

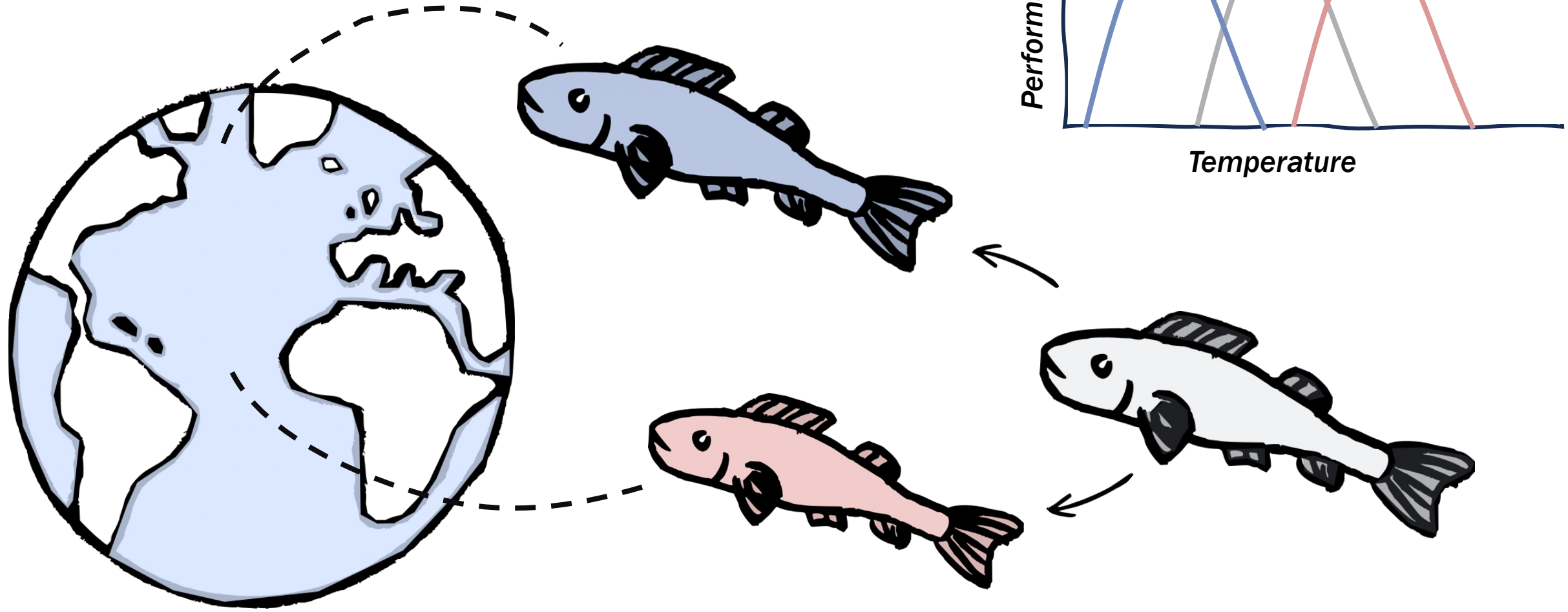
LOCAL ADAPTATION TO SALINITY STRESS: PHYSIOLOGICAL AND TRANSCRIPTOMIC INSIGHTS

Alexandra Hahn

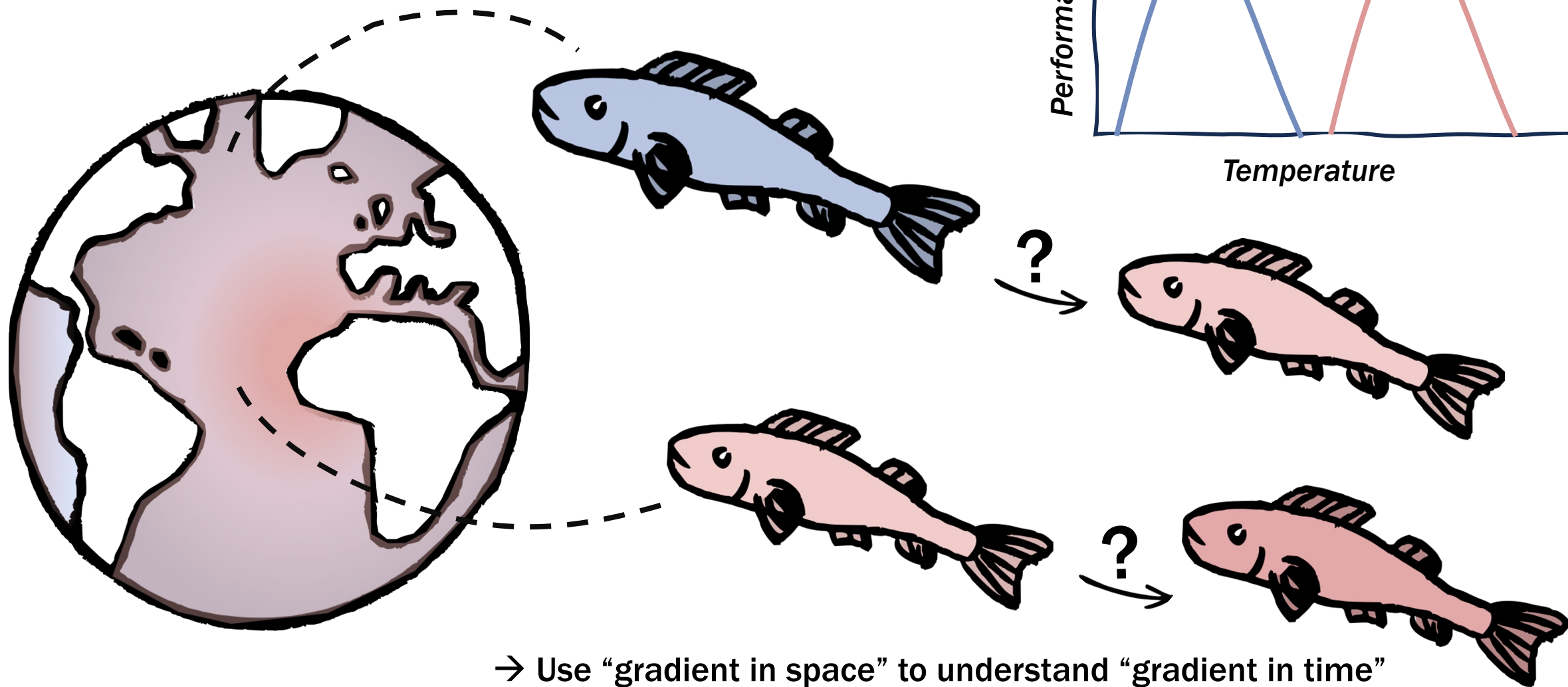
EMB Science Webinar - 20th March 2025



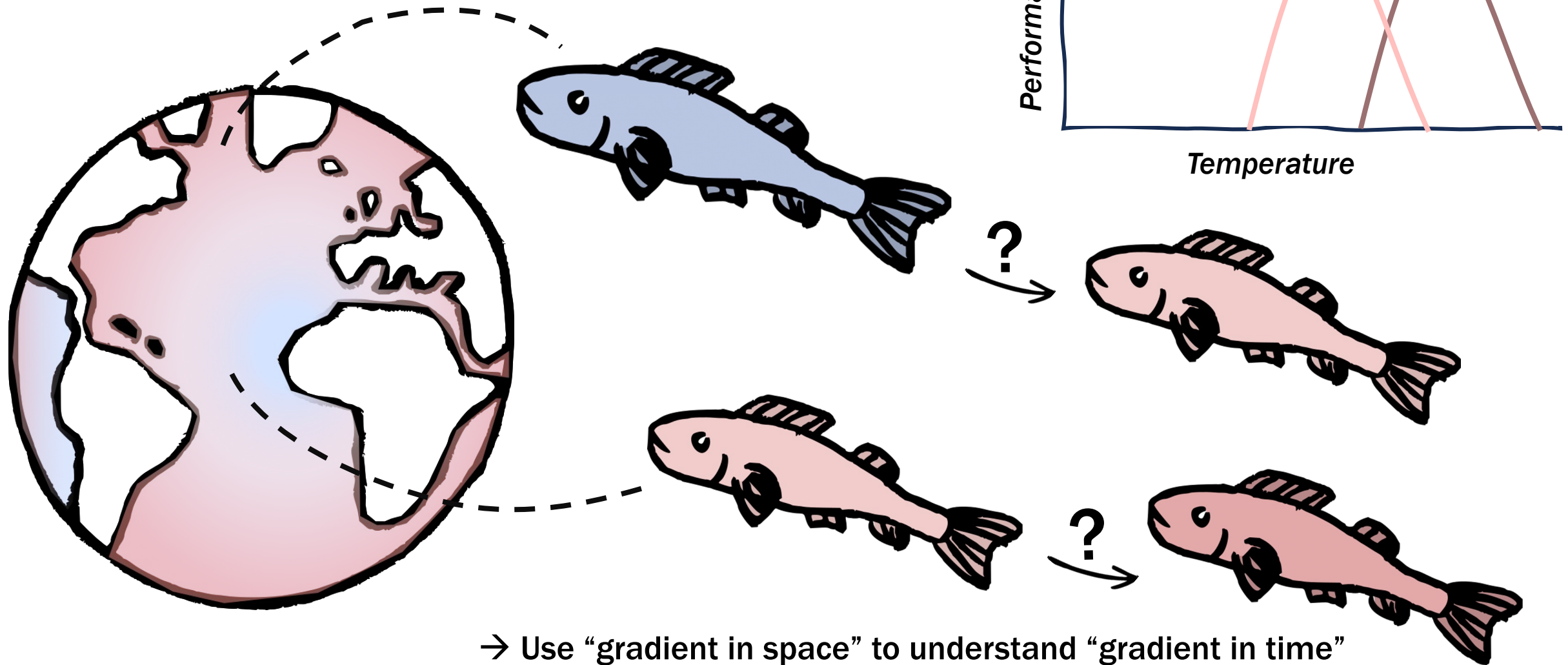
WHAT IS LOCAL ADAPTATION?



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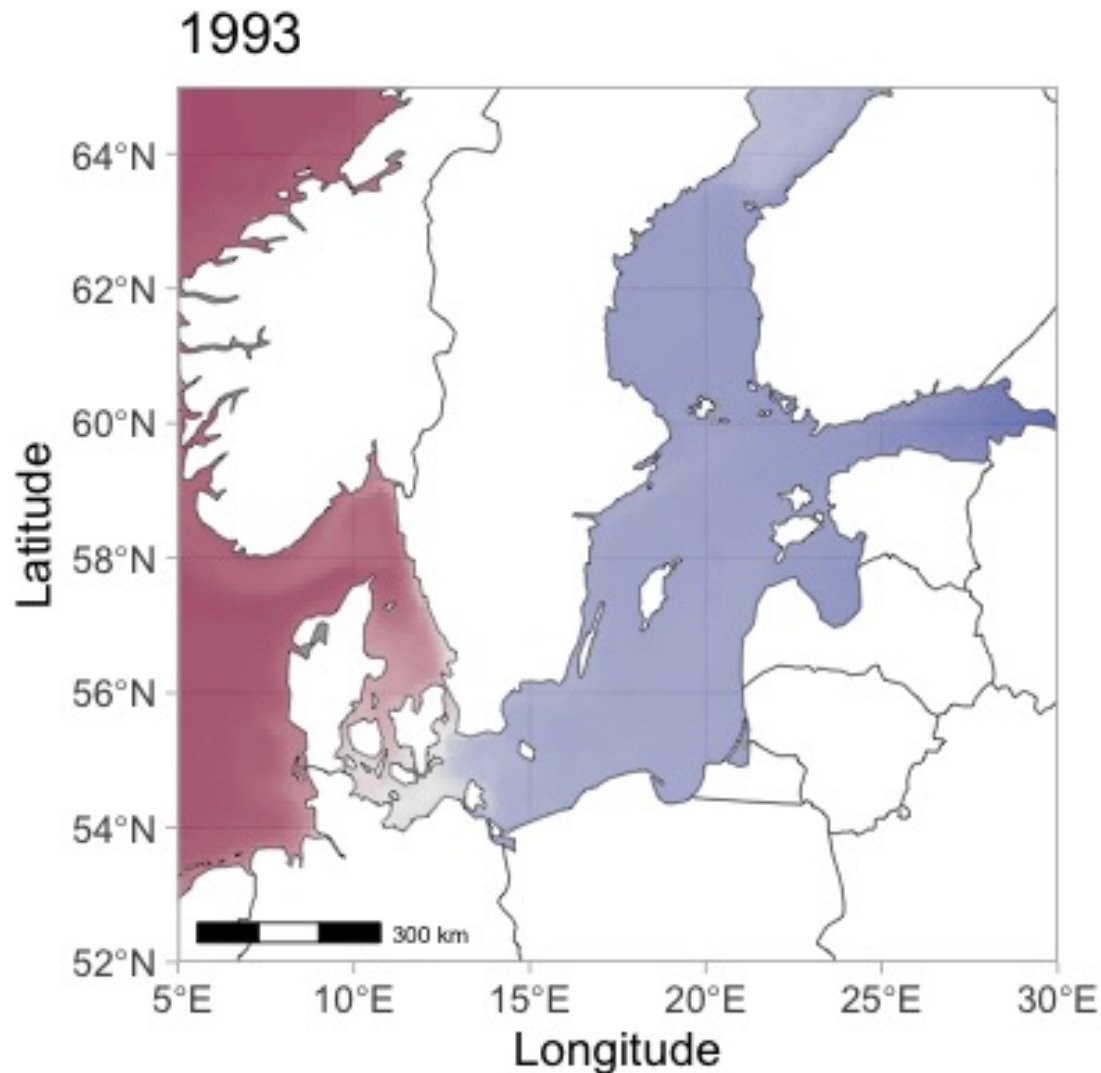


WHAT IS LOCAL ADAPTATION?

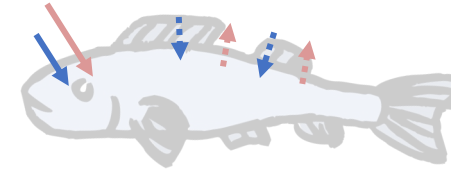


→ Use “gradient in space” to understand “gradient in time”

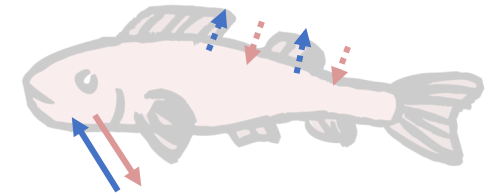
THE BALTIC SEA AND ADAPTATION



FRESHWATER



MARINE



→ ion flux → water flux → passive ———→ active

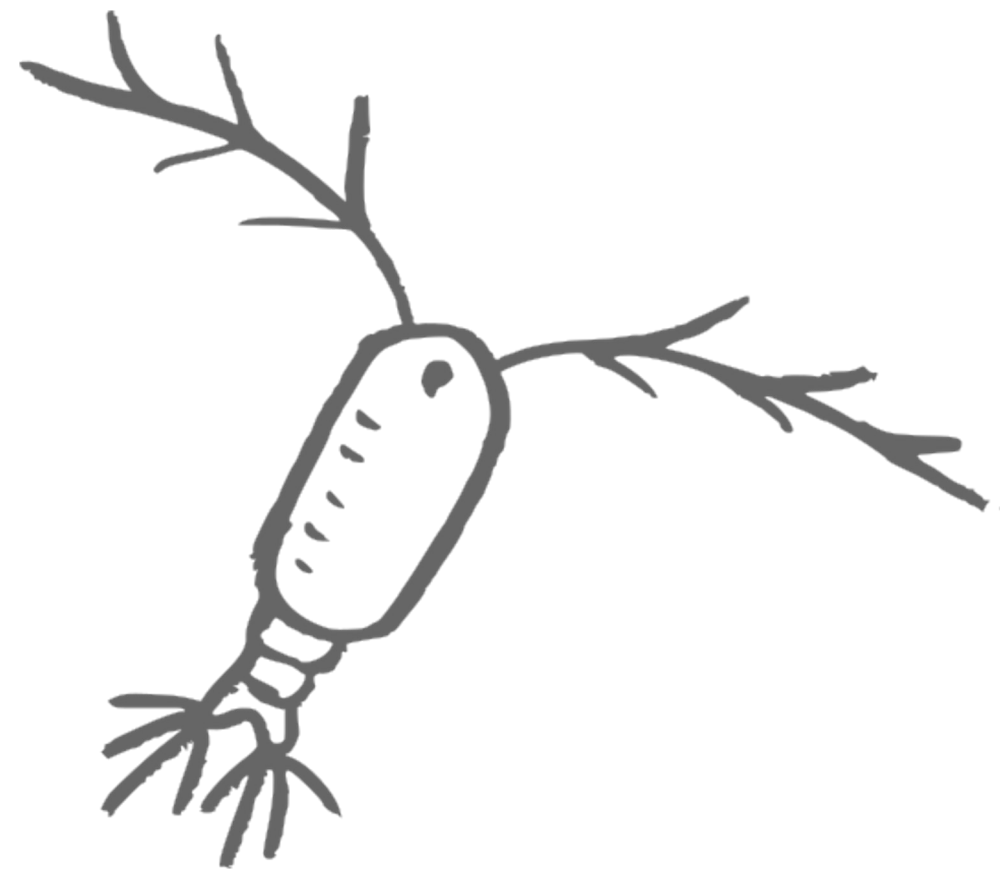
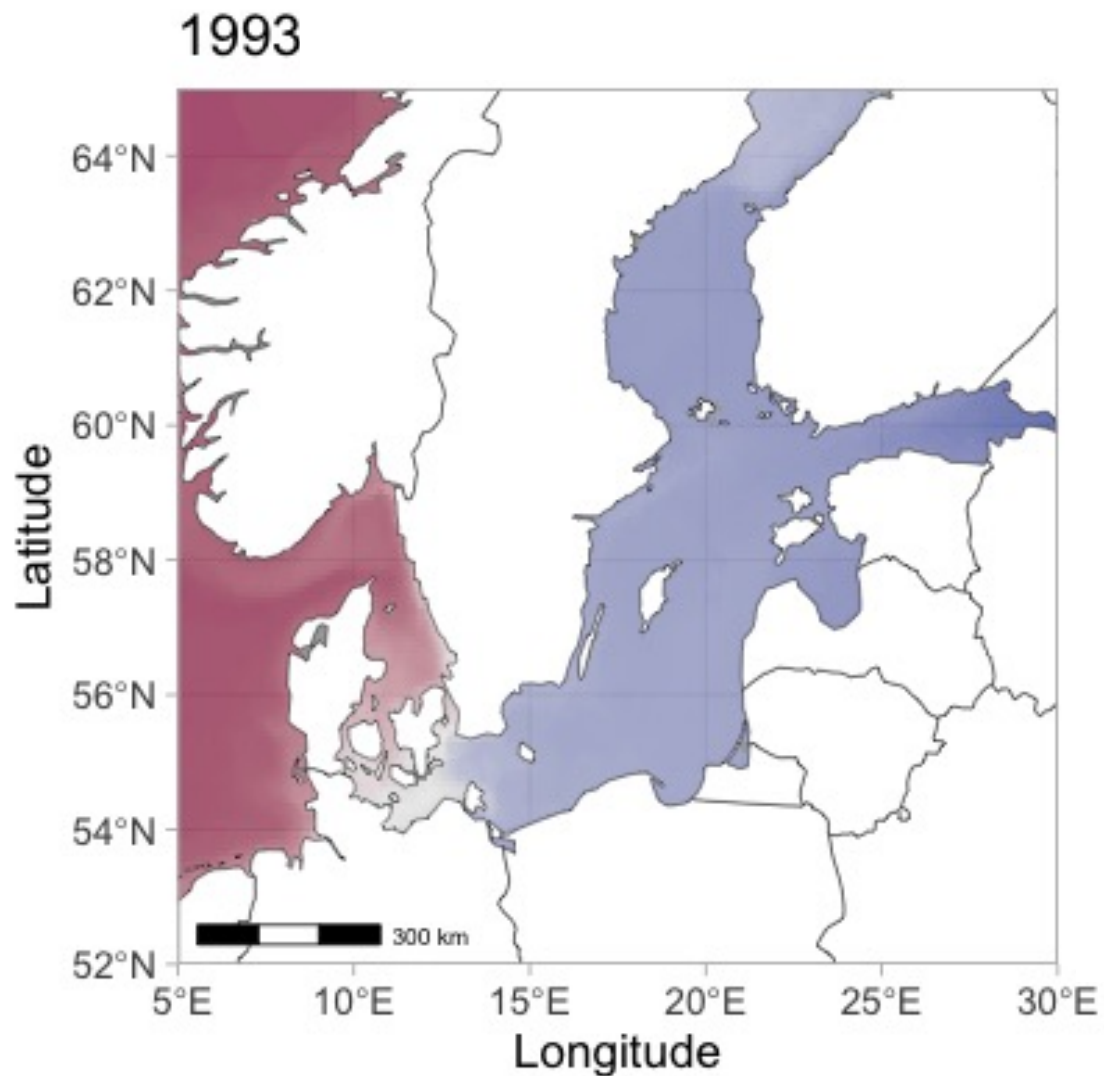
Salinity gradient forces organisms to compensate or adapt

→ Local adaptation shown in Baltic mussels, fish, and phytoplankton

→ Understanding adaptive capacity and population structure helps to predict how climate change might affect organisms

Study system: Coastal copepod *Acartia tonsa*

THE BALTIC SEA AND ADAPTATION



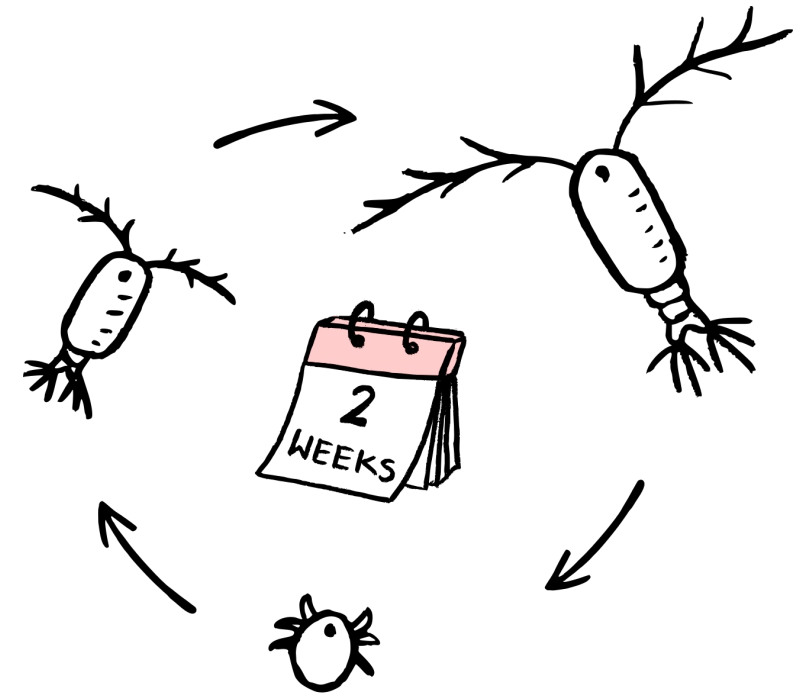
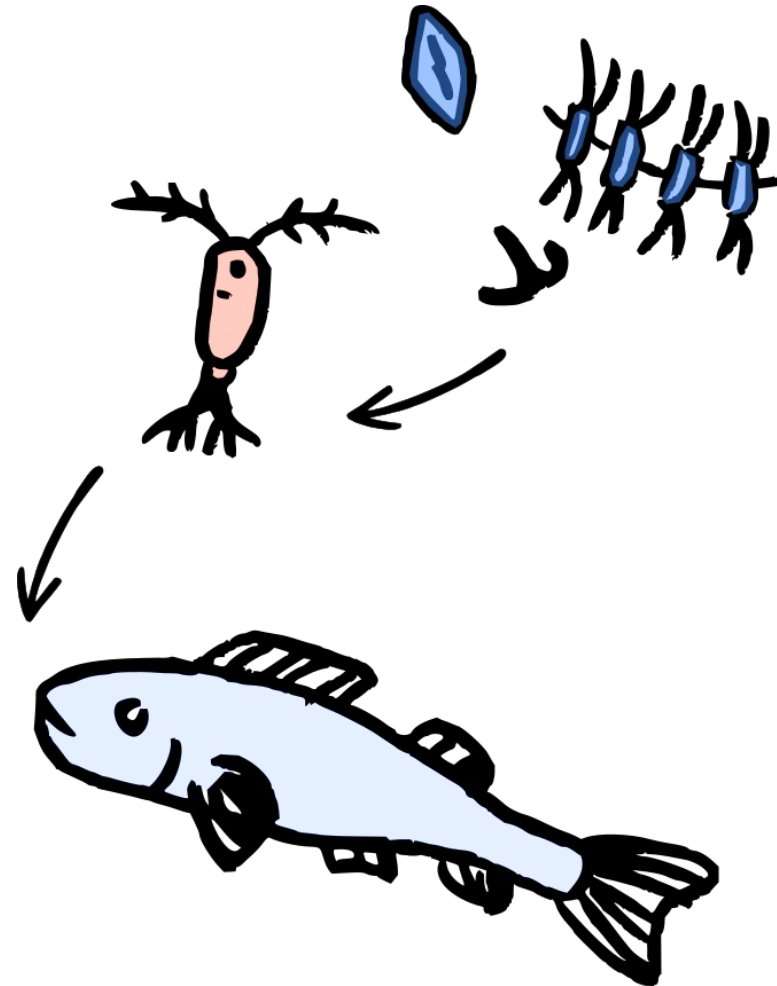
Study system: Coastal copepod *Acartia tonsa*

WHY COPEPODS?

ECOLOGICAL IMPORTANCE

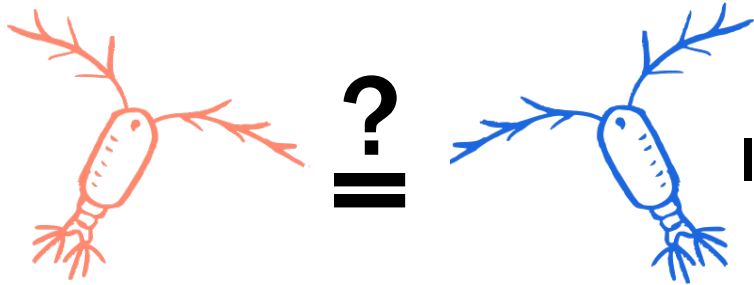


GLOBAL DISTRIBUTION

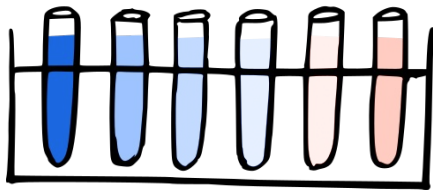


SHORT LIFECYCLE

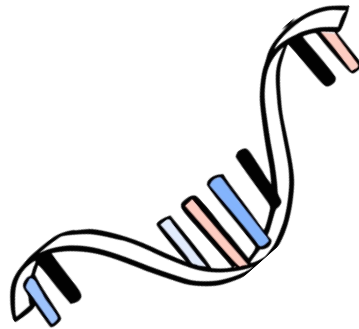
STUDY QUESTION



Is *Acartia tonsa* locally adapted to salinity in the Baltic Sea?



FITNESS EXPERIMENTS

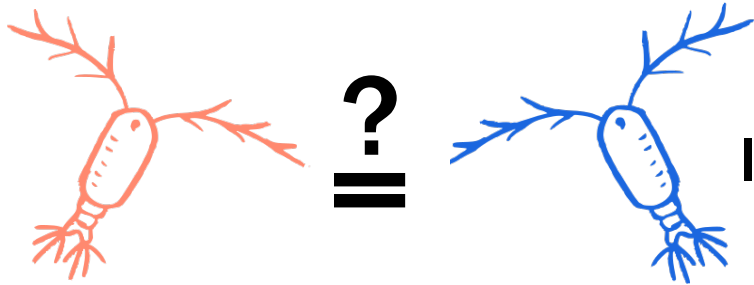


GENE EXPRESSION

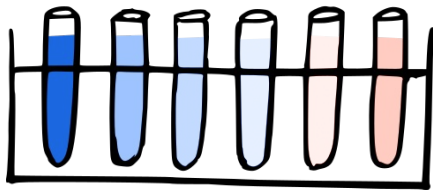


POPULATION GENOMICS

STUDY QUESTION



Is *Acartia tonsa* locally adapted to salinity in the Baltic Sea?



FITNESS EXPERIMENTS



GENE EXPRESSION

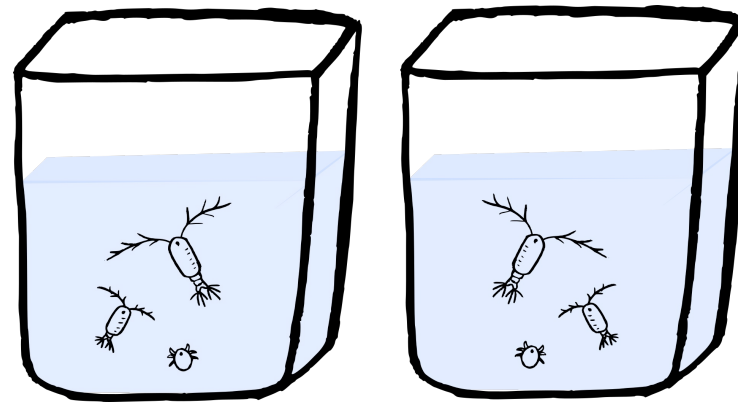
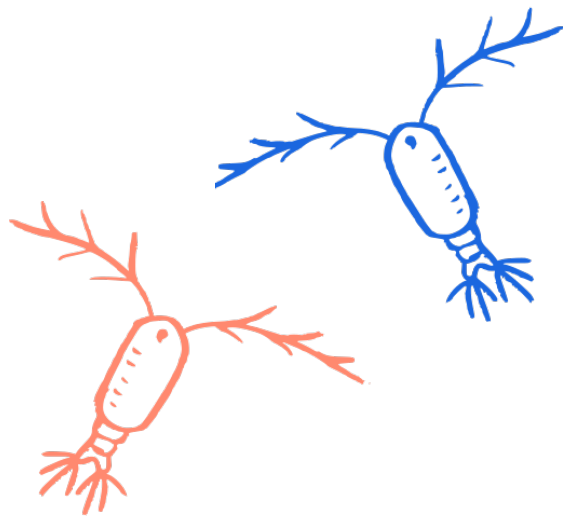
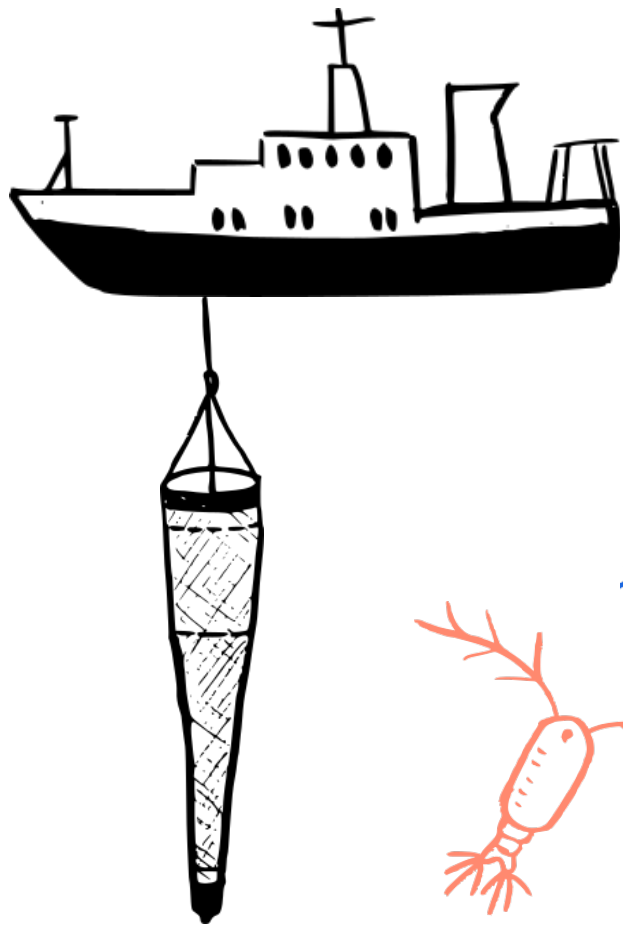


POPULATION GENOMICS

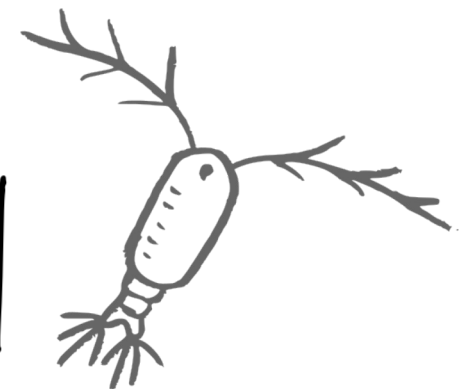
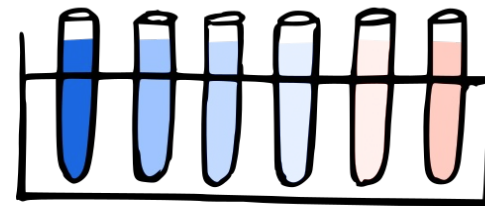
PHYSIOLOGY

Is there a difference in fitness at low salinity?

EXPERIMENTAL APPROACH

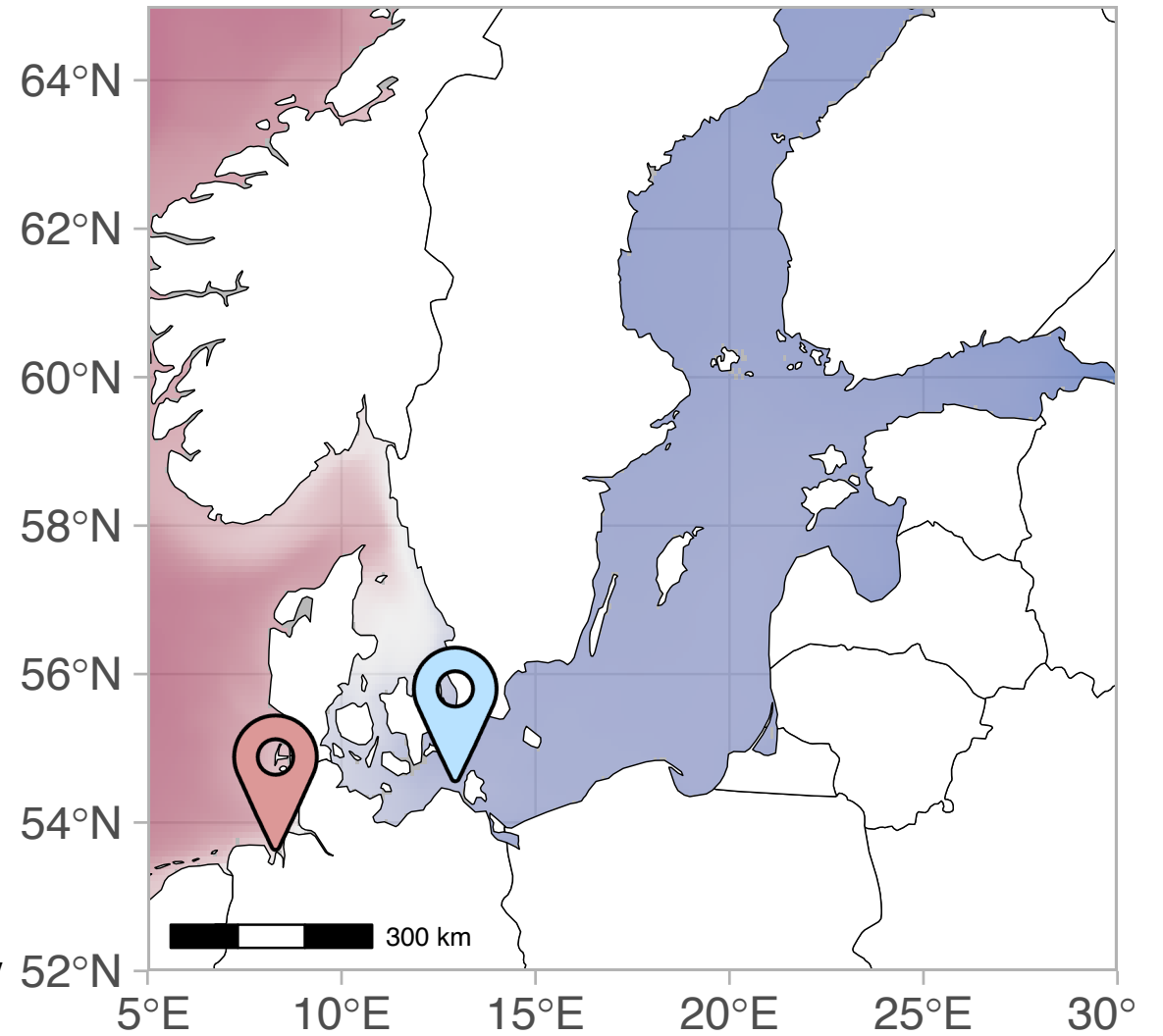
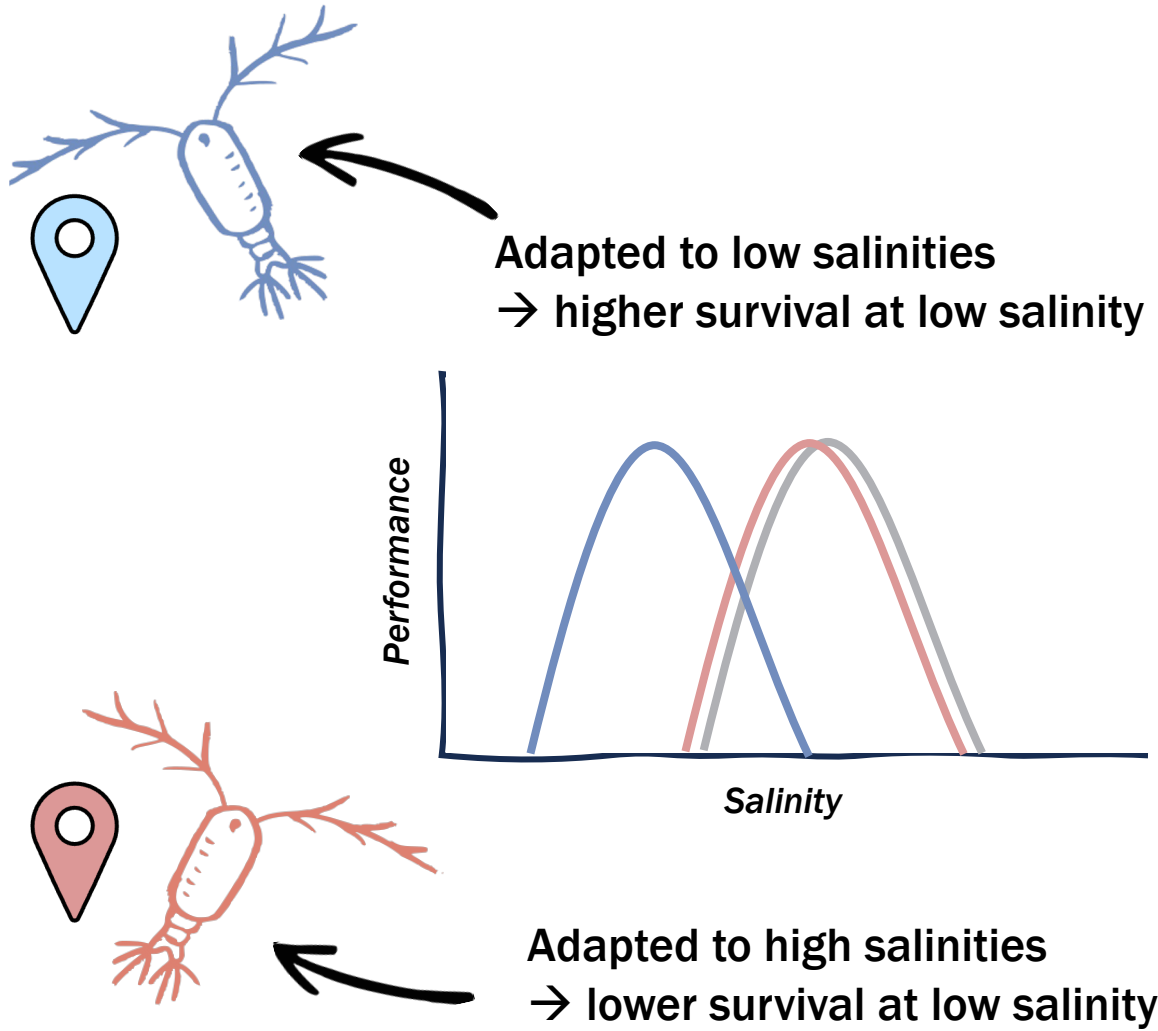


Expose to low salinity

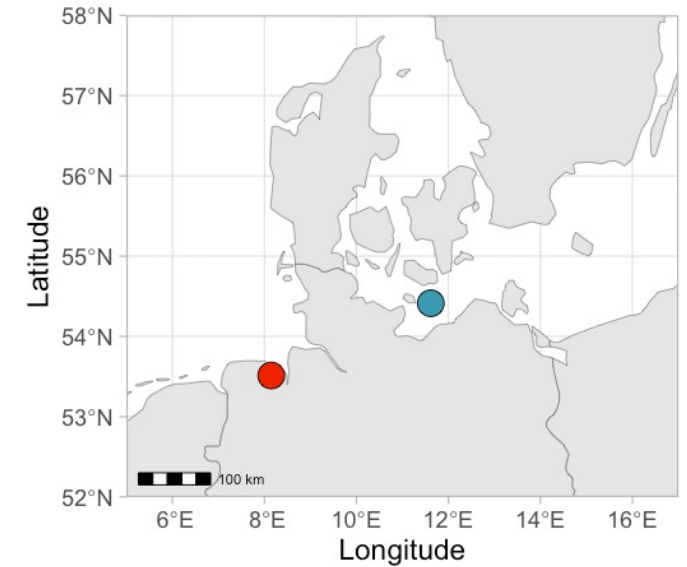
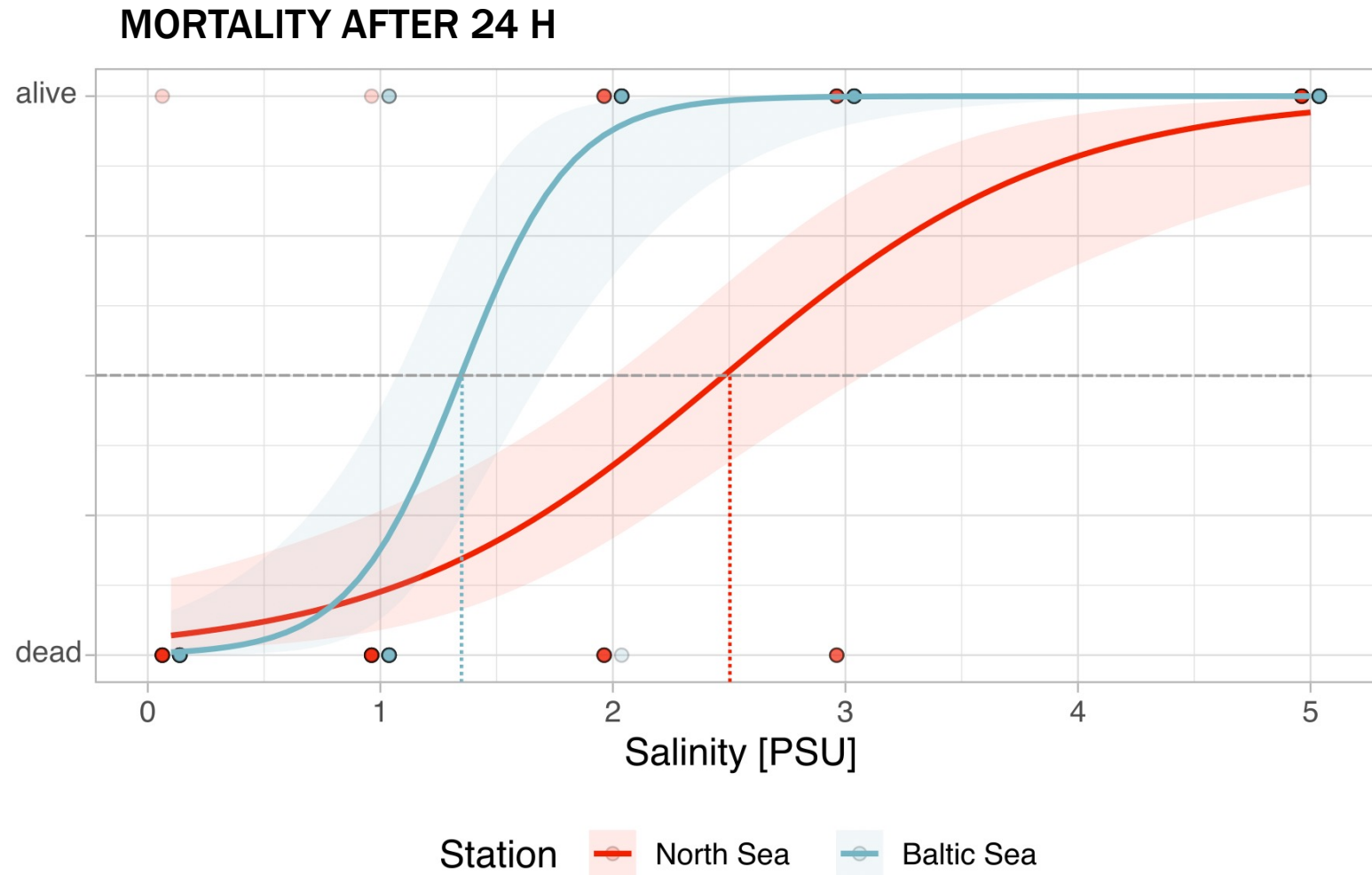


How many copepods survive?

EXPECTATIONS



HIGHER SURVIVAL IN BALTIC COPEPODS

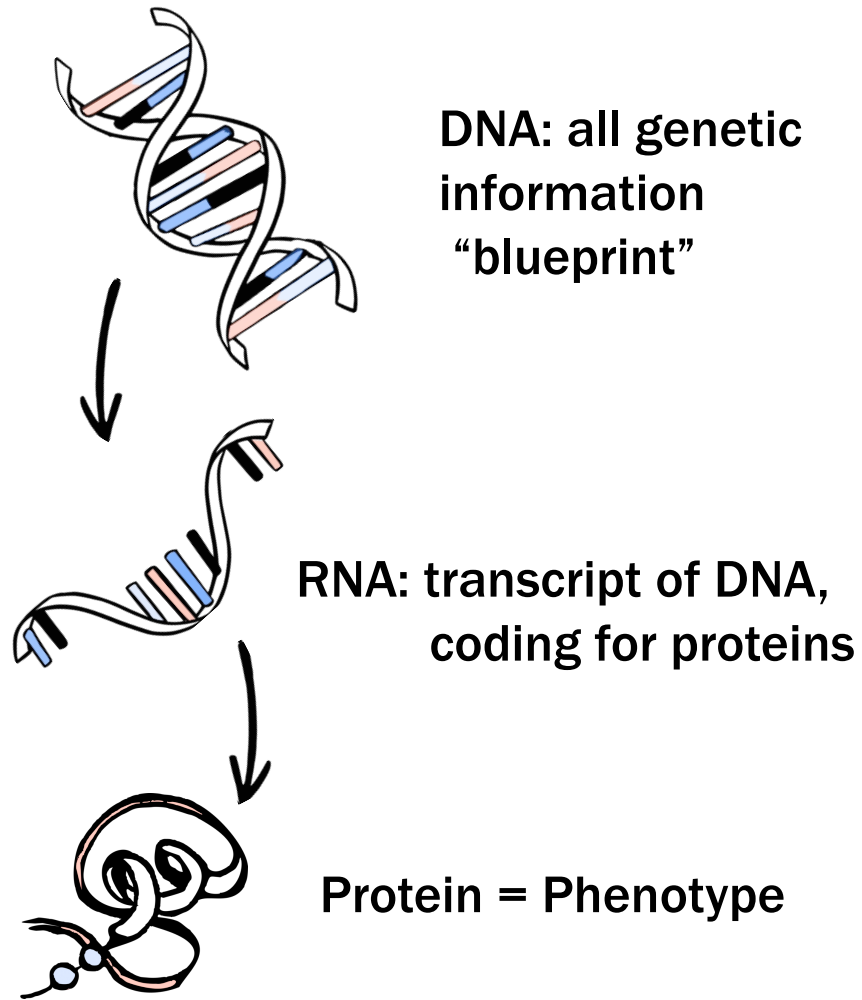


→ What are the mechanisms behind salinity tolerance?

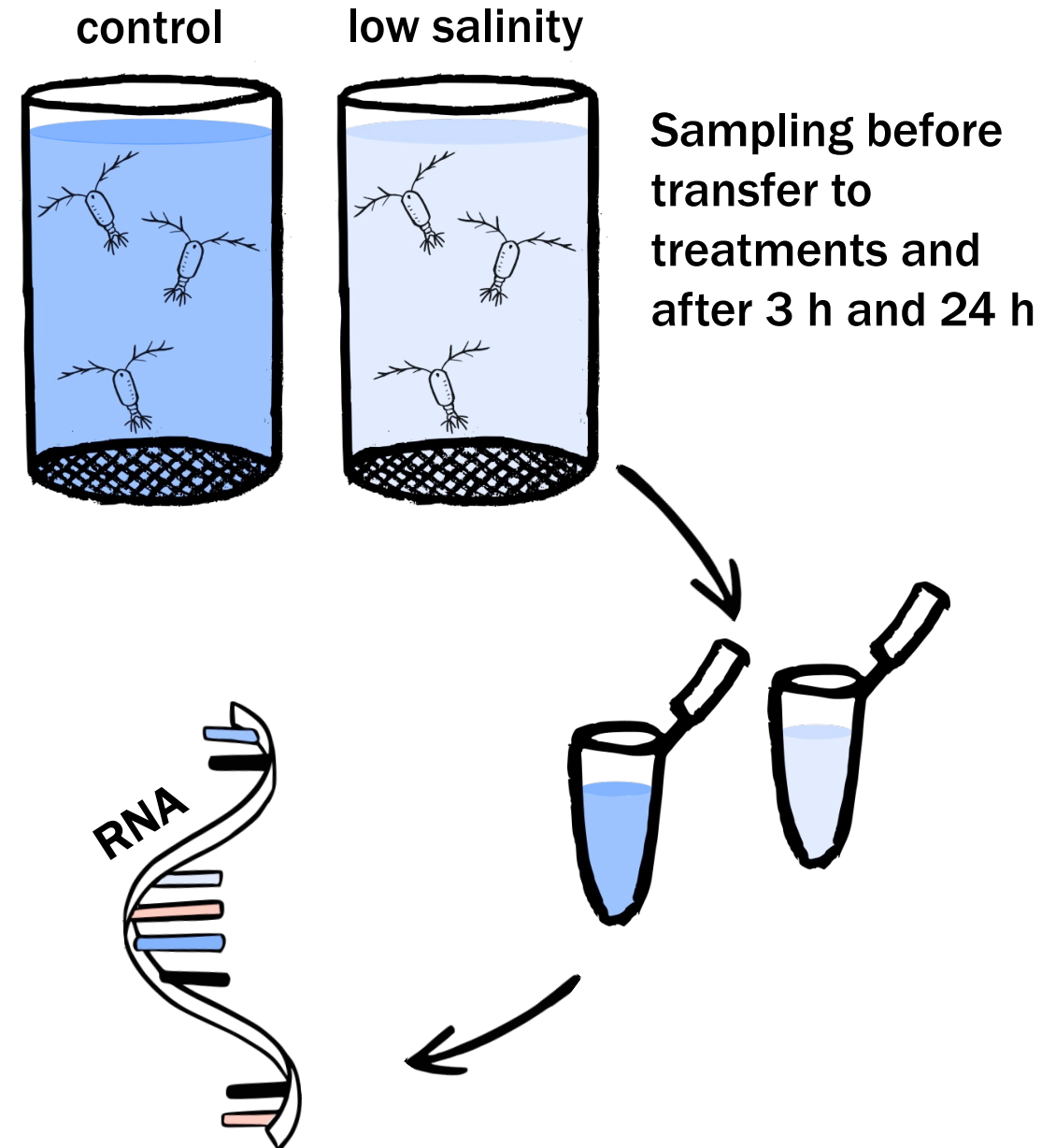
TRANSCRIPTOMICS

How does gene expression differ between populations?
Is there a common/unique response to low salinity stress?

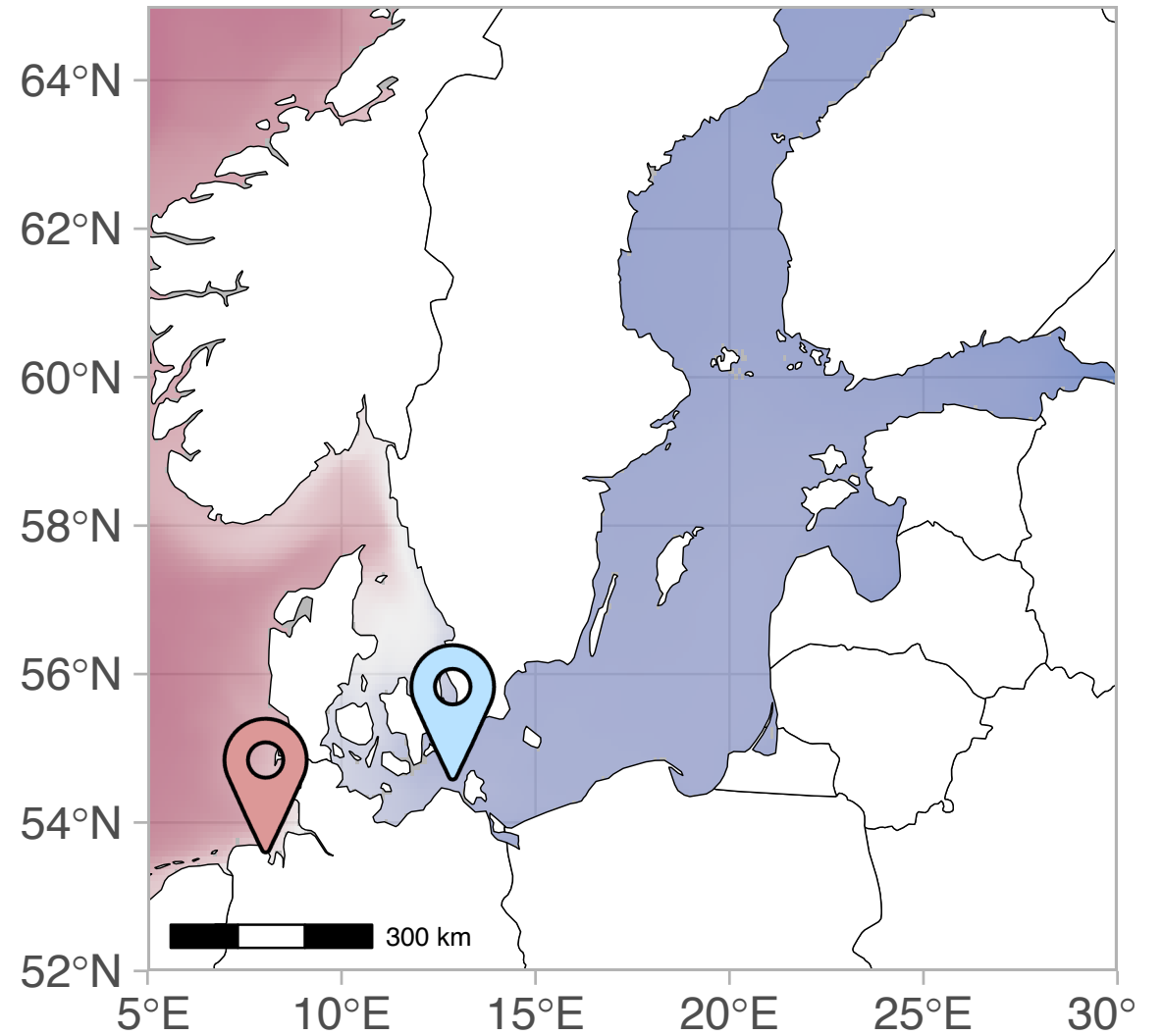
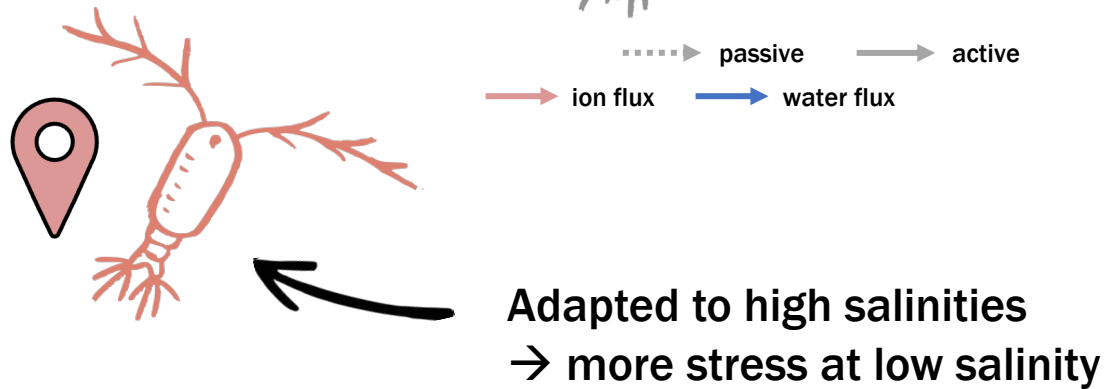
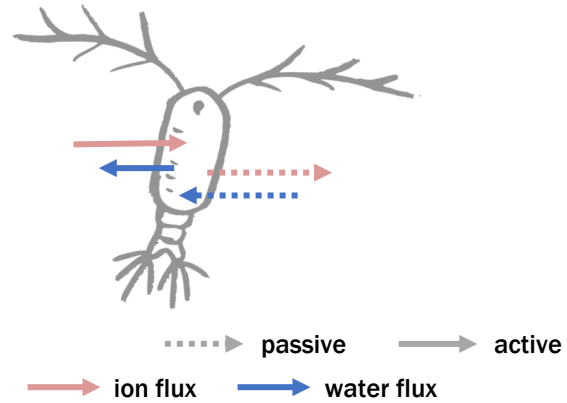
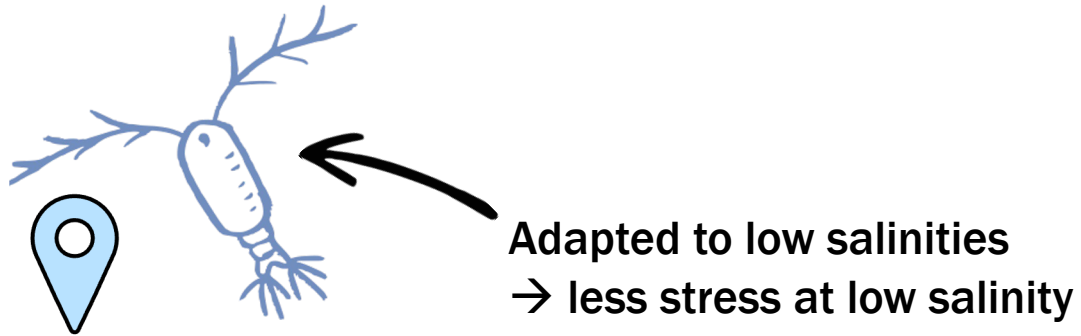
CONCEPT AND EXPERIMENTAL SETUP



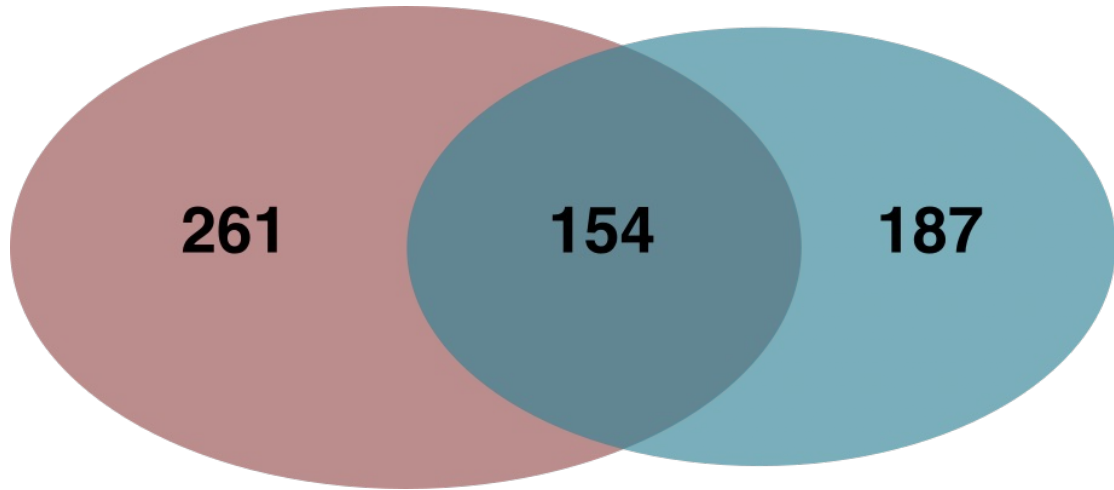
Transcriptomics = Analysis of all RNA transcripts at a given time to understand what processes are active or inactive



EXPECTATIONS

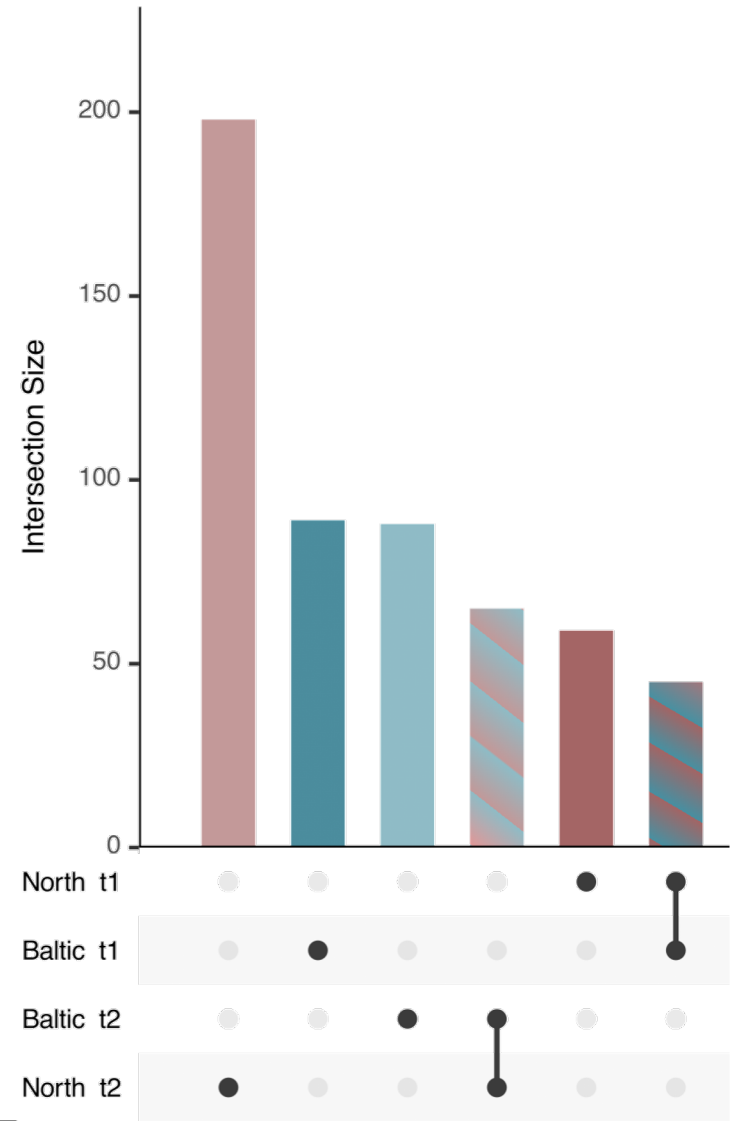
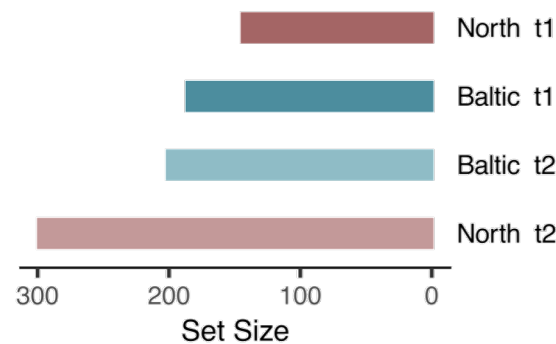
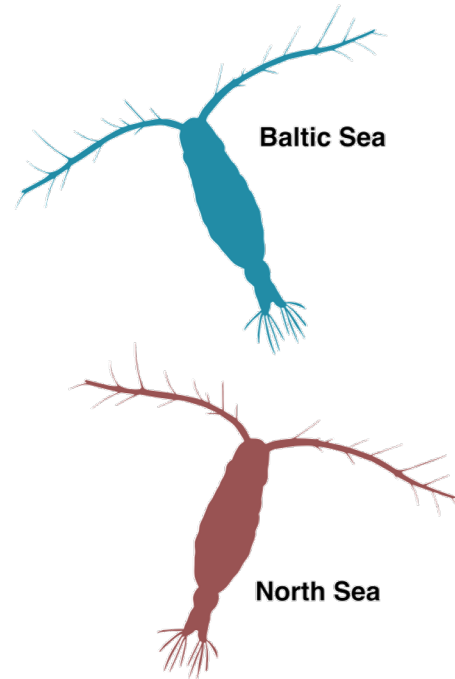


SOME SHARED BUT MANY UNIQUE TRANSCRIPTS

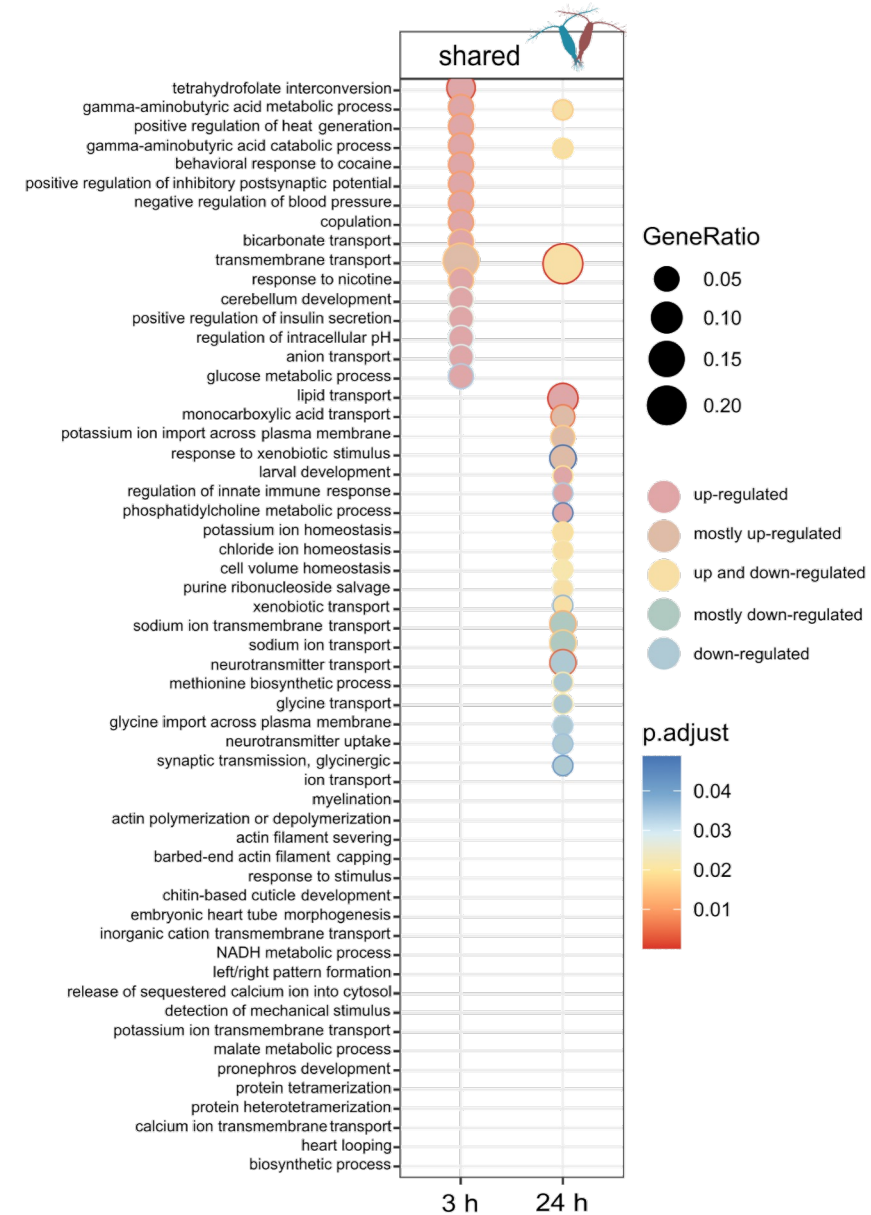


North Sea copepods have more differentially expressed genes, especially after 24 h

→ Struggle to conform



COMMON RESPONSE TO LOW SALINITY STRESS

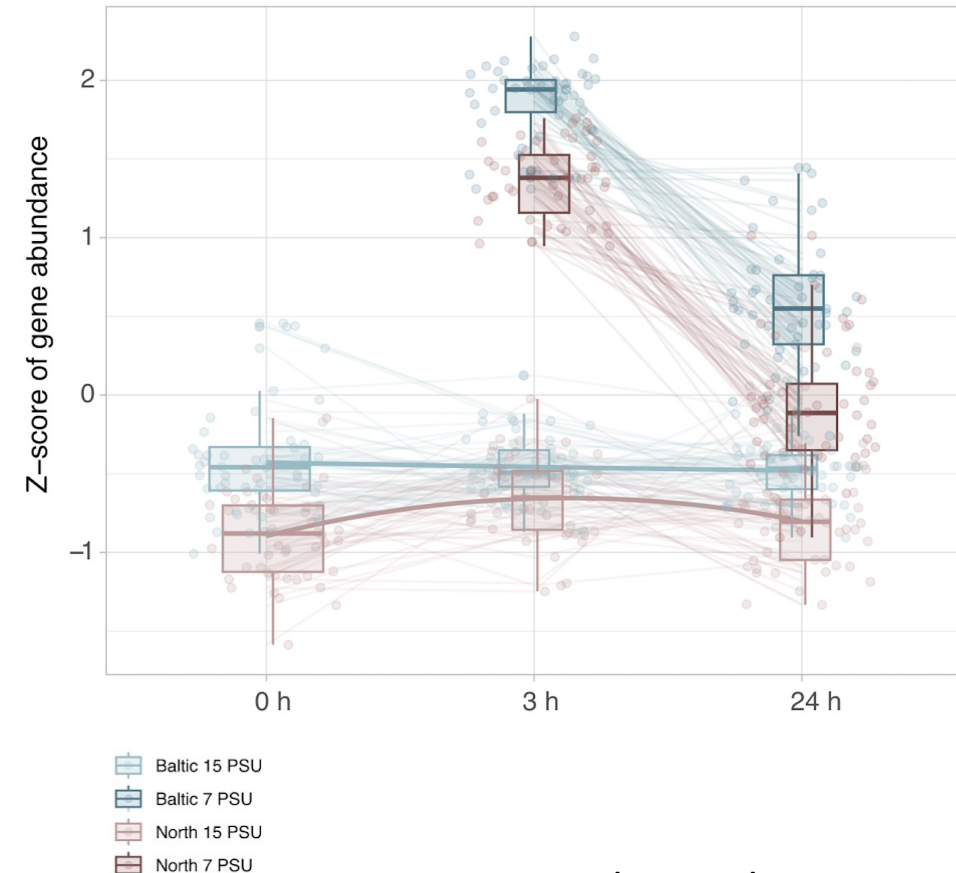
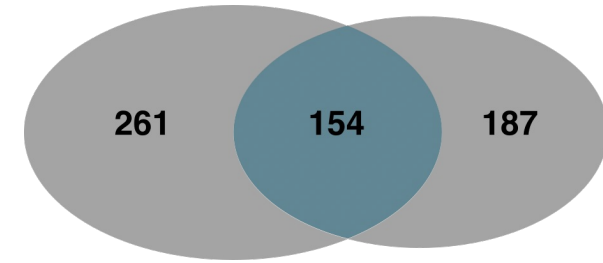


After 3 hours:

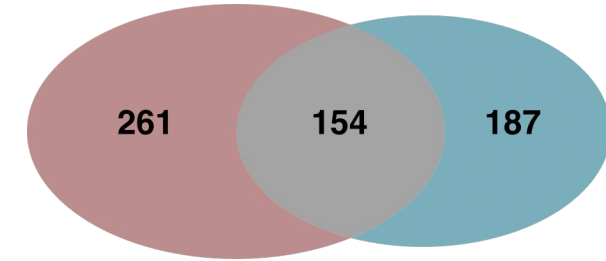
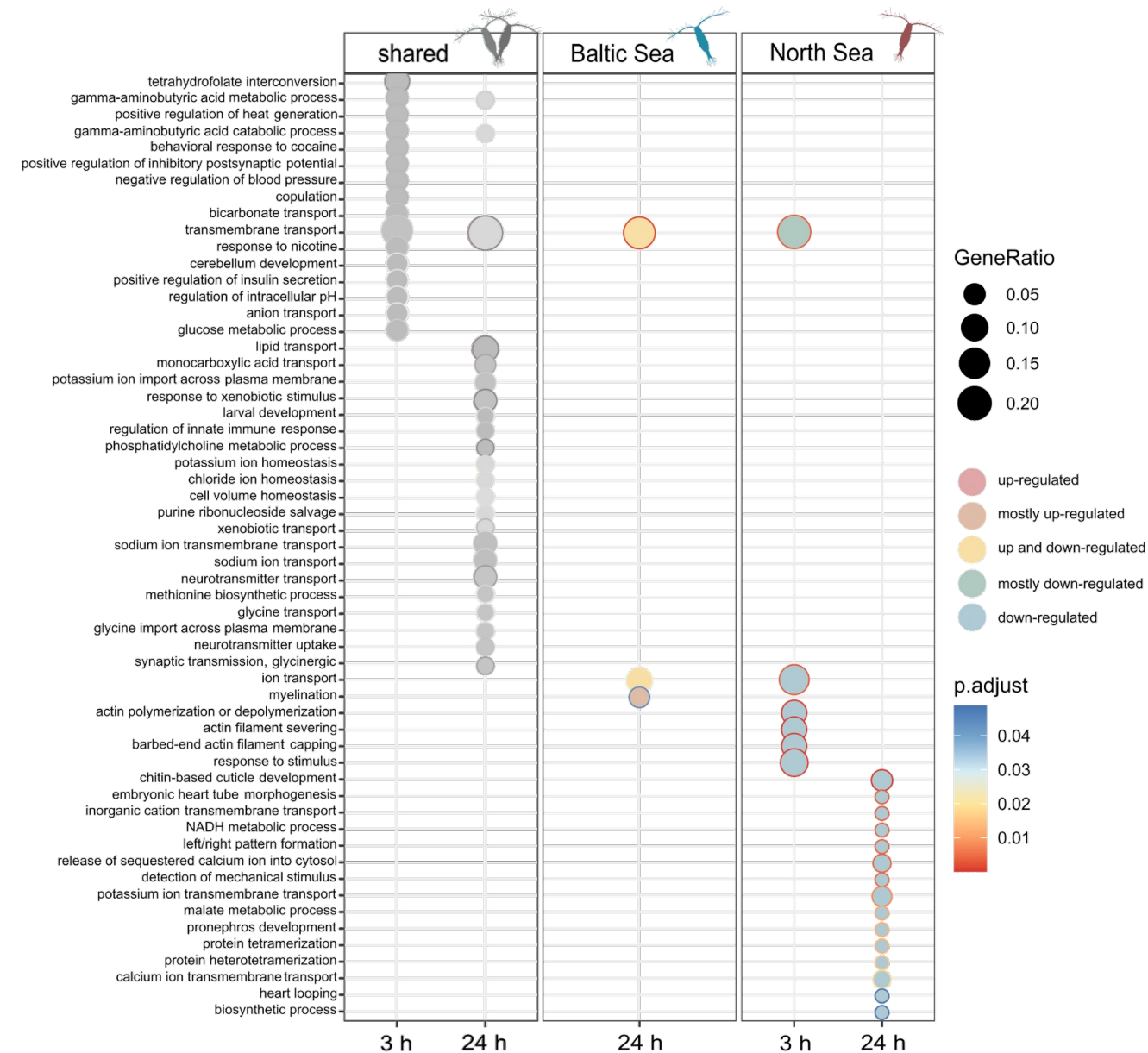
- Metabolic regulation
- Neurological and behavioral regulation
- Ion transport and pH regulation

After 24 hours:

- Various transport functions
- Homeostasis
- Not only up-regulation but more nuanced response



UNIQUE RESPONSE TO LOW SALINITY STRESS



Baltic Sea:

→ Few additional functions, none for t1

North Sea:

→ Many additional functions, time differentiation

→ Mostly downregulation, suppression of metabolic functions to conserve energy

→ Struggle to comply with low salinity

TAKE HOME MESSAGES

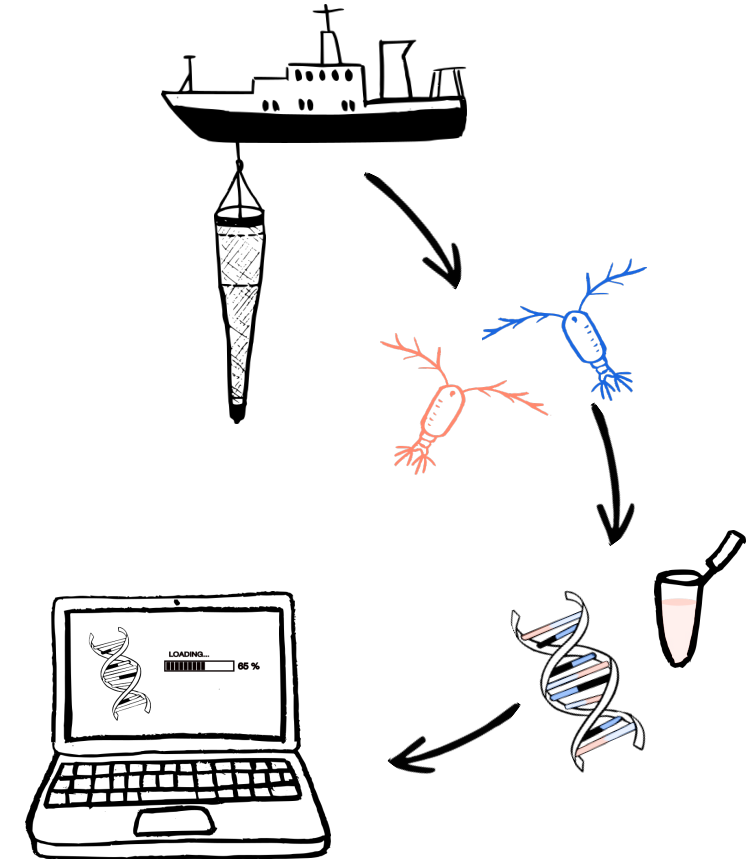
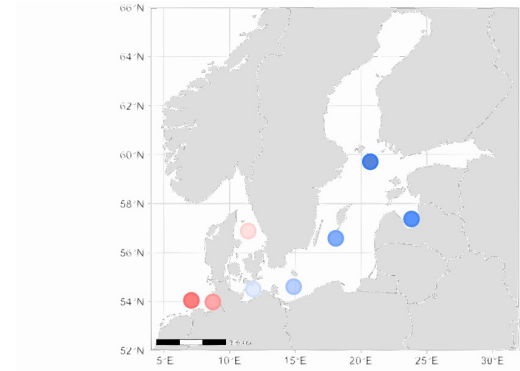
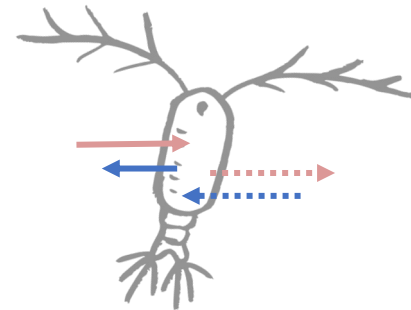
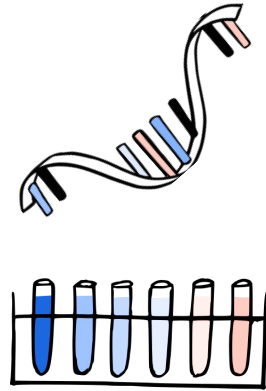
Differences in gene expression and fitness

→ Local adaptation

Tolerance to low salinity in both populations

→ Osmoregulatory strategy

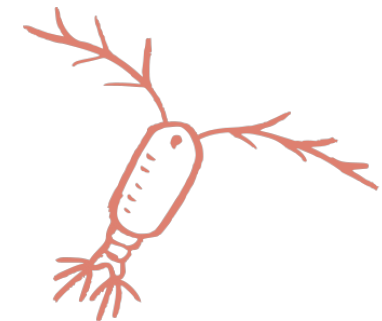
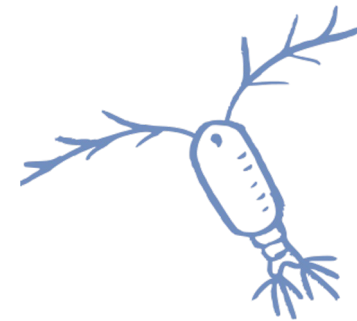
ONGOING: population genomics to identify genetic basis and verify findings



OUTLOOK

**High resilience to environmental stressors is evident in both populations
→ ensure survival and buy time for adaptations to occur**

Adaptive potential could help copepods cope with future environmental change



QUESTIONS?

THANKS TO

Reid Brennan

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Frank Melzner

Jasmin Renz

Meike Stumpp

Sheena Chung

Gianina Consing

HPC @ CAU

RD3 @ GEOMAR

Scientists and crew aboard
cruise AL580