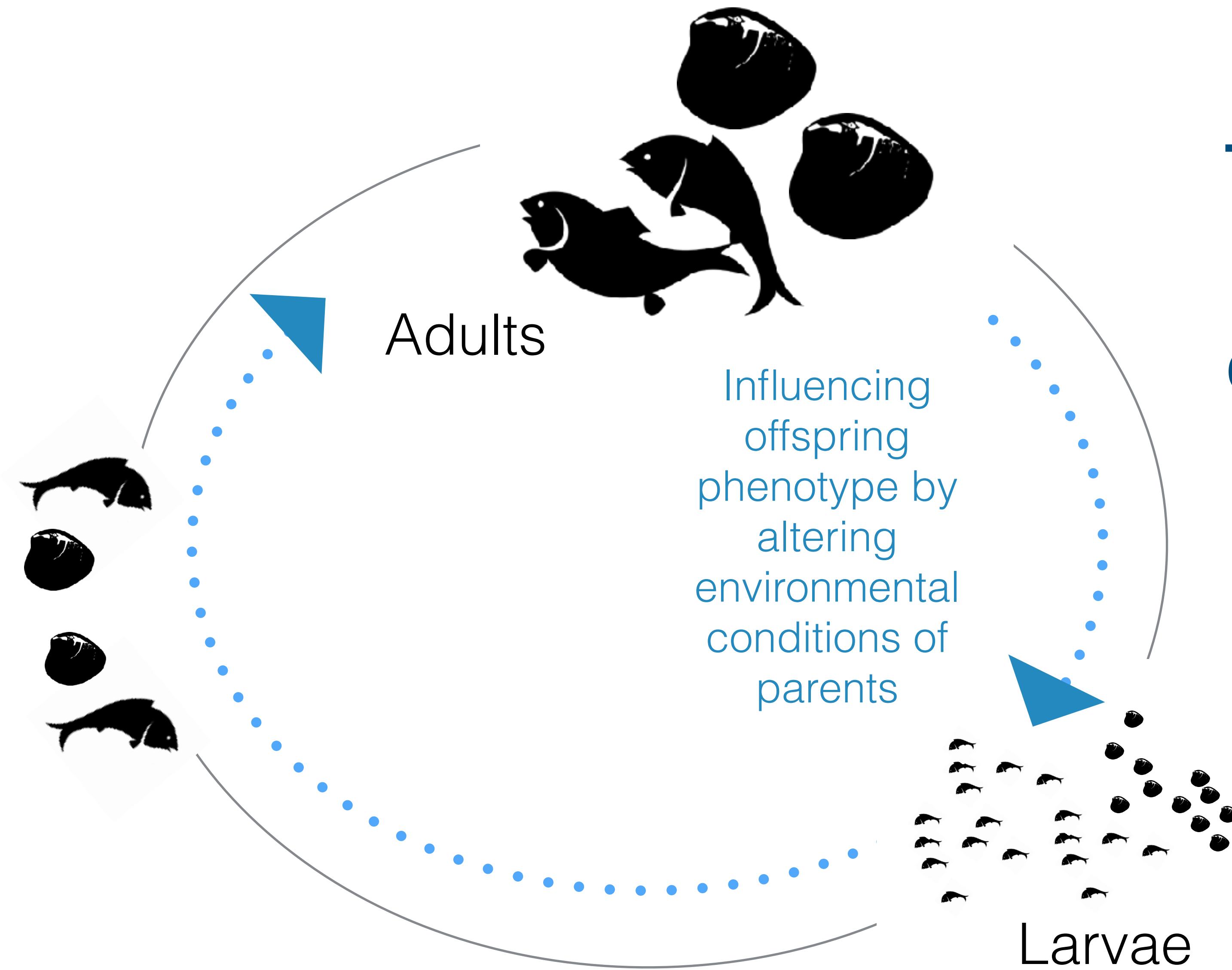
GEODUCKS CLAMS



GEODUCKS CLAMS

► Does conditioning to low pH confer tolerance within a generation?

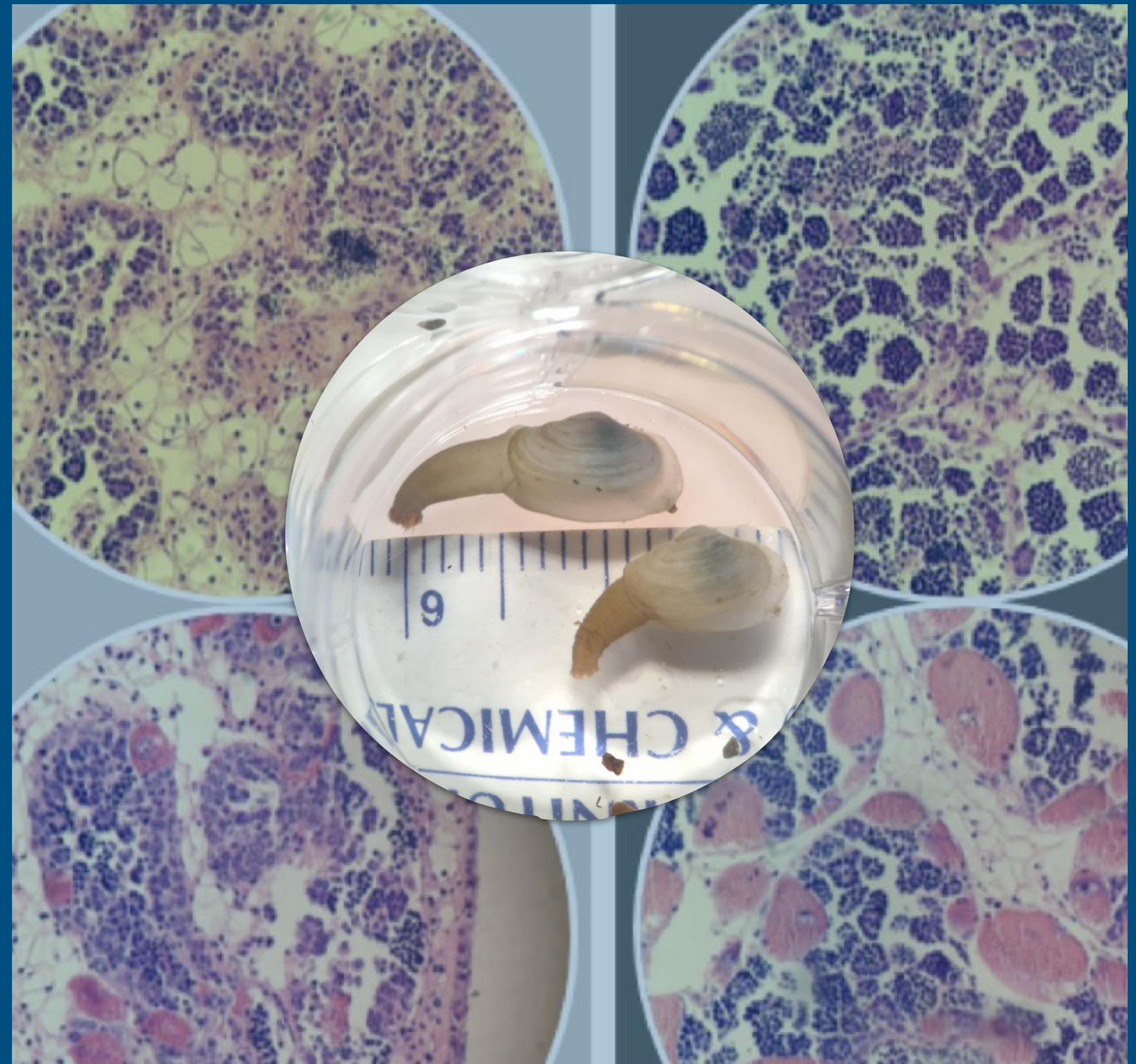




**Transgenerational
Plasticity**
Carry-over effects

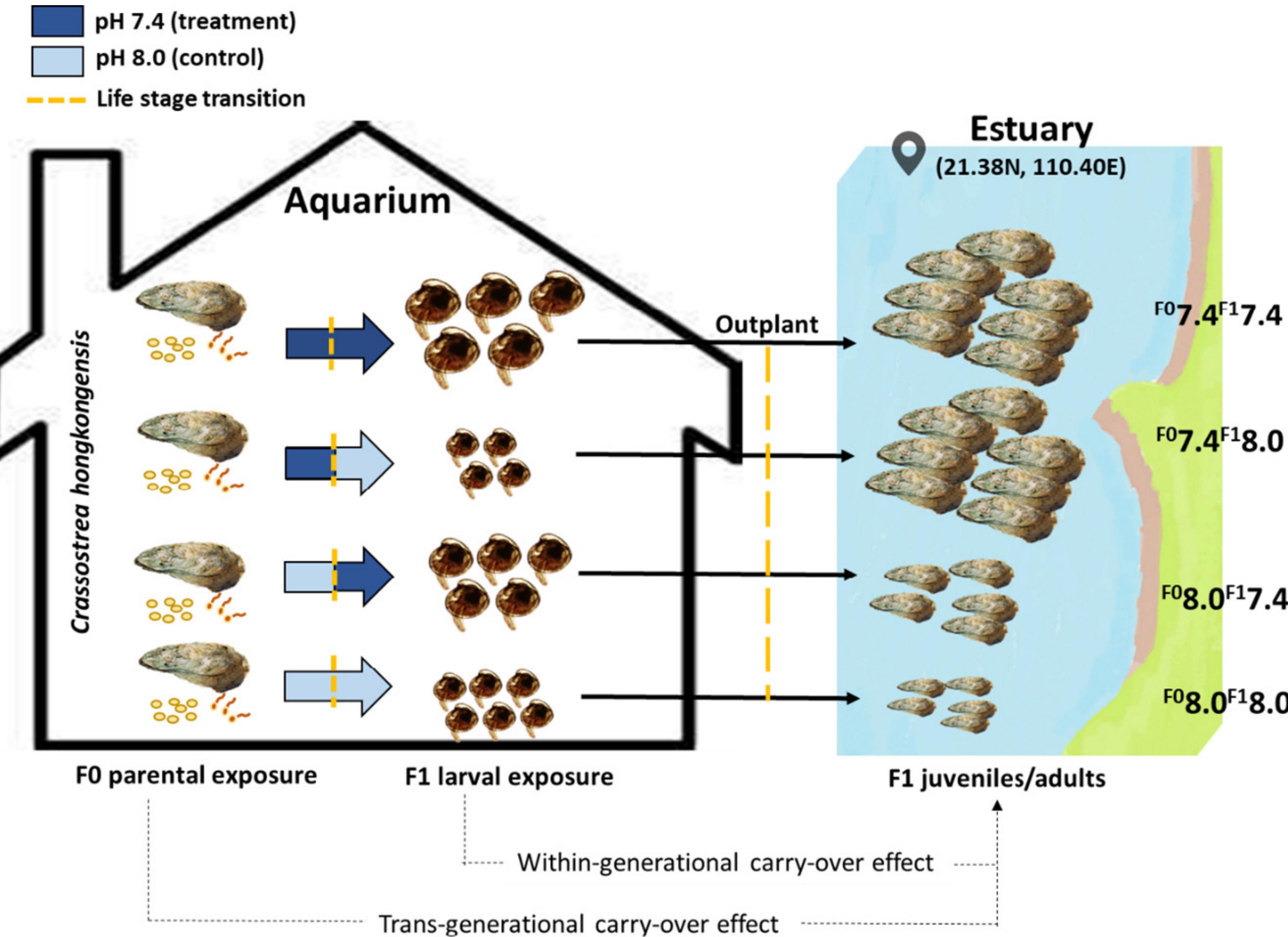
INFLUENCE OF PARENTAL CONDITIONS

- ▶ Selection (various generations)
- ▶ Germ cells are present
- ▶ Maternal provisioning
- ▶ Paternal role?
- ▶ Beneficial versus detrimental
- ▶ *Mechanisms at play*



Crassostrea hongkongensis

Transgenerational responses to seawater pH in the edible oyster, with implications for the mariculture of the species under future ocean acidification



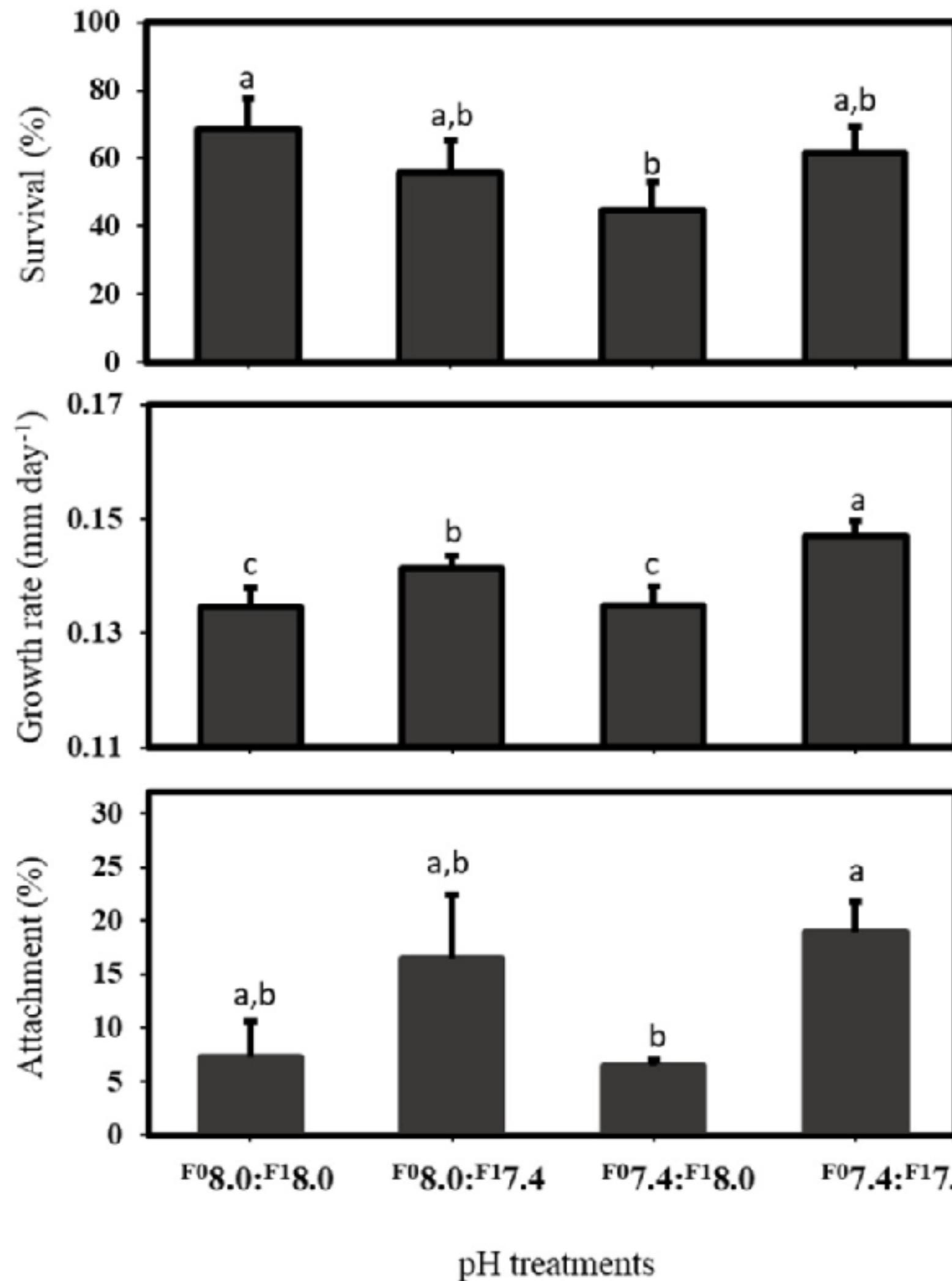
Crassostrea *hongkongensis*

Transgenerational responses to
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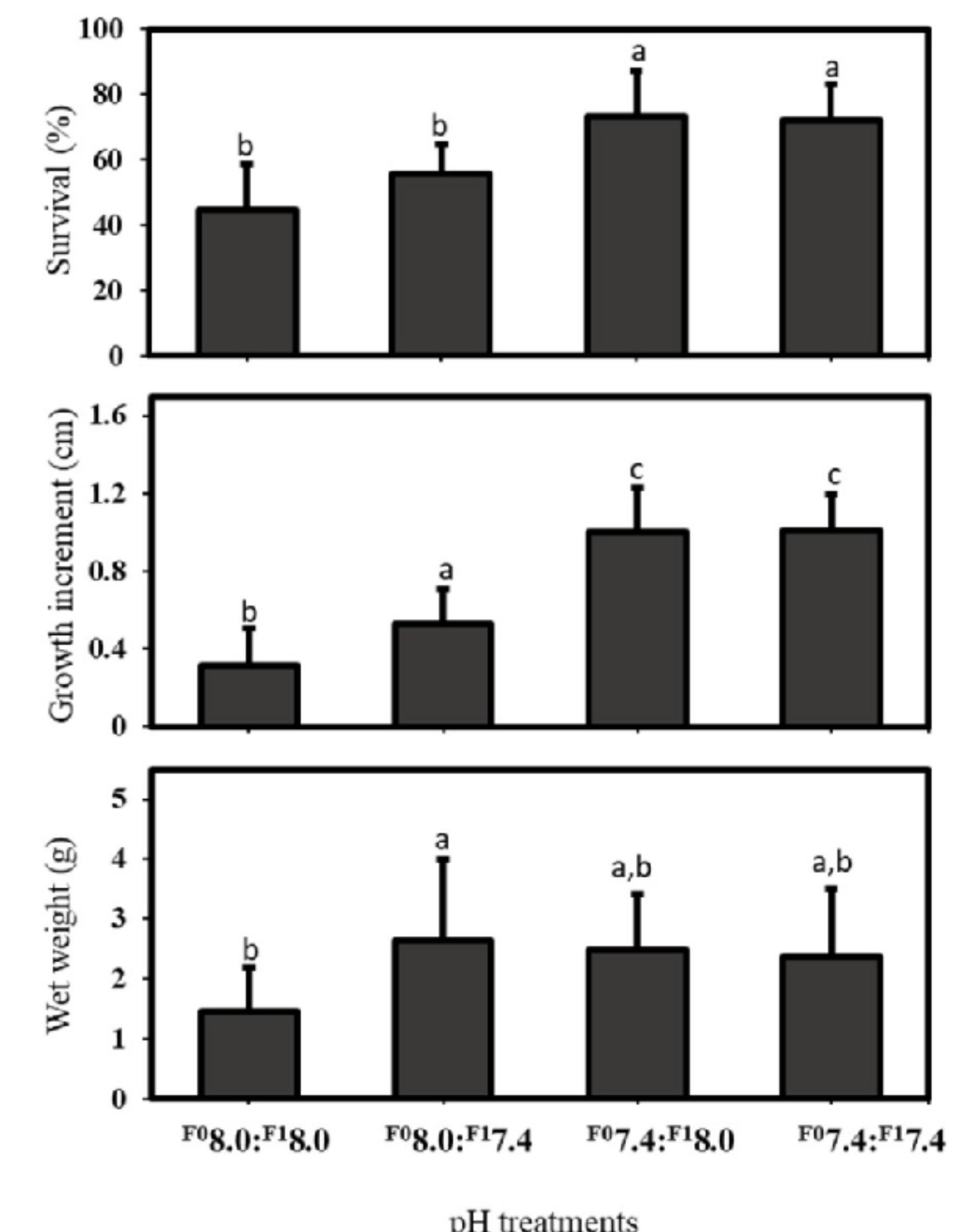
Yong-Kian Lim¹, Xin Dang¹, Vengatesen Thiagarajan  



A. F₁ larvae



B. F₁ juveniles



EFFECTS OF TEMPERATURE AND OA IN OLYMPIA OYSTER POPULATIONS

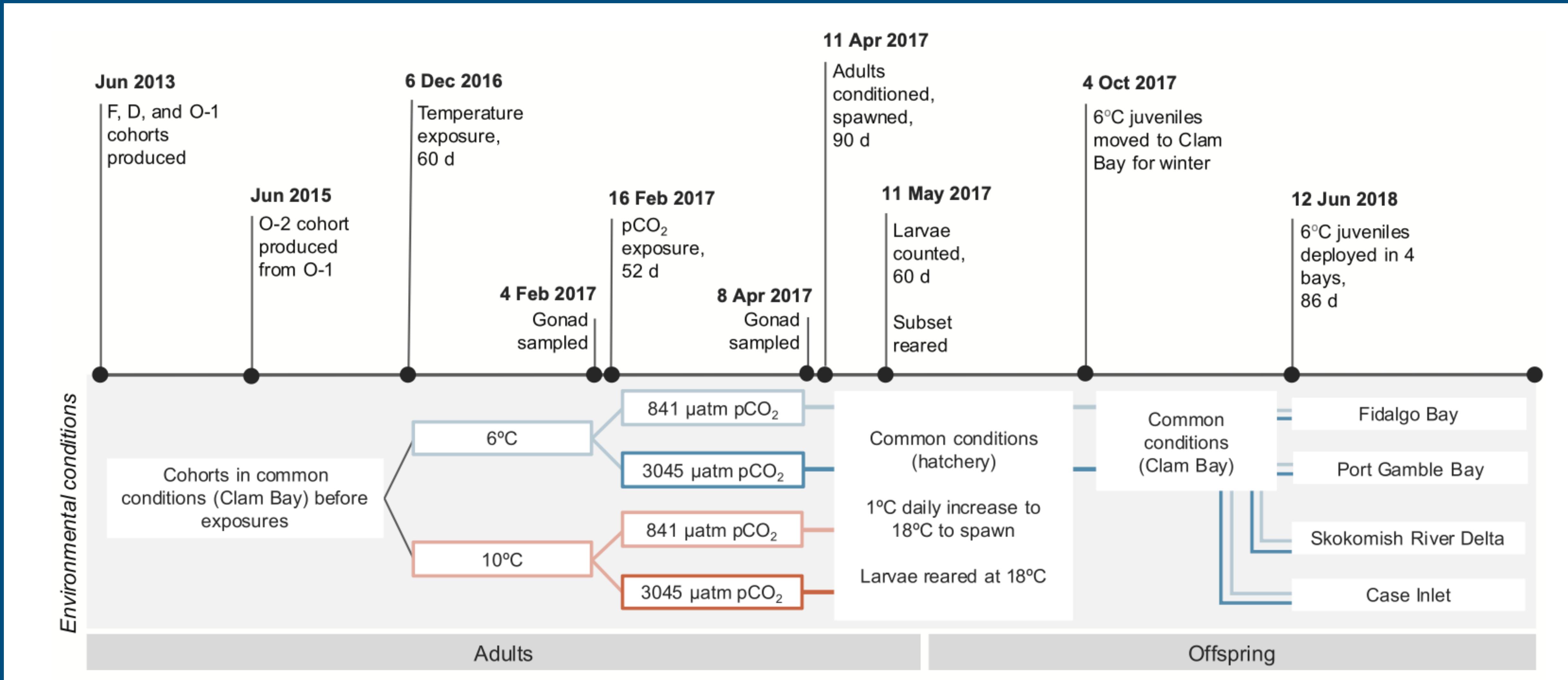


- ▶ Oysters were held at two temperature regimes (6°C and 10°C) for 60 days in December
- ▶ A differential pCO₂ exposure was carried out after the temperature treatment ended. Held at ambient pCO₂ (841 µatm) or high pCO₂ (3045 µatm) for 52 days, during the Winter.

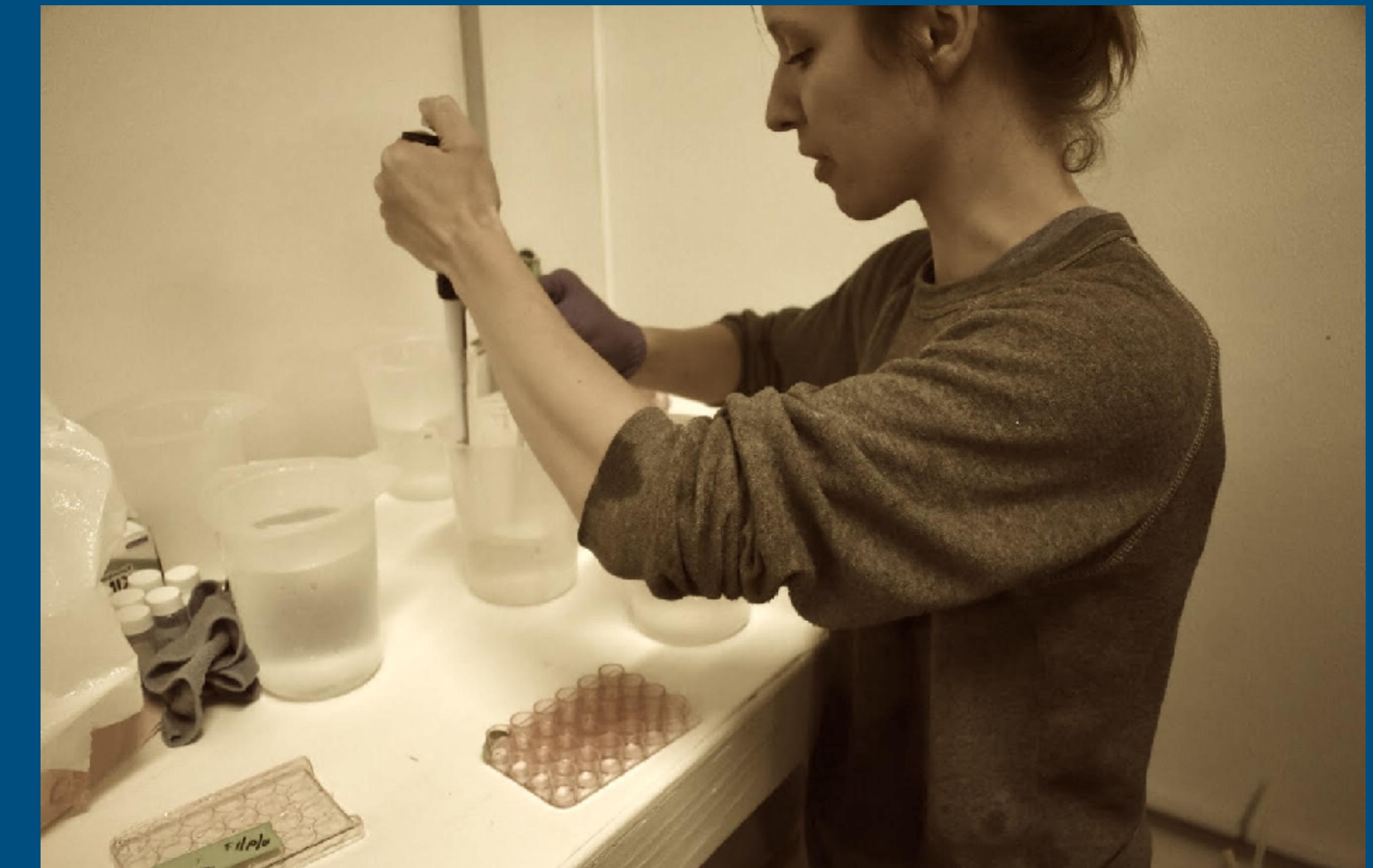
Carryover effects of temperature and pCO₂ across multiple Olympia oyster populations

LAURA H. SPENCER,¹ YAAMINI R. VENKATARAMAN,¹ RYAN CRIM,² STUART RYAN,² MICAH J. HORWITZ,³ AND STEVEN B. ROBERTS^{1,4}

TEXT



EFFECTS OF TEMPERATURE AND OA IN OLYMPIA OYSTER POPULATIONS

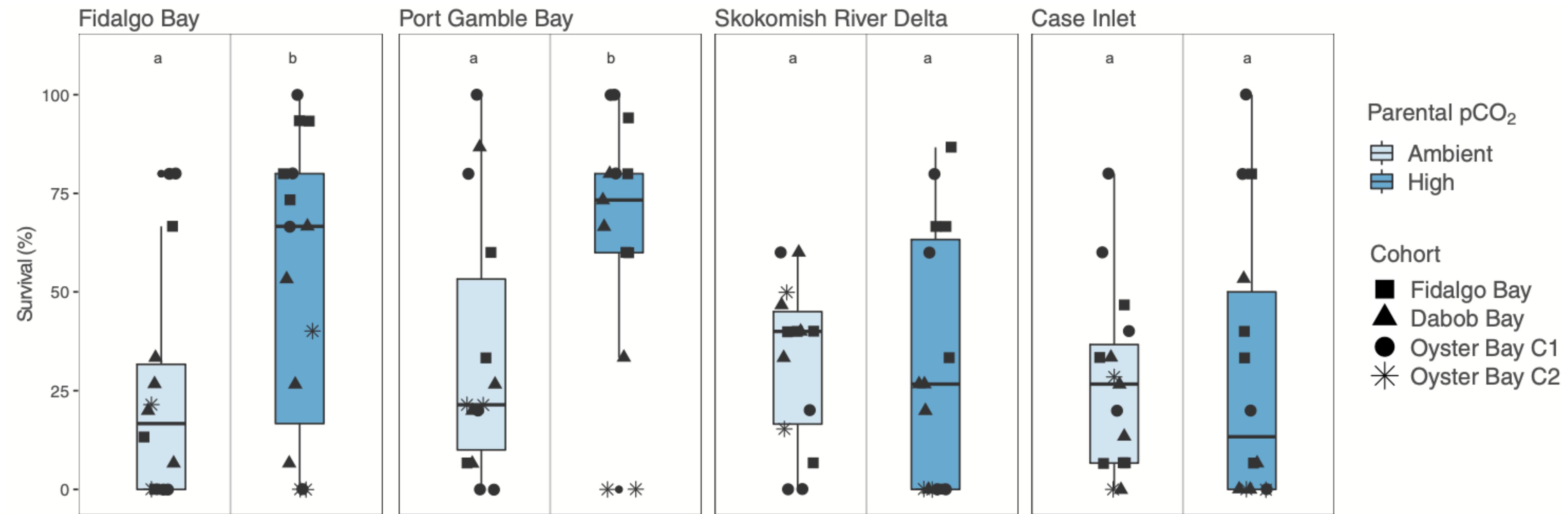


EFFECTS OF TEMPERATURE AND OA IN OLYMPIA OYSTER POPULATIONS



- ▶ Larval release occurred earlier in warm-exposed oysters
- ▶ Winter warming conditions increased larval production
- ▶ No effects on larval survival were detected
- ▶ **Juveniles of parents exposed to elevated pCO₂ had higher survival rates in the natural environment**

EFFECTS OF OA IN OLYMPIA OYSTER POPULATIONS



Latent effects of winter warming on Olympia oyster reproduction and larval viability

Laura H. Spencer ^{a,*}, Erin Horkan ^b, Ryan Crim ^b, Steven B. Roberts ^a

^a University of Washington, School of Aquatic and Fishery Sciences, 1122 NE Boat St, Seattle, WA 98105, United States

^b Puget Sound Restoration Fund, 8001 NE Day Rd W, Bainbridge Island, WA 98110, United States

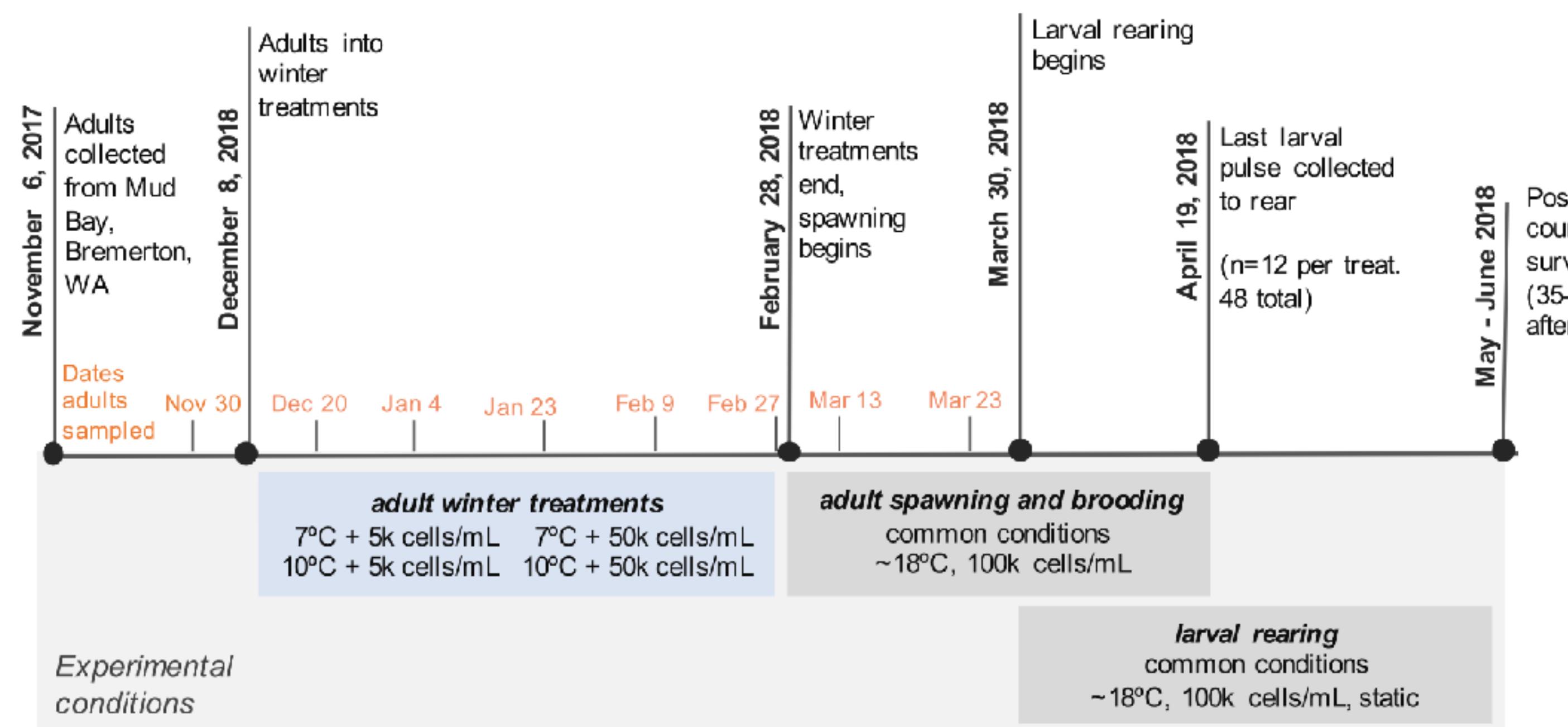
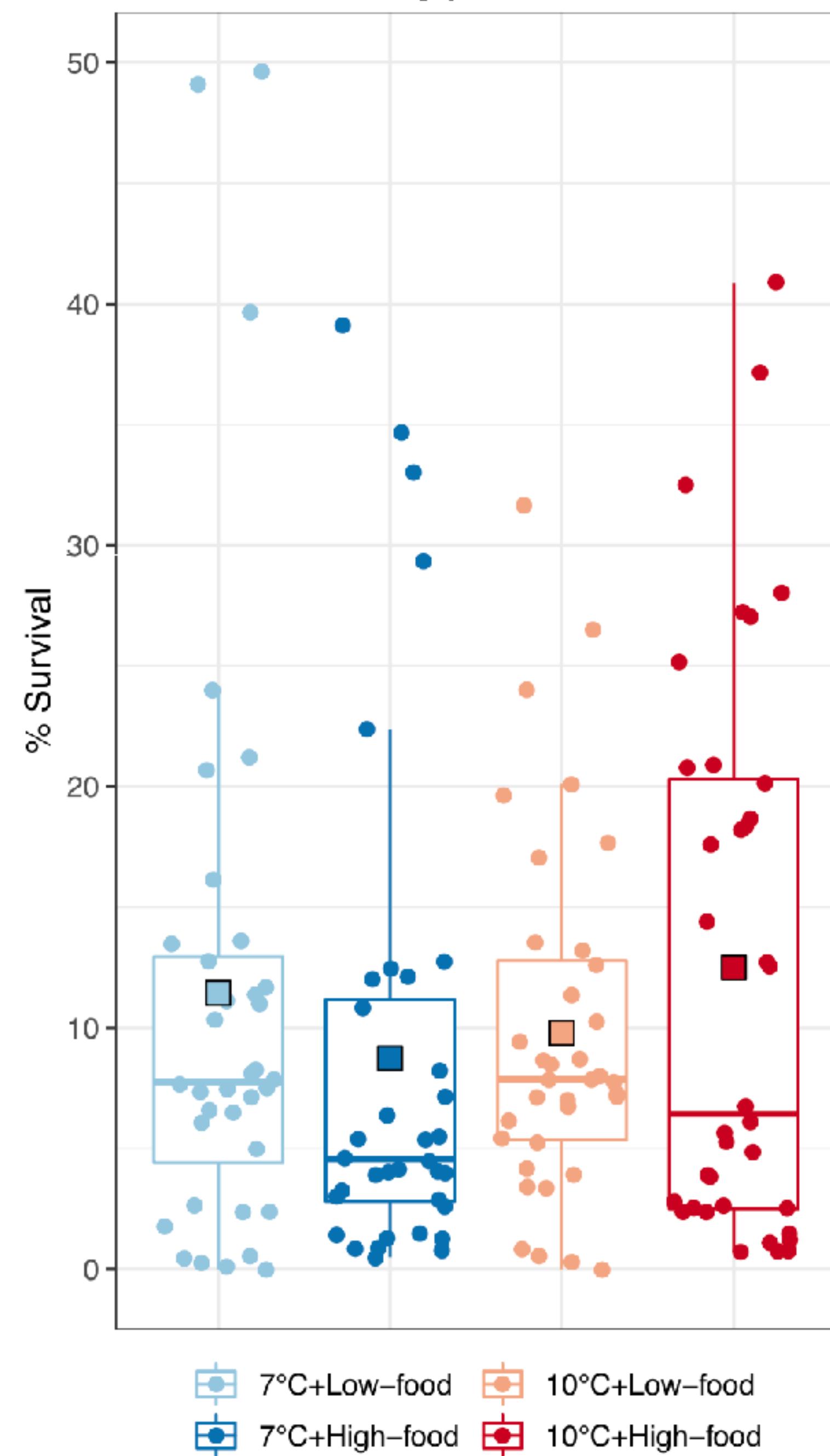


Fig. 1. Experimental timeline. Cells/mL indicate the concentrations of live algae given to oysters at various stages.

Larval survival by parental treatment



Environmental Memory



CAN BE BENEFICIAL
STAGE DEPENDENT
ACROSS GENERATION
IMPLICATIONS ...

Environmental Memory



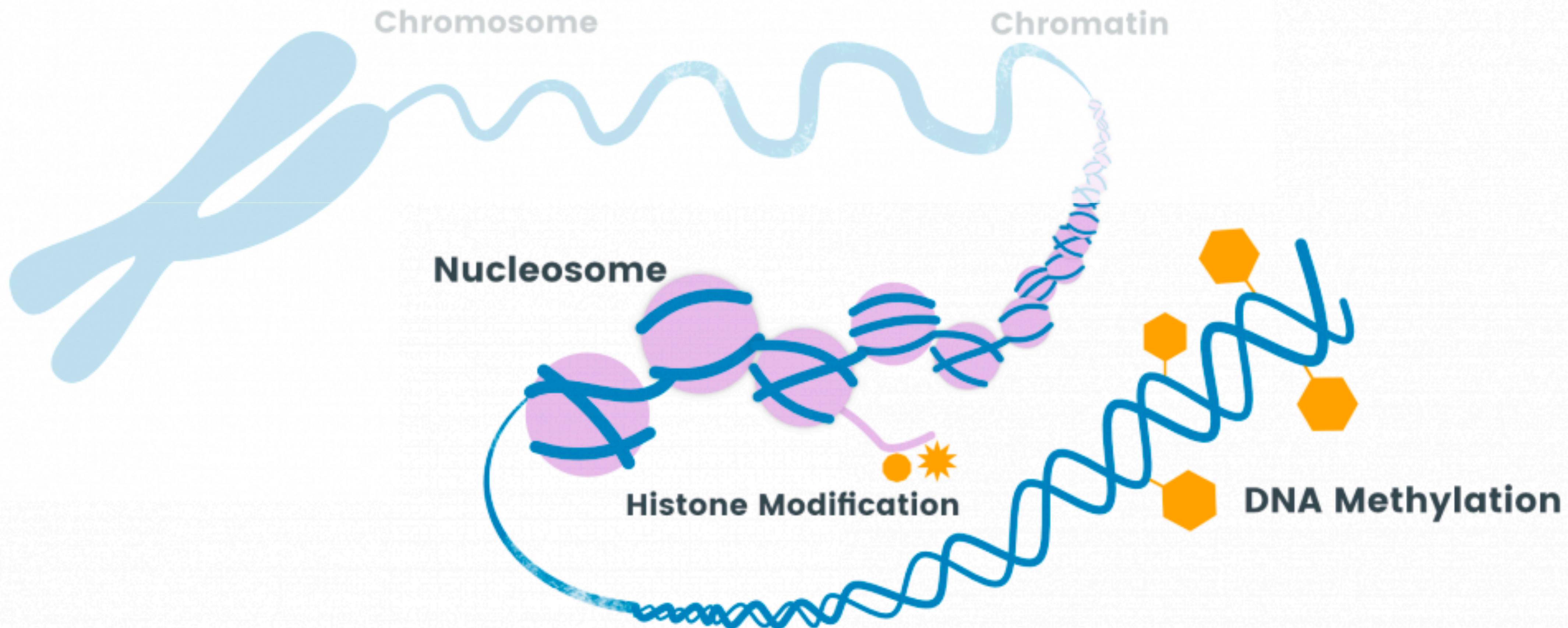
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IMPLICATIONS . . .

Epigenetic Mechanisms

Epigenetics

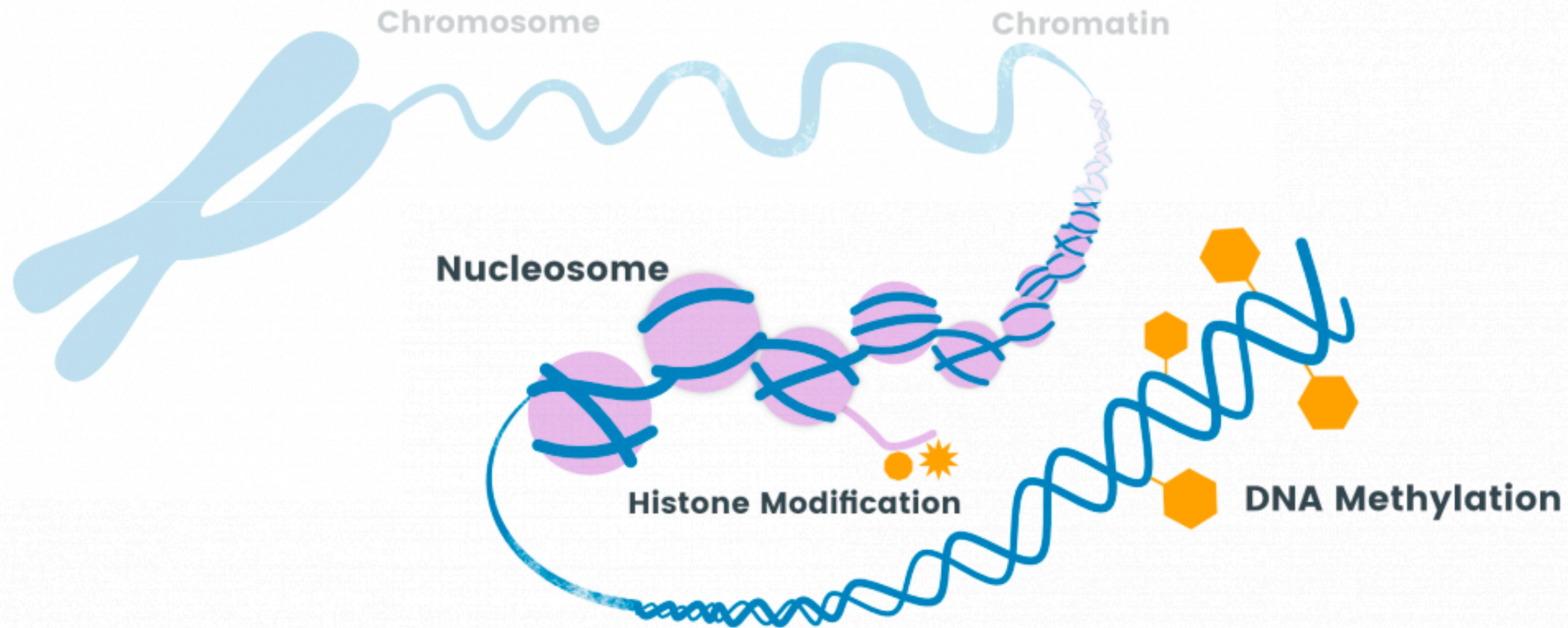
WHAT IS EPIGENETICS?

ALTERS THE PHENOTYPE (WITHOUT CHANGING DNA CODE); HERITABLE

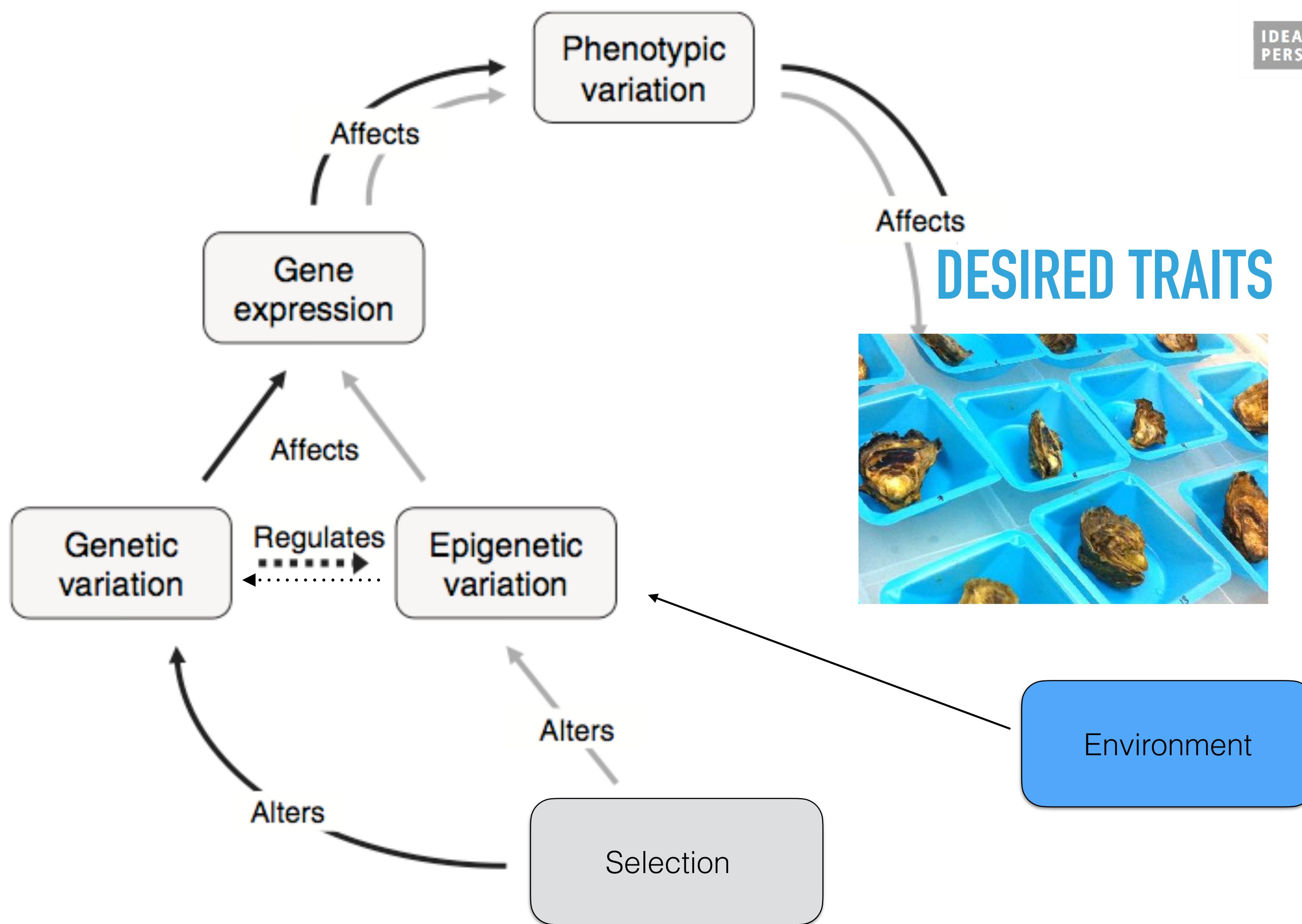


WHAT IS EPIGENETICS?

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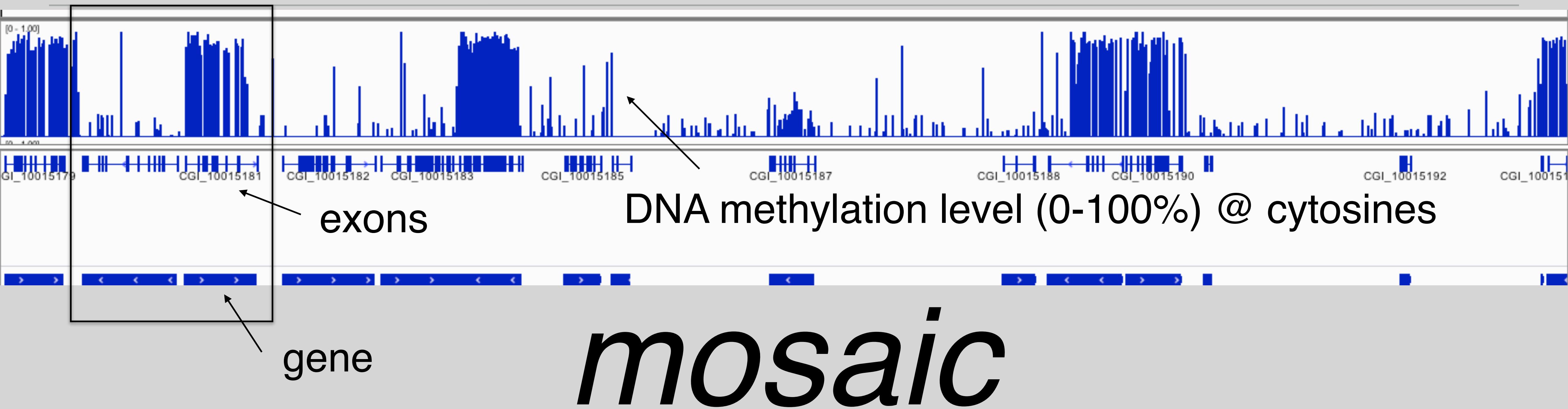
CAN BE INDUCED WITH ENVIRONMENTAL MANIPULATION



IDEA AND
PERSPECTIVE

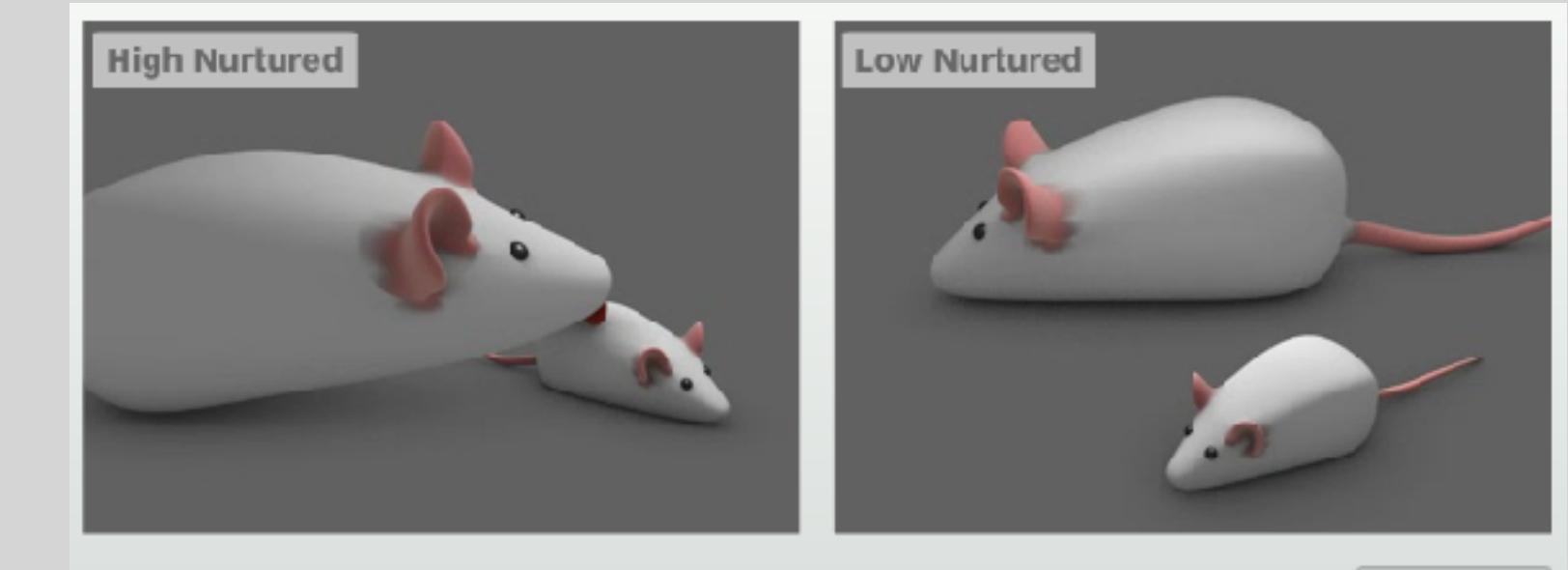
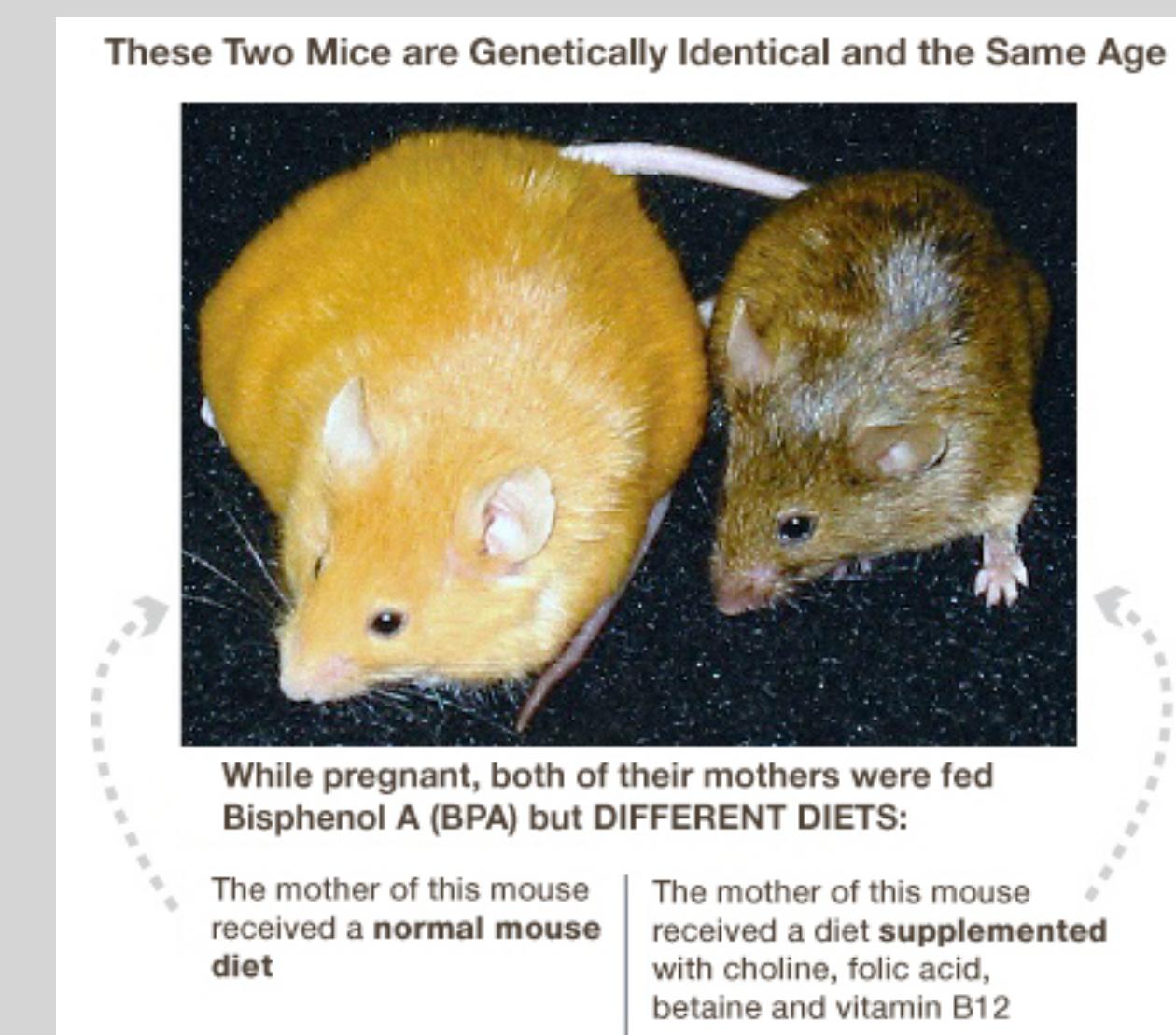
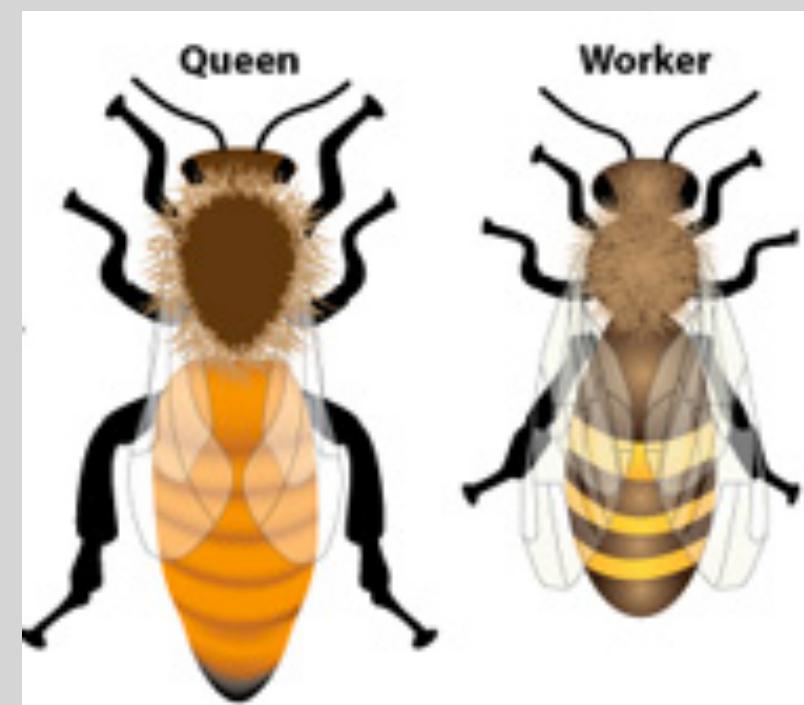
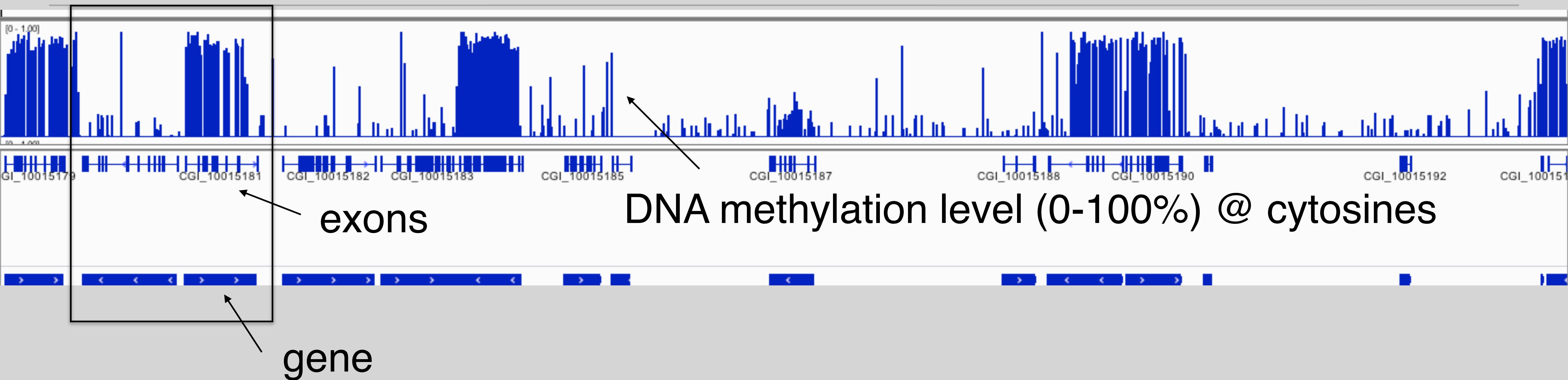
Epigenetics for ecologists

METHYLATION LANDSCAPE IN MARINE INVERTEBRATES



associated with gene bodies

METHYLATION LANDSCAPE IN MARINE INVERTEBRATES



Four Dimensionalities



Reliable Transcription

Spurious Transcription

Targeted Regulation

Stochastic Regulation

Four Dimensionalities

- Evolutionary
- Life History Driven
- Constitutive

Reliable Transcription

Spurious Transcription

- Distinct Lineage
- Experiential
- Inducible

Targeted Regulation

Stochastic Regulation

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Reliable Transcription

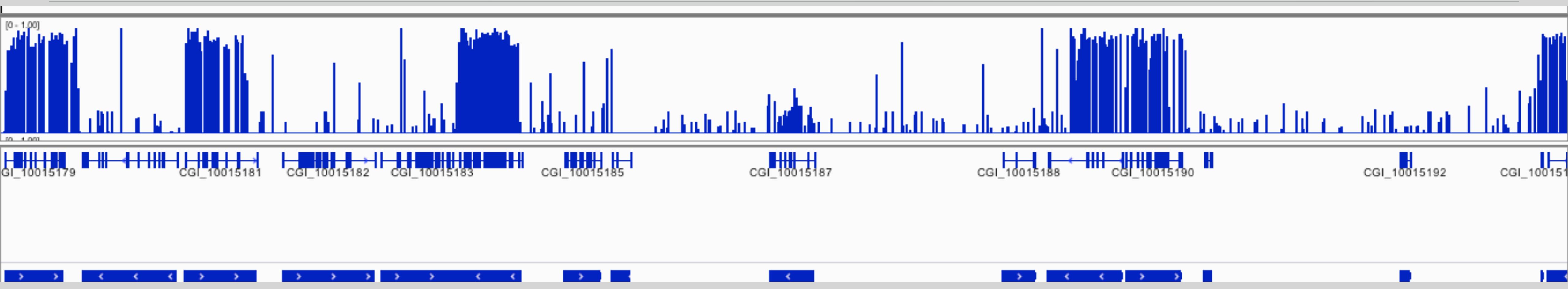
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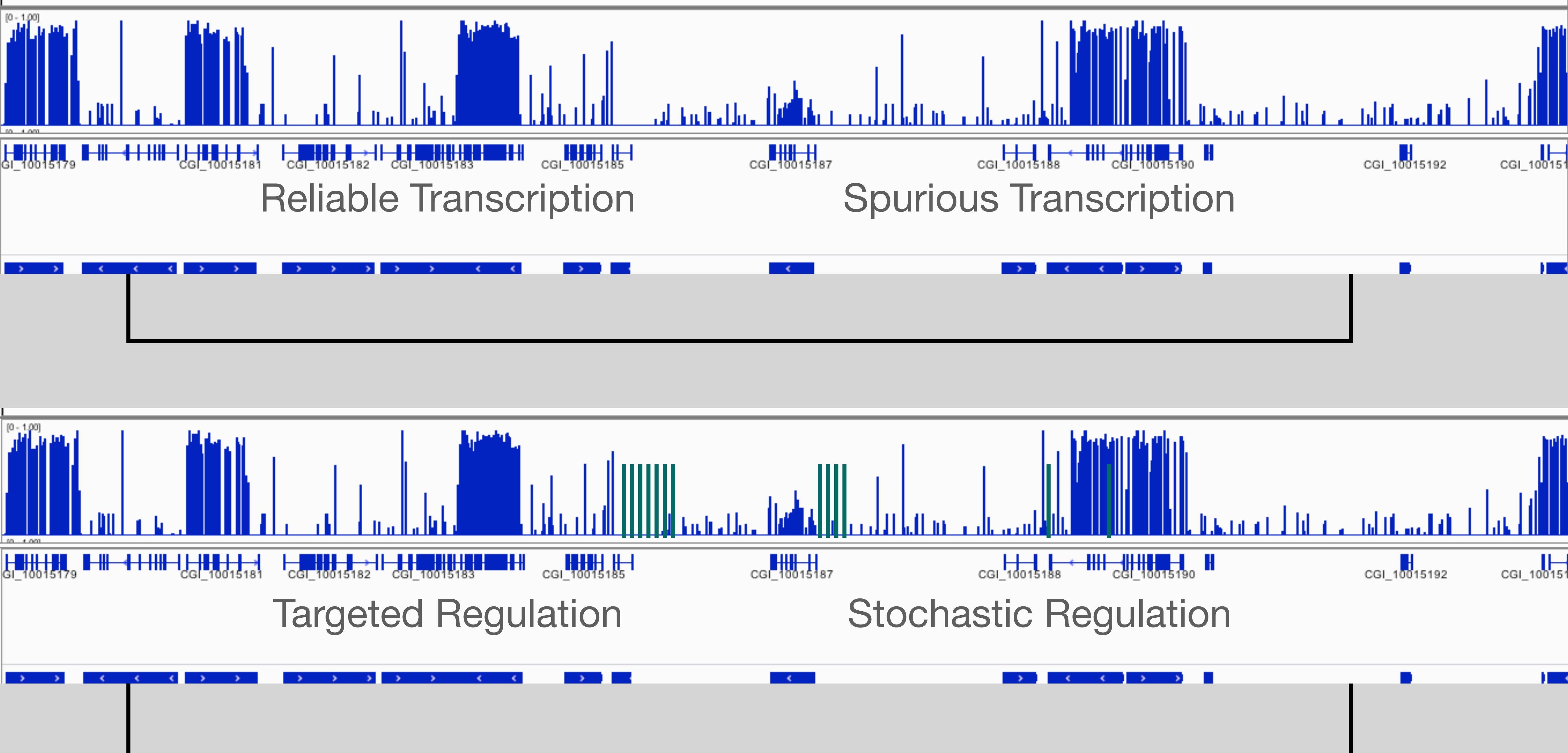
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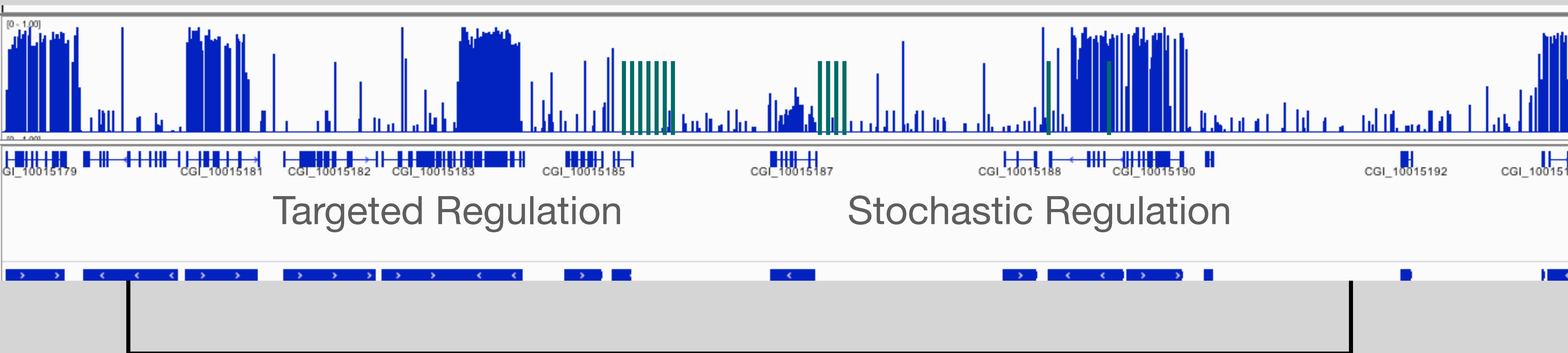
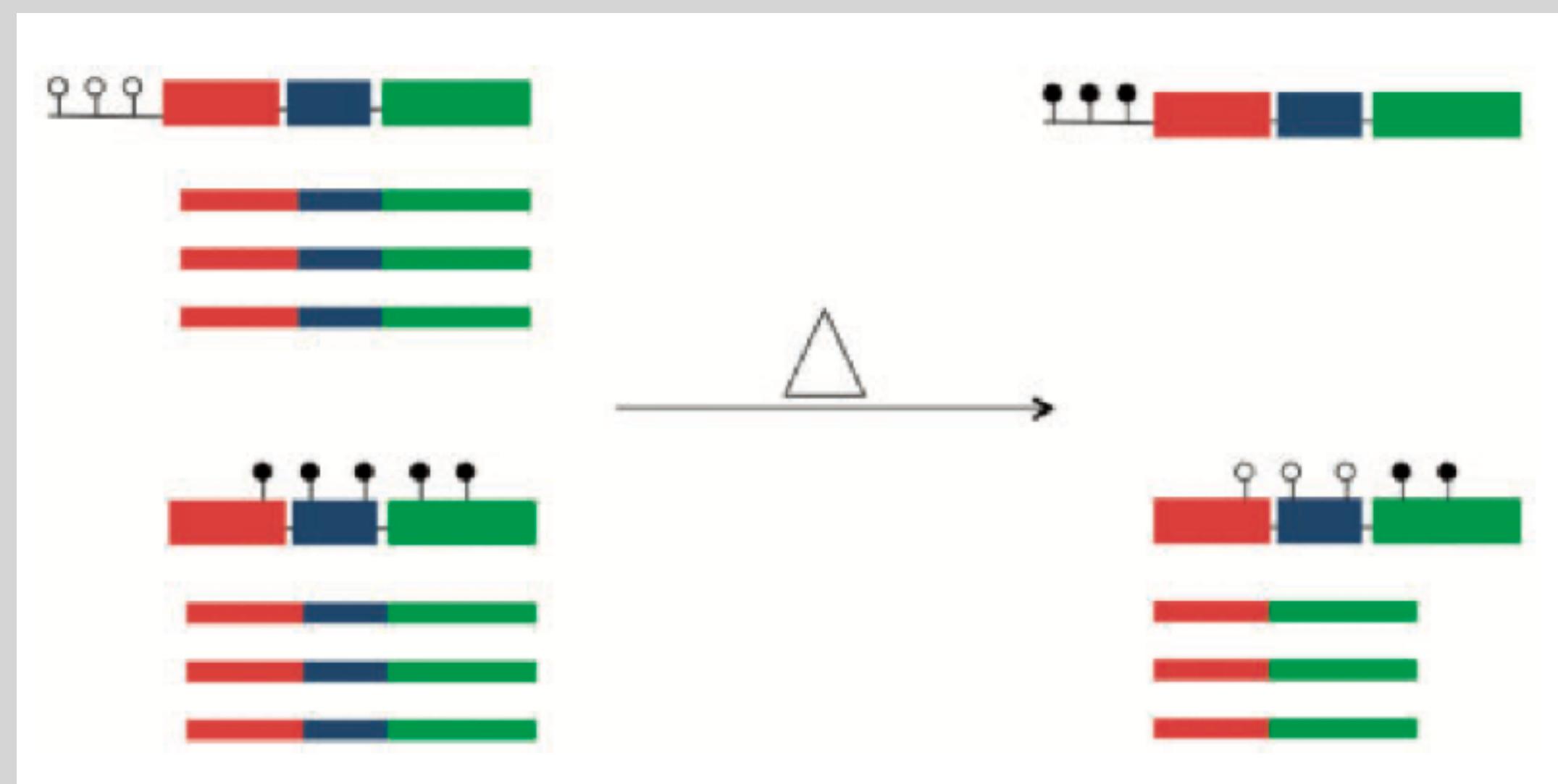
METHYLATION LANDSCAPE IN MARINE INVERTEBRATES



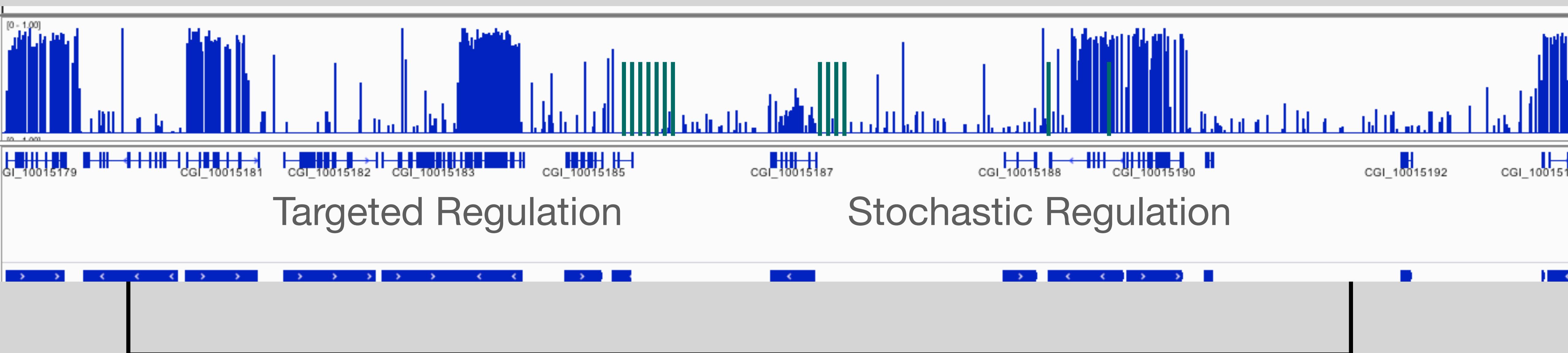
METHYLATION LANDSCAPE IN MARINE INVERTEBRATES



METHYLATION LANDSCAPE IN MARINE INVERTEBRATES

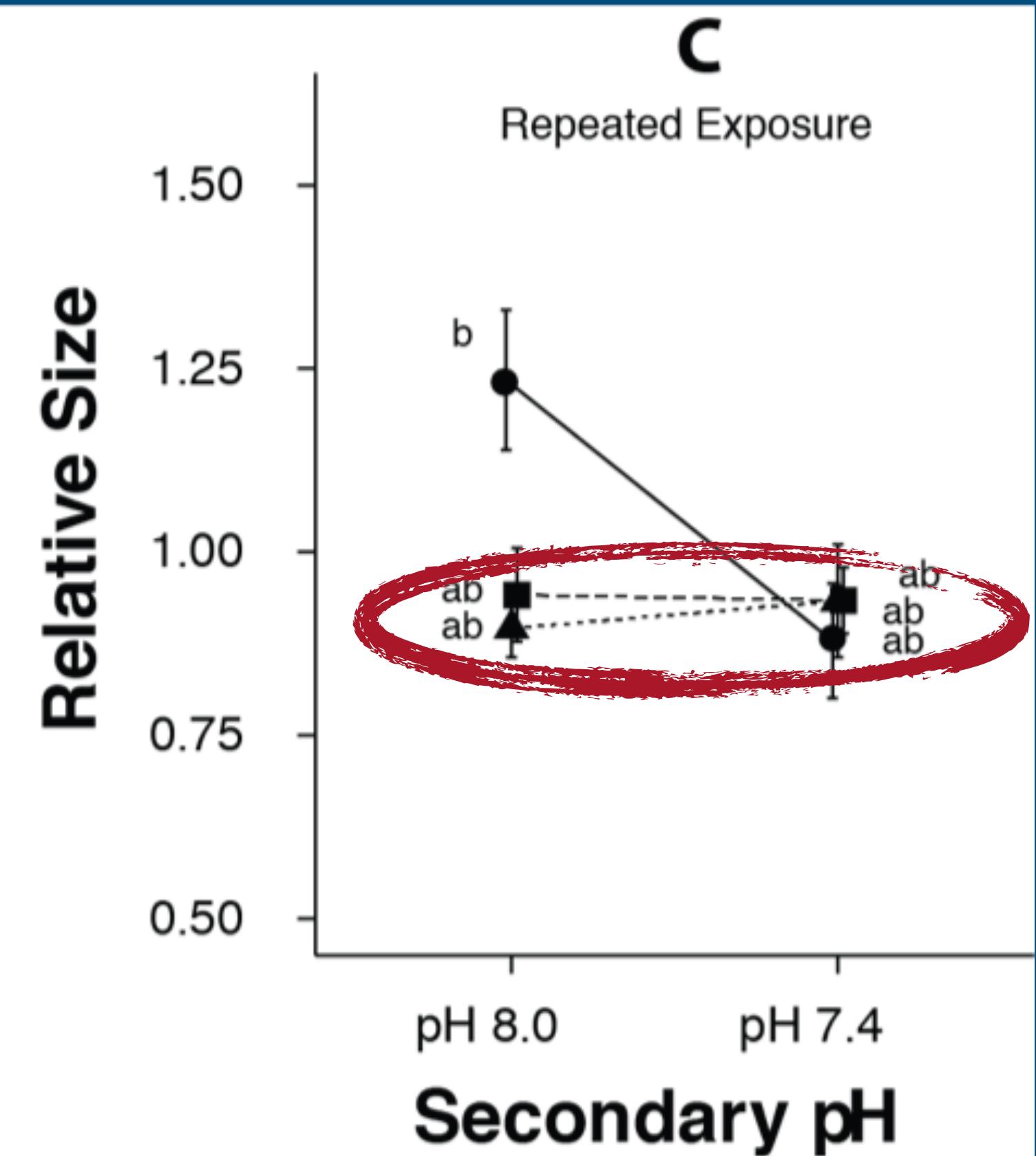
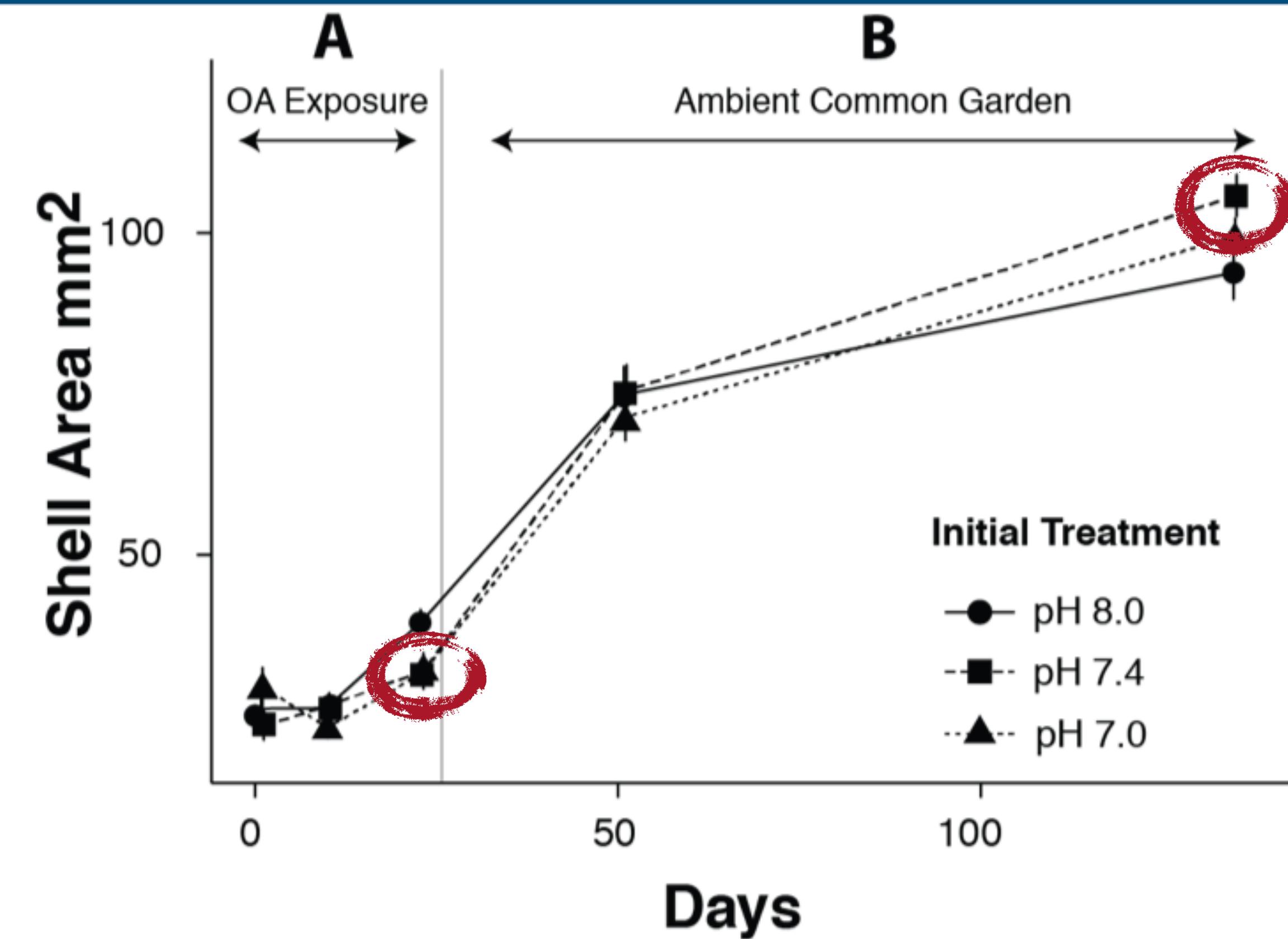


METHYLATION LANDSCAPE IN MARINE INVERTEBRATES



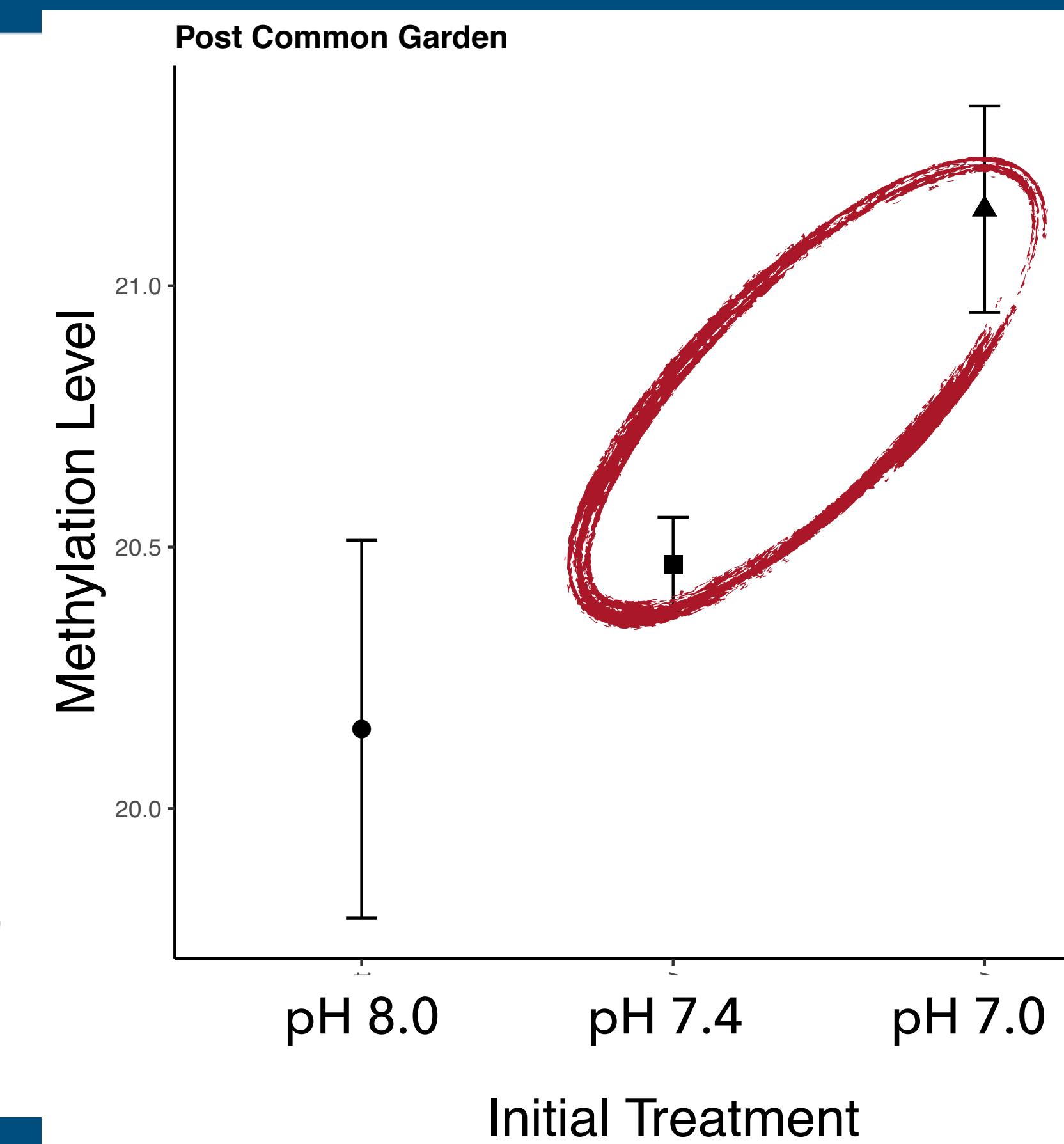
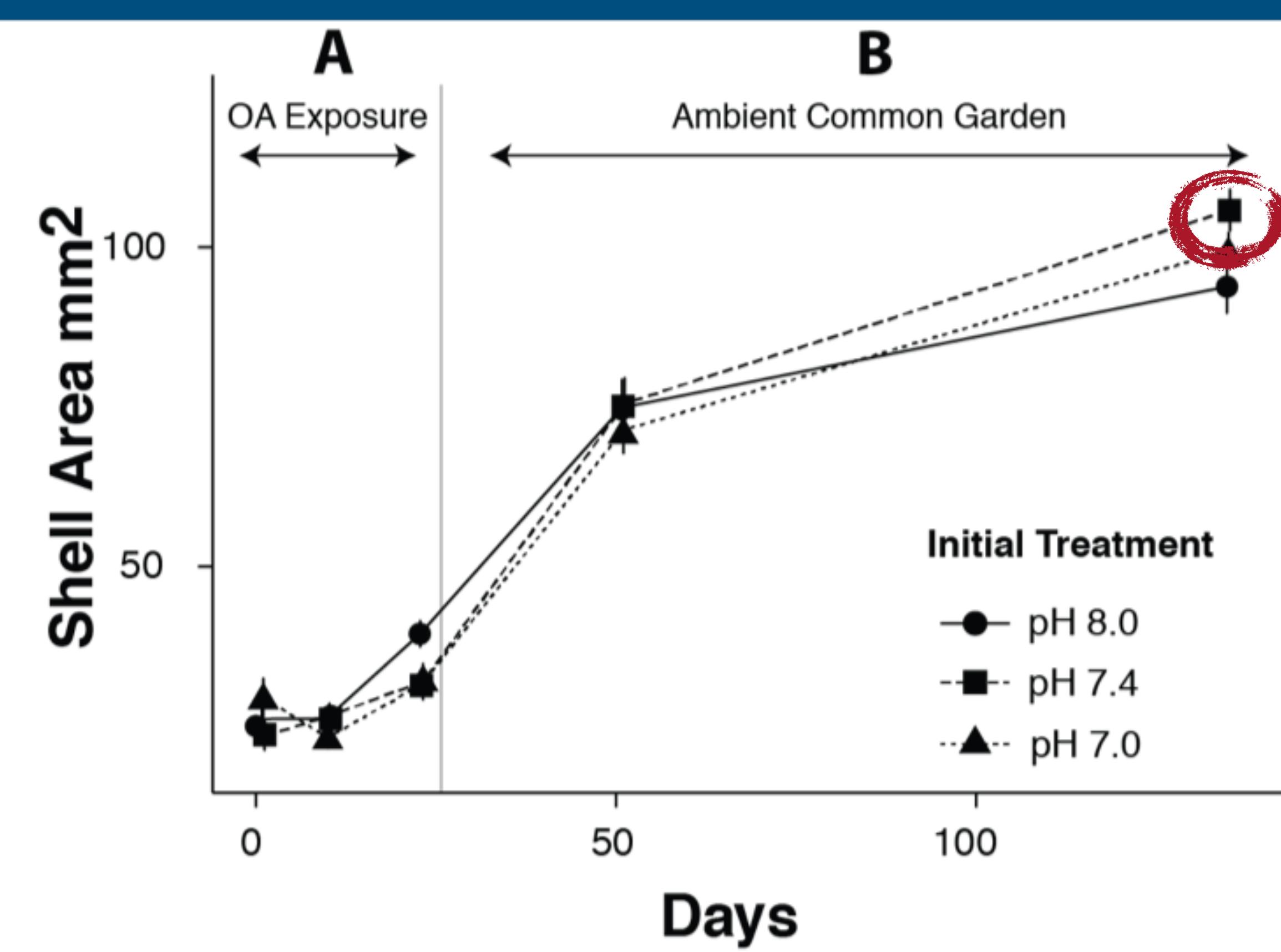
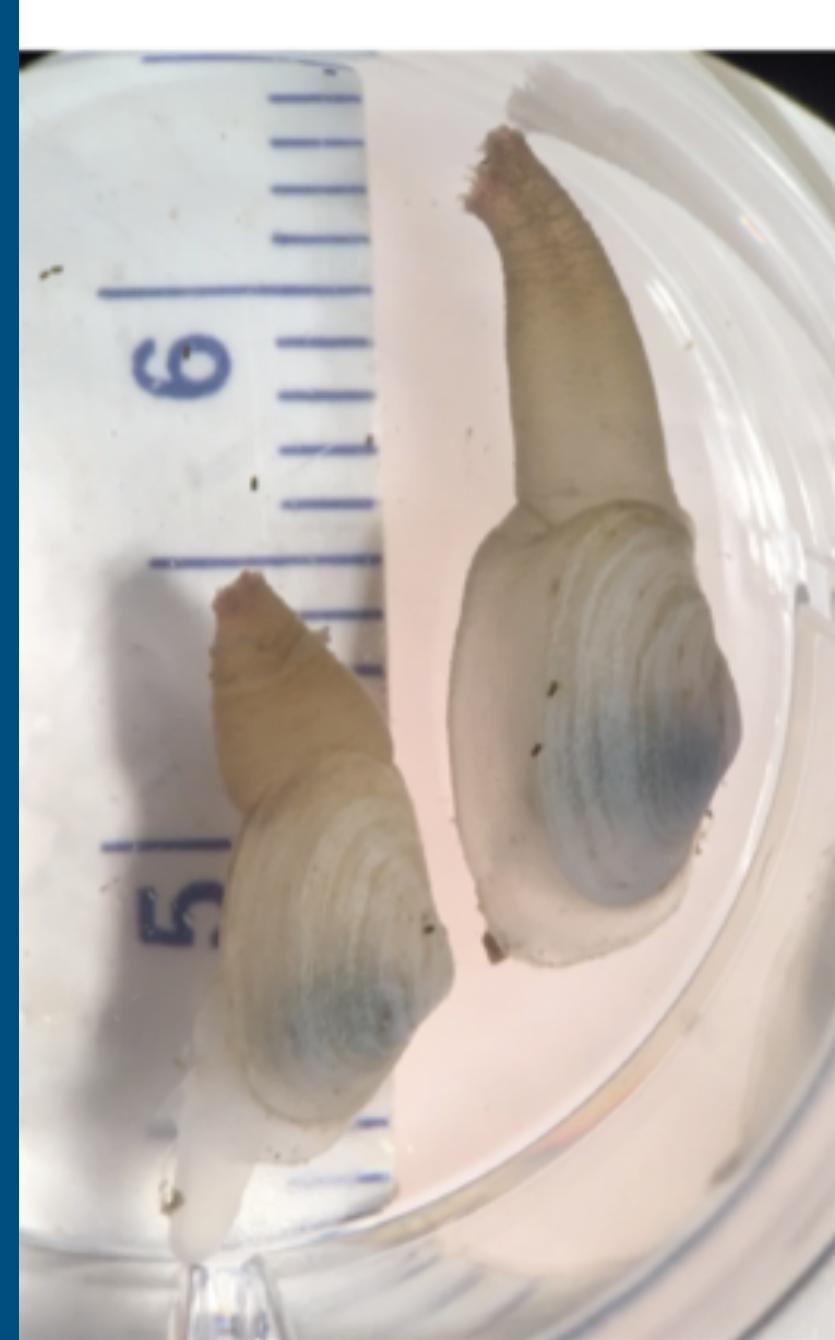
GEODUCKS AND OA

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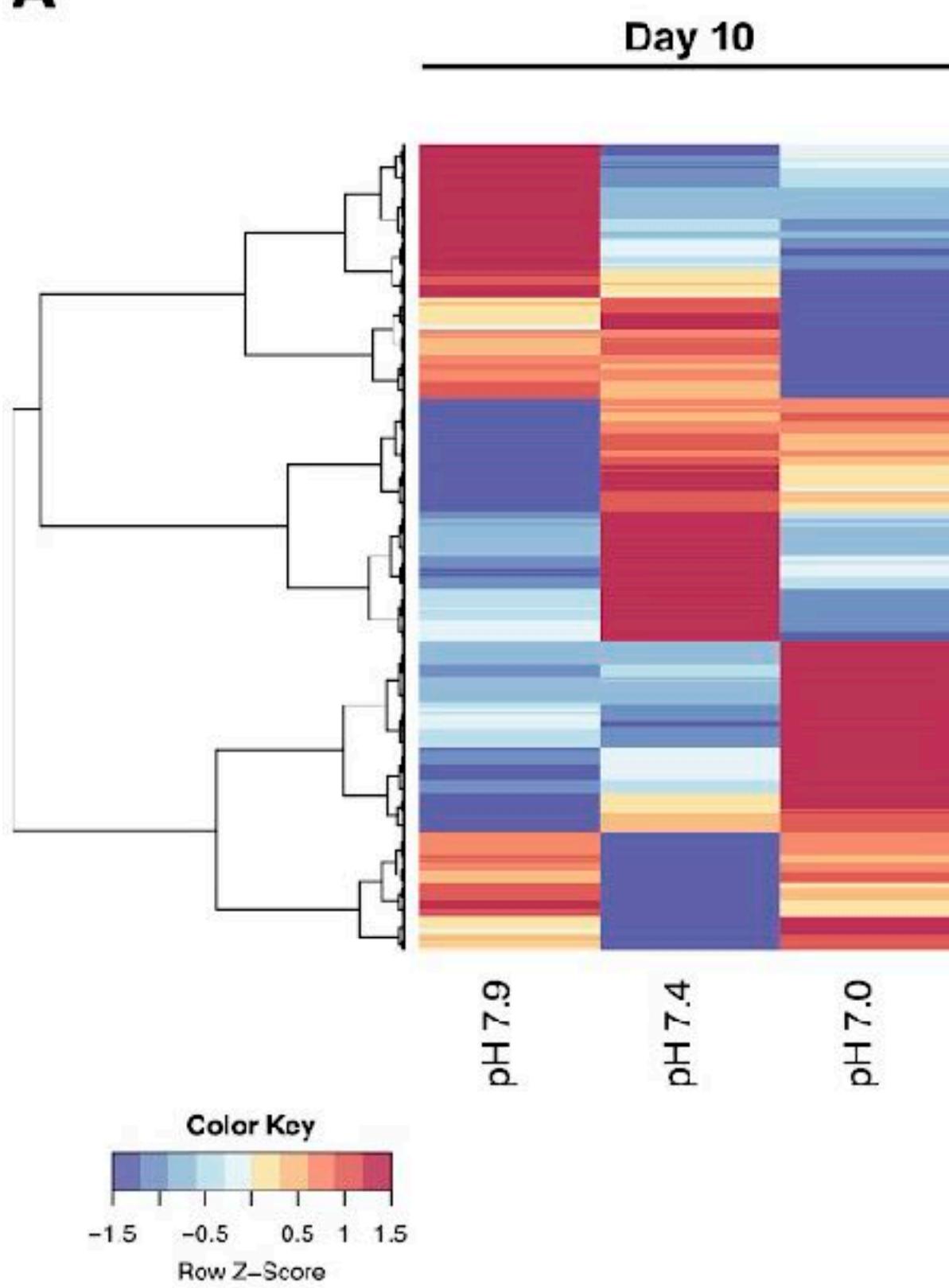
GEODUCKS AND OA

DNA METHYLATION

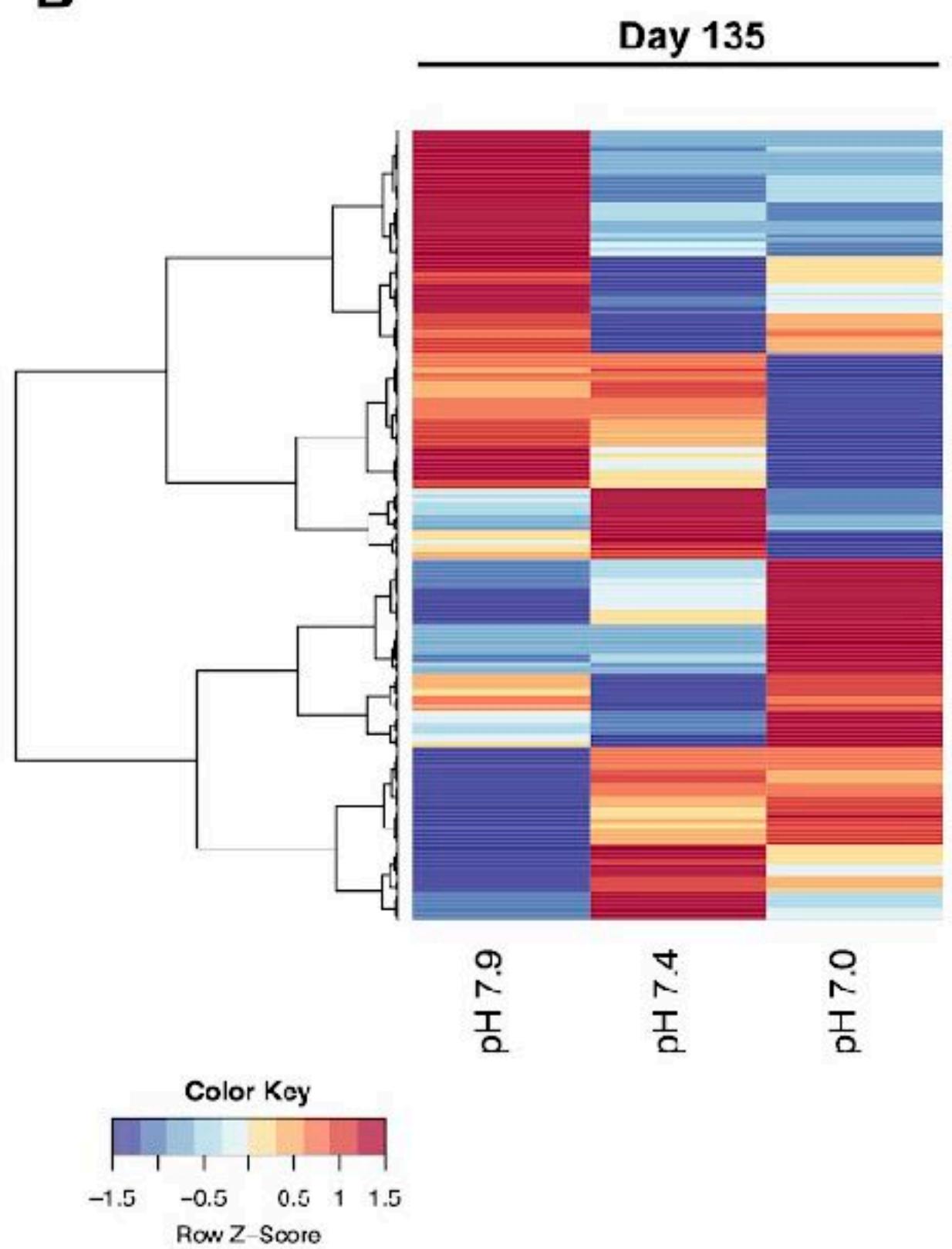


GEODUCKS AND OA

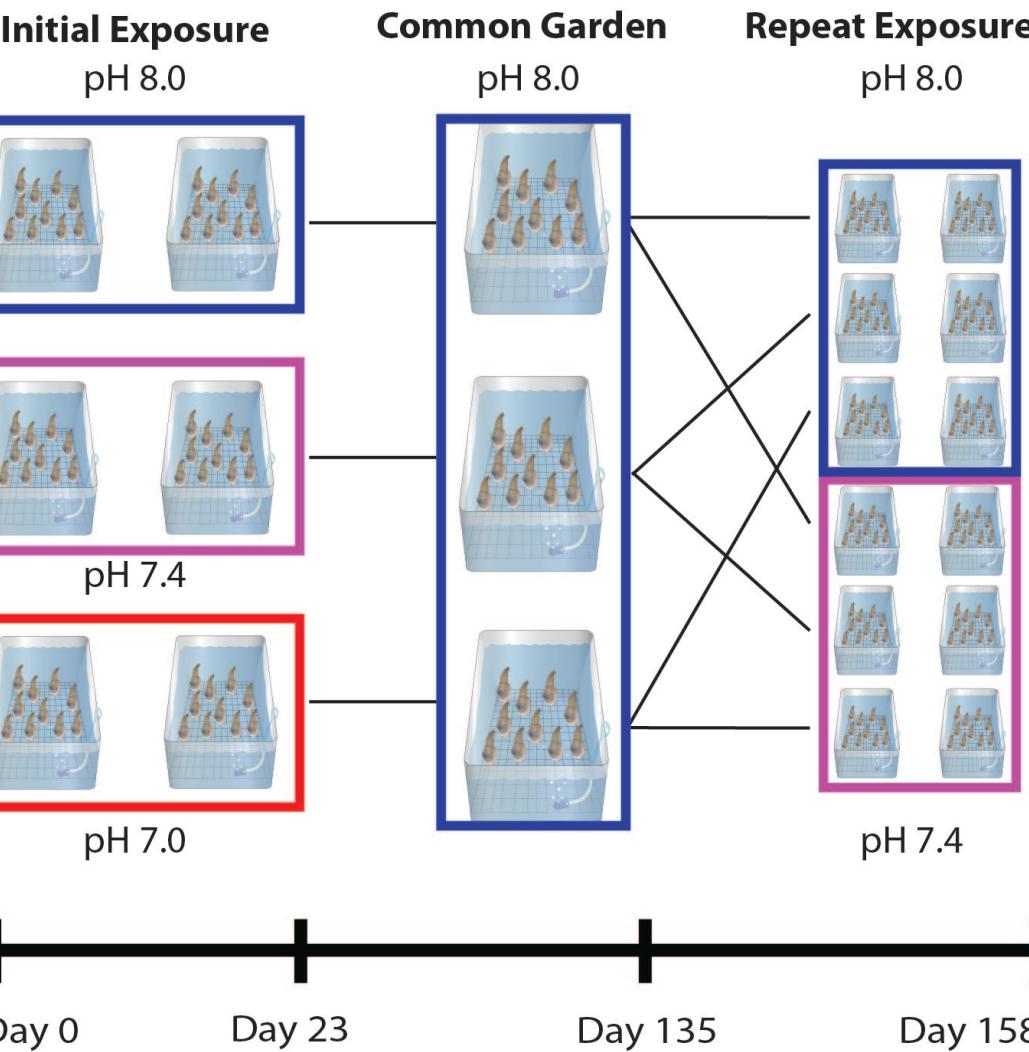
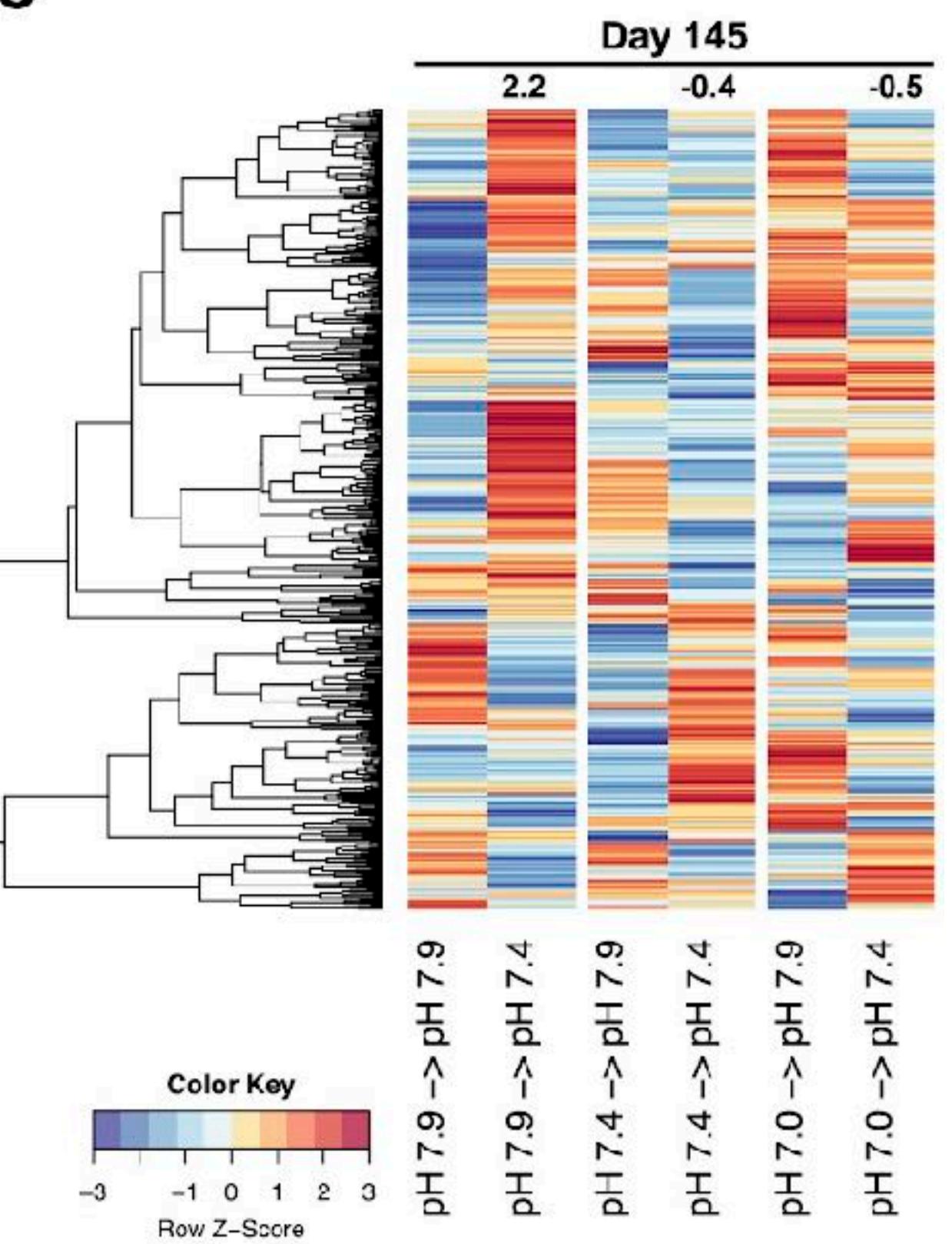
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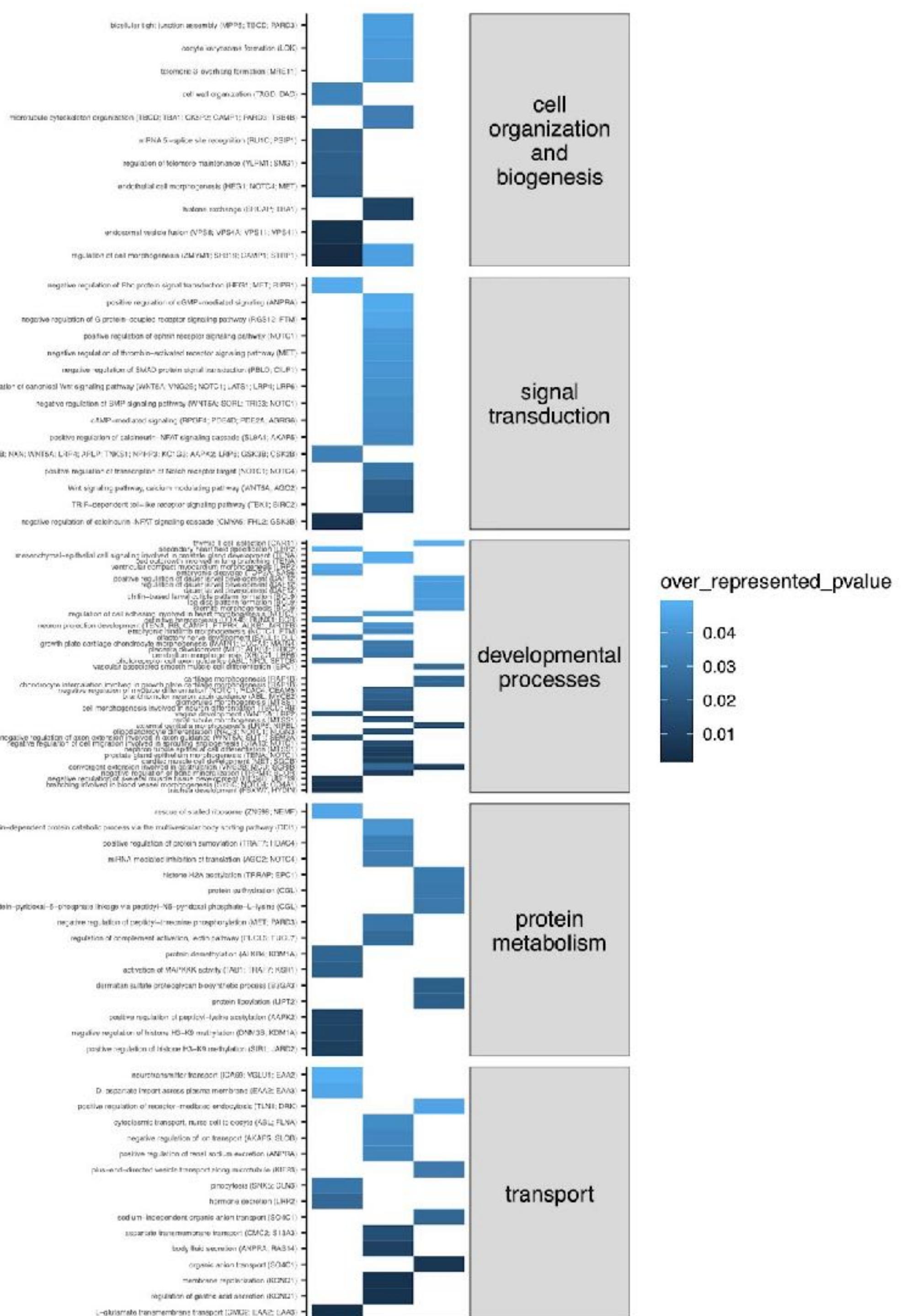
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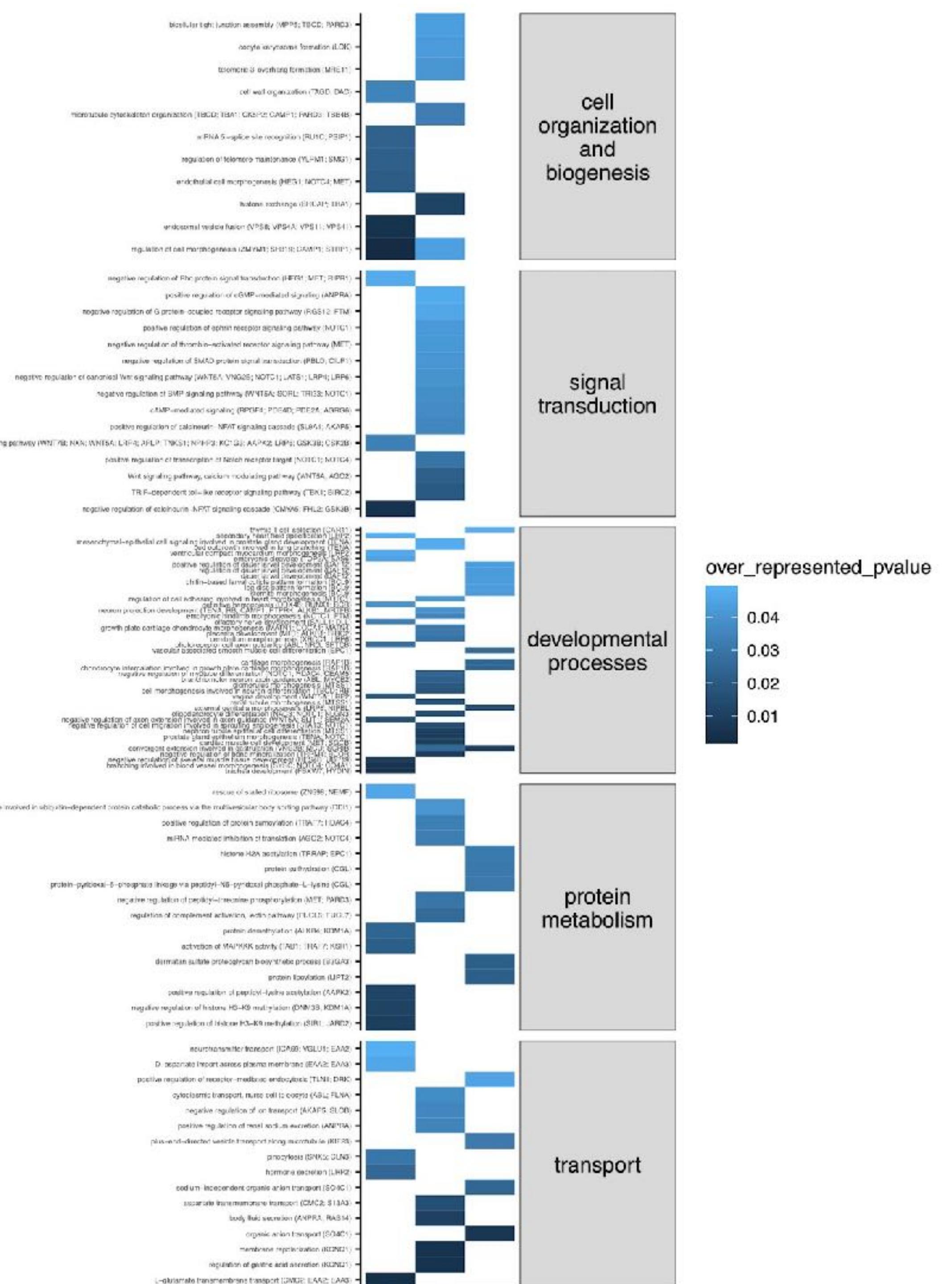
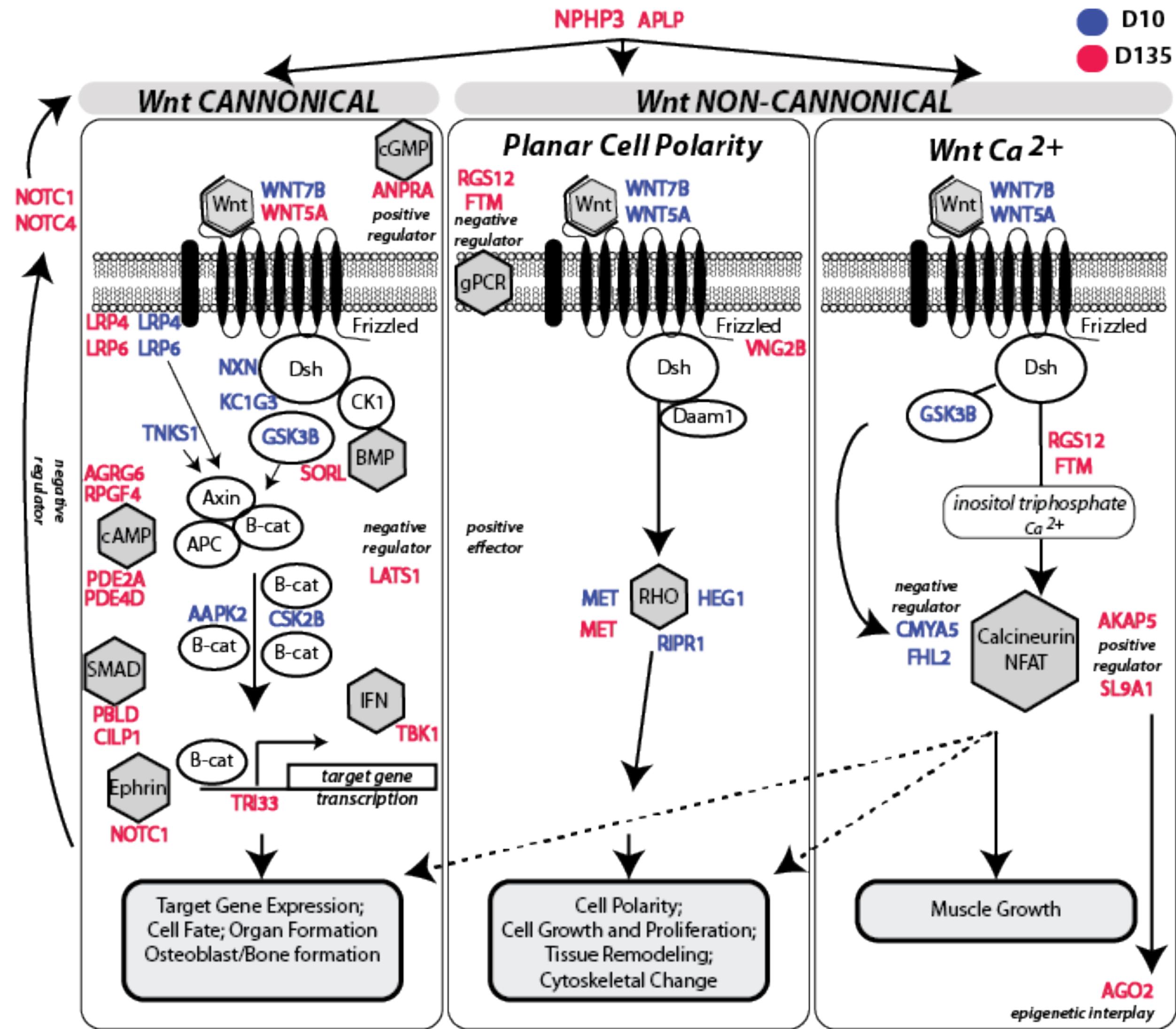
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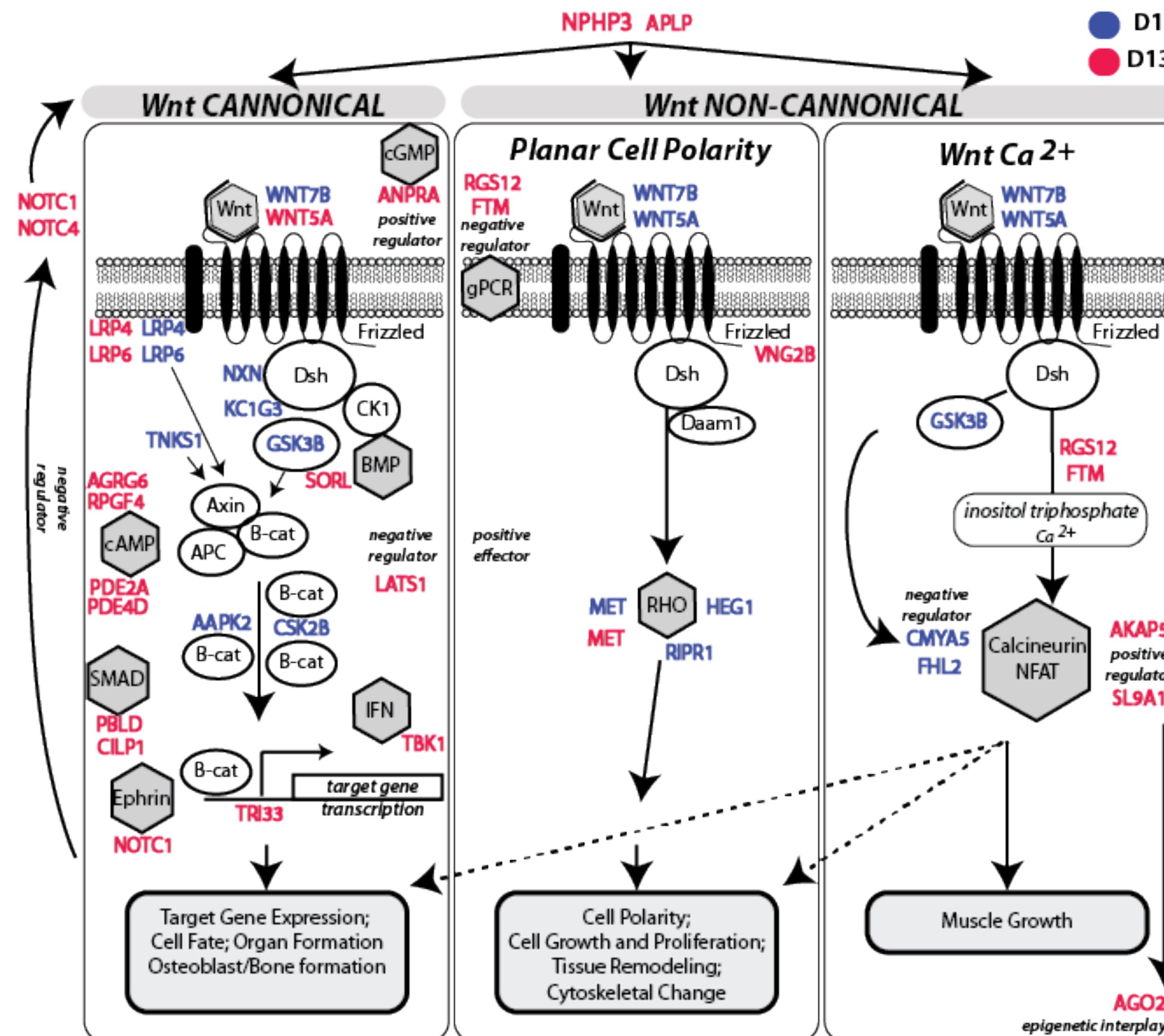
GEODUCKS AND OA



GEODUCKS AND OA



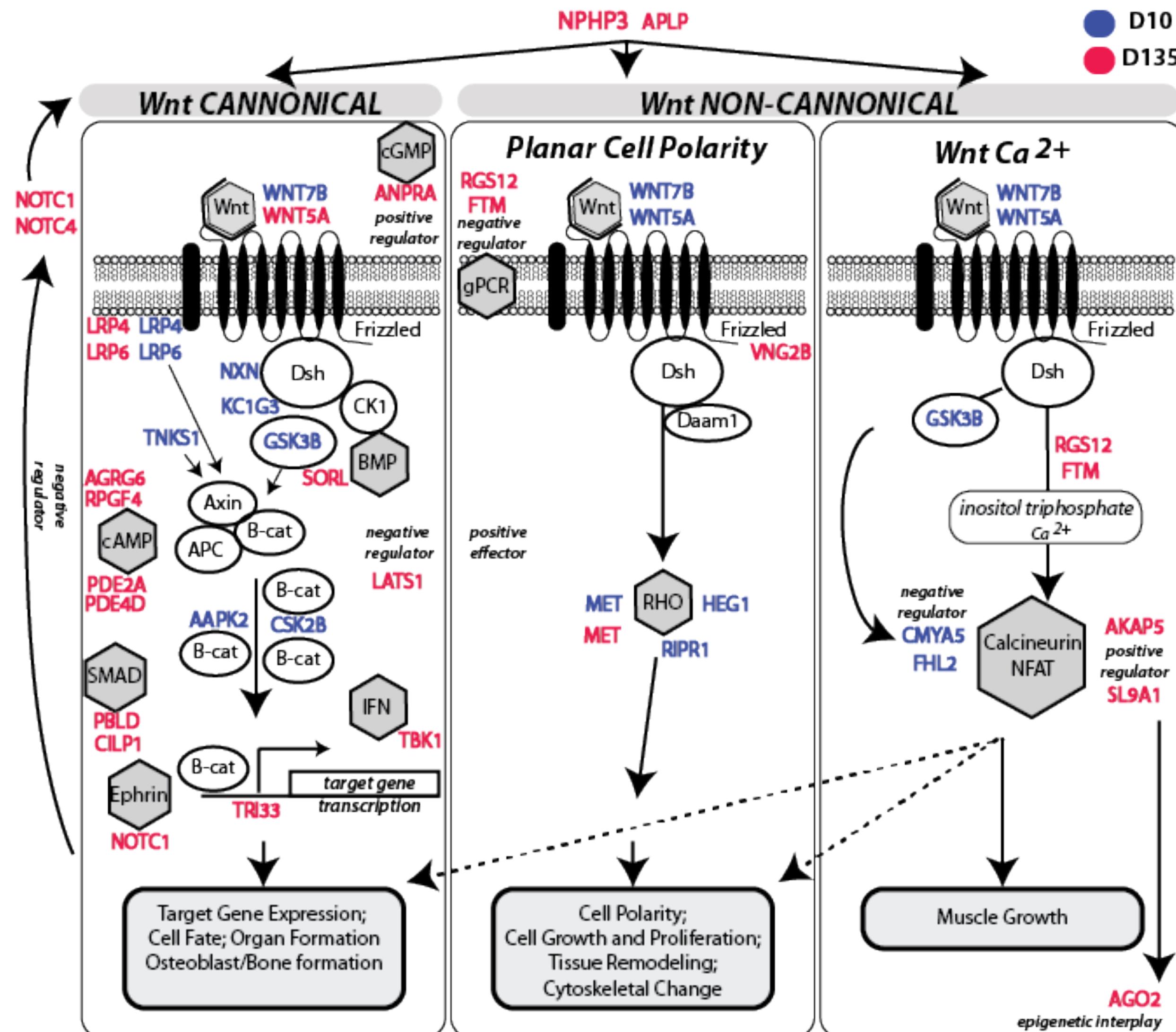
GEODUCKS AND OA



Hollie M. Putnam, Shelly A. Trigg, Samuel J. White, Laura H. Spencer, Brent Vadopalas, Aparna Natarajan, Jonathan Hetzel, Erich Jaeger, Jonathan Soohoo, Cristian Gallardo-Escárate, Frederick W. Goetz, Steven B. Roberts

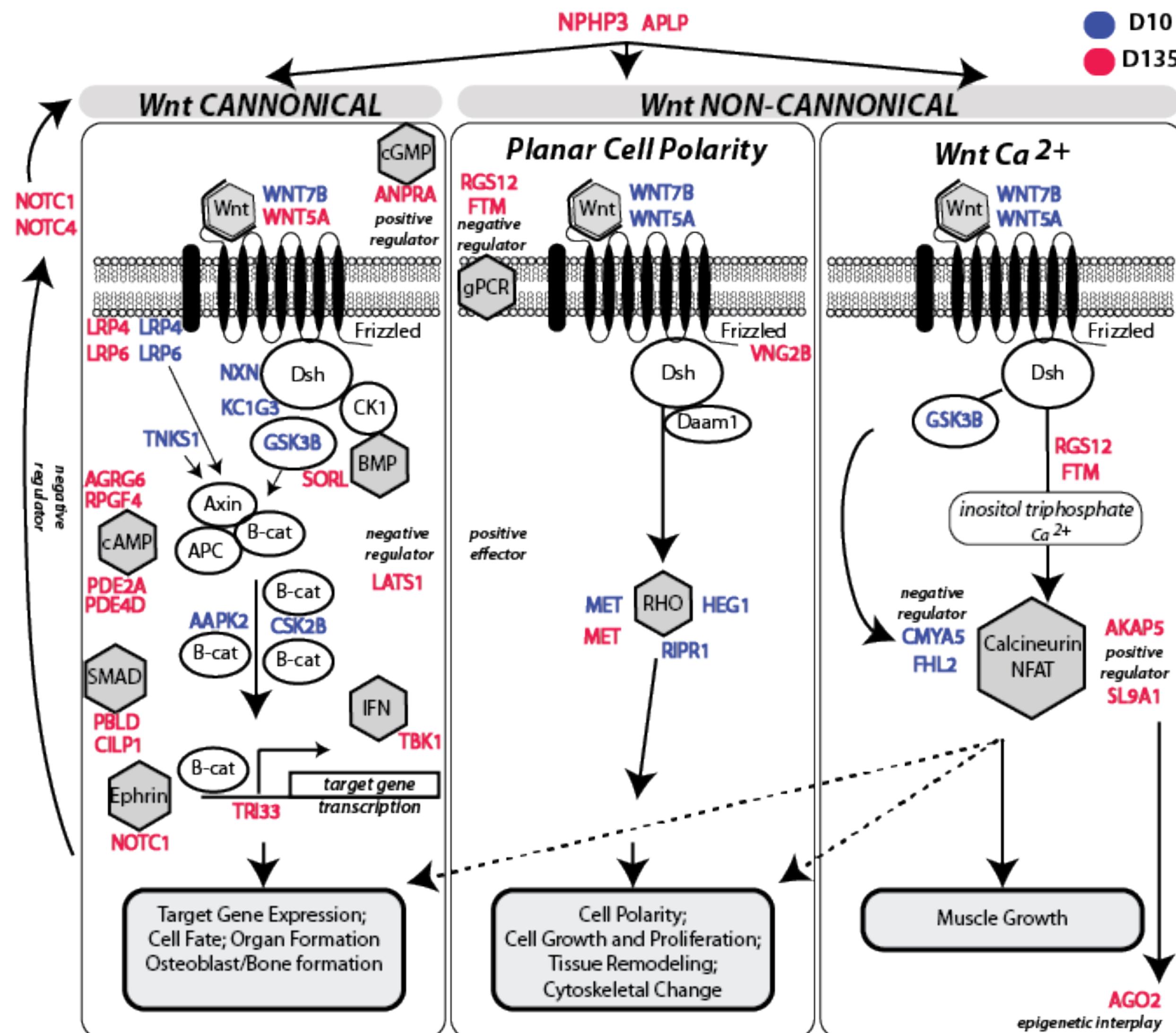
doi: <https://doi.org/10.1101/2022.06.24.497506>

GEODUCKS AND OA



Following four months of ambient common-garden conditions, **juveniles initially exposed to low pH compensatorily grew larger**, with DNA methylation indicative of these phenotypic differences, demonstrating epigenetic carryover effects persisted months after initial exposure.

GEODUCKS AND OA



Functional enrichment analysis of differentially methylated genes revealed regulation of signal transduction through widespread changes in the **Wnt signaling pathways that influence cell growth, proliferation, tissue and skeletal formation, and cytoskeletal change.**

Early-life Priming

Hardening

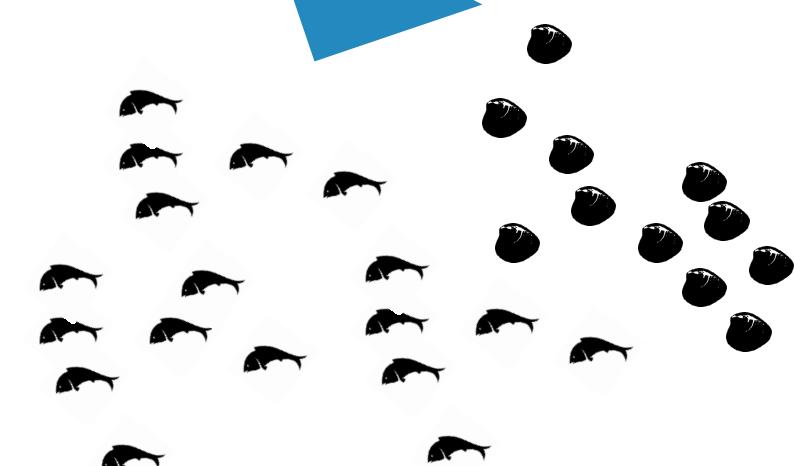


Adults

Influencing adult phenotype by altering early life environment



Influencing offspring phenotype by altering environmental conditions of parents



Larvae

Transgenerational Plasticity

Carry-over effects

Can we see an imprint in parents transmitted to the offspring?

Early-life Priming

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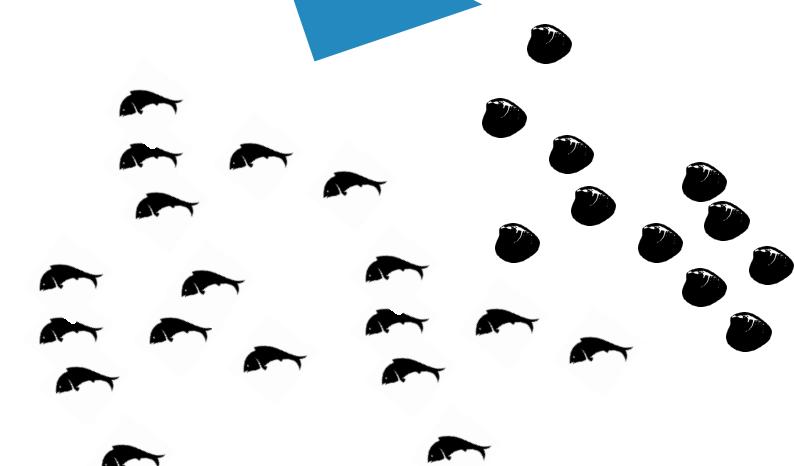


Adults

Influencing adult phenotype by altering early life environment



Influencing offspring phenotype by altering environmental conditions of parents



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Can we see an imprint in parents transmitted to the offspring?

NO



EXCITING?
COMPLEX
'LAYER' OF RESILIENCE



EXCITING? COMPLEX 'LAYER' OF RESILIENCE

Genetic Linkage

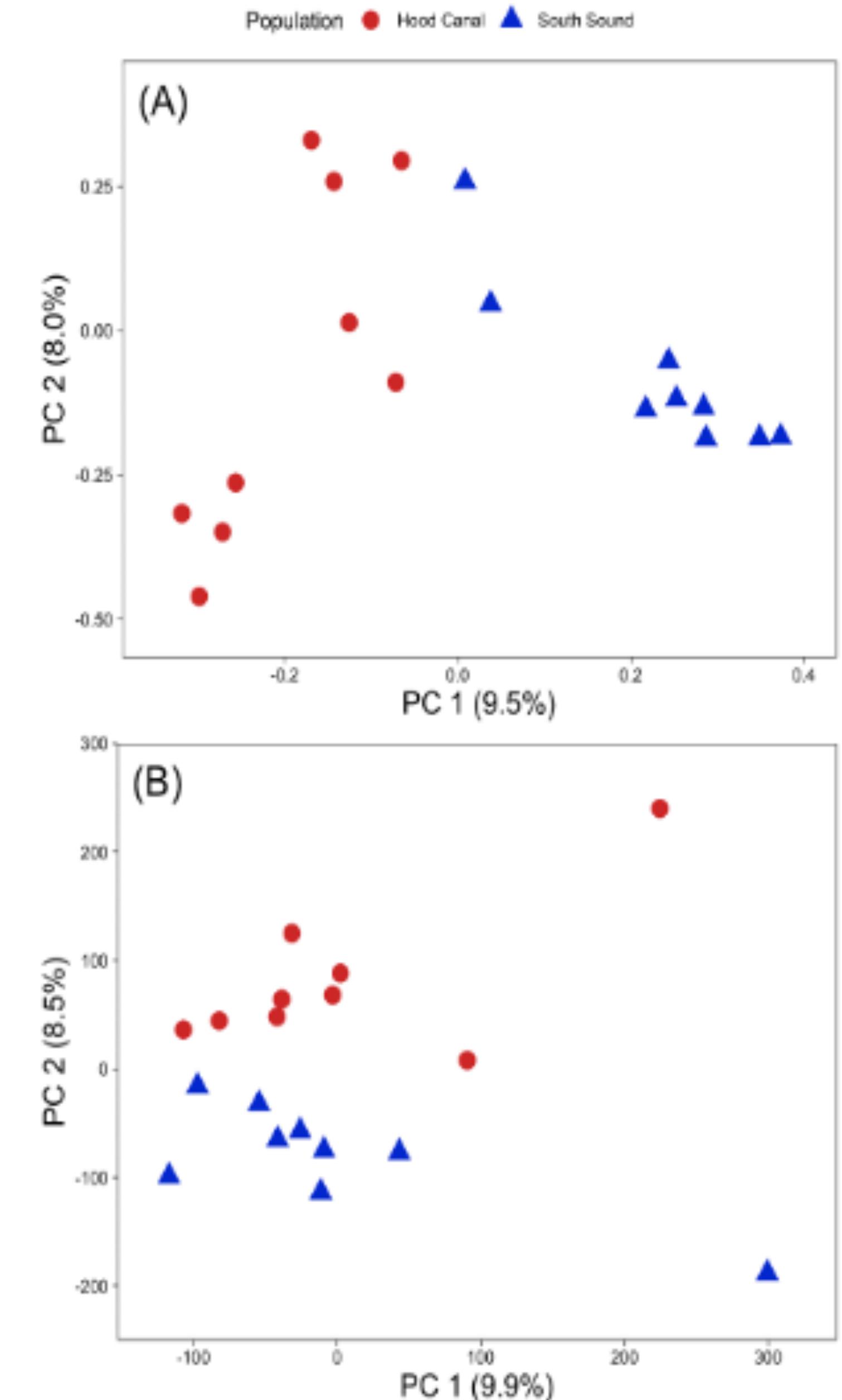
Epigenetic and Genetic Population Structure is Coupled in a Marine Invertebrate

Katherine Silliman ^{1,†}, Laura H. Spencer ^{2,†}, Samuel J. White², and Steven B. Roberts ^{2,*}

First characterization of genome-wide DNA methylation patterns in the oyster genus *Ostrea*

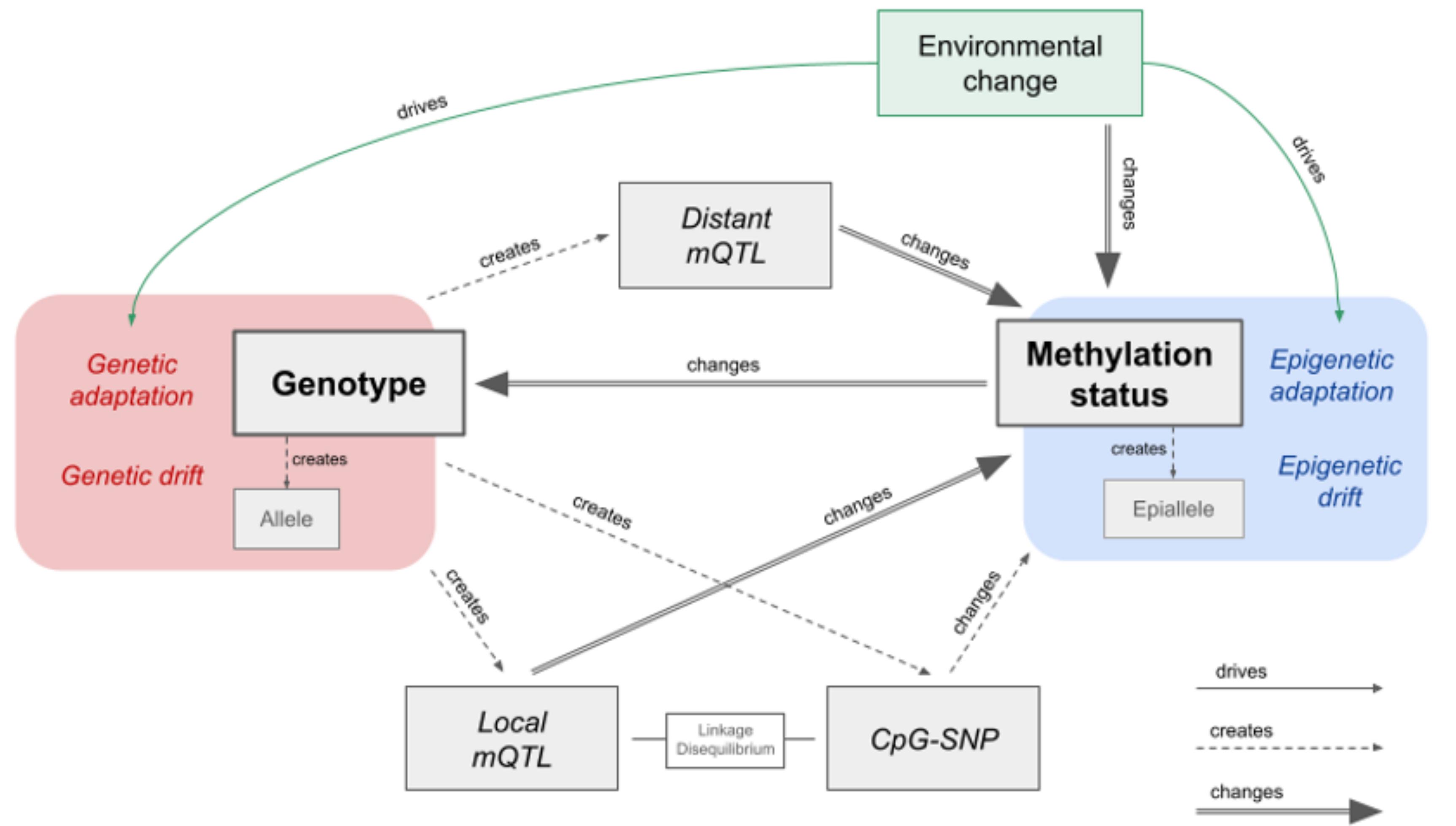
Identified 3,963 differentially methylated loci between populations. Clear coupling between genetic and epigenetic patterns of variation, **with 27% of variation in inter-individual methylation differences explained by genotype.**

Underlying this association are both direct genetic changes in CpGs (CpG-SNPs) and genetic variation with indirect influence on methylation (mQTLs).



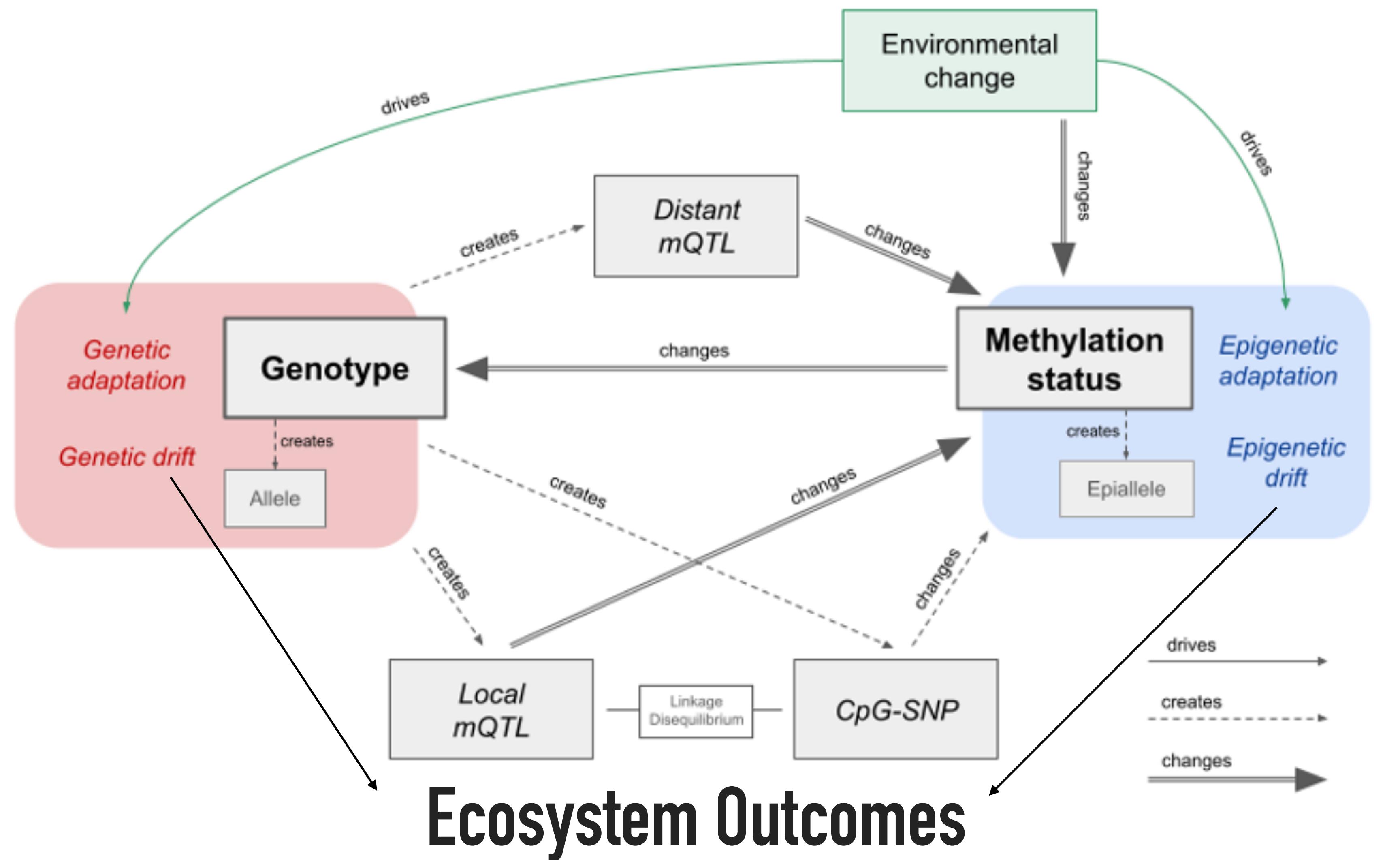
IMPLICATIONS

EPIGENETIC AND GENETIC POPULATION STRUCTURE



IMPLICATIONS

EPIGENETIC AND GENETIC POPULATION STRUCTURE



Open Science



Bioinformatic Approaches in non-model species

marineomics.io

Received: 29 March 2023 | Accepted: 11 September 2023

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APPLICATION

A dynamic web resource for robust and reproducible genomics in nonmodel species: marineomics.io

Samuel N. Bogan¹ | Jason Johns¹ | Joanna S. Griffiths² | Danielle Davenport³ |
Sara J. Smith^{4,5,6} | Sara M. Schaal⁷ | Alan Downey-Wall⁸ | Runyang Nicolas Lou^{9,10} |
Katie Lotterhos¹¹ | Megan E. Guidry¹² | Hanny E. Rivera¹³ | Joseph A. McGirr¹⁴ |
Jonathan B. Puritz¹² | Steven B. Roberts¹⁵ | Katherine Silliman^{16,17}

4 | Methods in Ecology and Evolution Open Access BRITISH
ECOLOGICAL
SOCIETY

MarineOmics

Best Principles Contributions ▾ Population Genomics ▾ Functional Genomics ▾ Genome-Phenome ▾ Panel Seminars Discussion Forum

Topics covered within RADSeq tutorial

RADseq

Katherine Silliman, Danielle Davenport

Description of tutorial steps

Setup for running code

Reduced Representation Sequencing (RADseq/GBS)

Considerations During Lab Work

Principles for Analyzing Your Data

Steps for a robust RAD analysis

First, look at the raw data!

Run an assembly pipeline

Evaluate potential sources of error

"Bad" samples

The power of PCA

Batch effects

Cryptic species/contamination/clones

Test a range of key parameters

Clustering threshold

Mapping parameters

If you would like to run the R code examples that are scattered throughout the guide (recommended but not required!), you will need to install some R packages. Only need to run this code once:

```
install.packages("tidyverse")
if (!requireNamespace("BiocManager", quietly = TRUE))
  install.packages("BiocManager")

BiocManager::install("SeqArray")
BiocManager::install("SNPRelate")
```

Example code to run using real data

Now load those packages, if using:

```
library(SeqArray) # efficient storage and filtering of genomic data
```

```
## Loading required package: gdsfmt
```

ACKNOWLEDGEMENTS

- ▶ Mackenzie Gavery (NOAA), Sam White (UW), Brent Vadopalas (UW), Shelly Wanamaker (GMGI), Sam Gurr (NOAA), Hollie Putnam (URI), Laura Spencer (UW), Katherine Silliman (NOAA), Yaamini Venkataraman (WHOI), Katie Lotterhos (NEU)

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