Consortium 1: Pelagic ecosystems

Adaptations to ocean acidification in mesozooplankton WP 1.14:

Evolutionary response to ocean acidification

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Introduction

Very few studies exist dealing with the evolutionary responses of species to ocean acidification (OA), even though the selective forces of OA are potentially strong. Hence, we investigate selection processes in zooplankton. Potential patterns of adaptation to OA will be uncovered by integrating field observations, laboratory experiments and genetic analyses.

Long-term selection experiment

To investigate the adaptability of mesozooplankton to OA a long-term selection experiment with Acartia tonsa a common copepod in the North Sea were set-up.

- 100 I plastic tanks with artificial sea water (ASW; 32 psu) at 18° C and constant darkness
- 15.000-20.000 individuals per tank
- pCO₂ selection lines (SL): 800 μatm A, B, C; 200 μatm D, E, F
- ph-value: 200 tanks: 8.5 \pm 0.2; 800 tanks: 8.0 \pm 0.2
- running time until now: 12 months / 24 generations

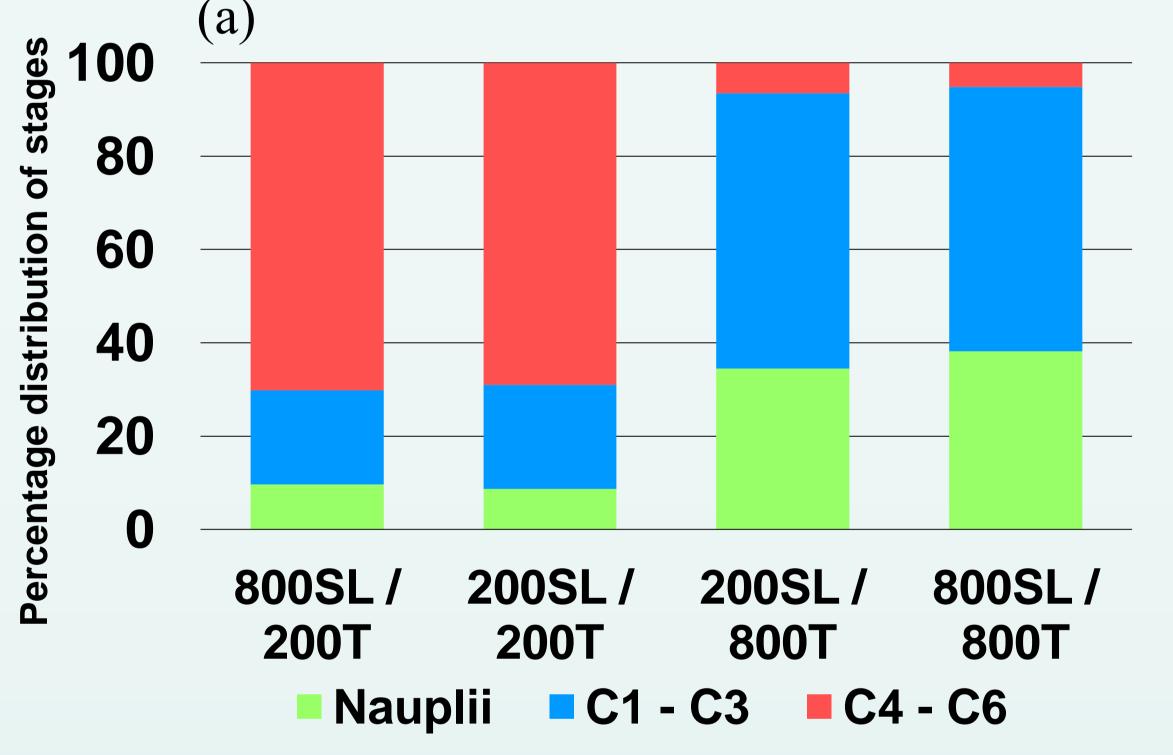






Fig.1 (a-c) Copepod and food cultures. 1(a) Pyrenomonas salina cultures bubbled with different CO₂ concentrations 1(b) Chemostats of the backup algae cultures 1(c) Aeration system of the copepod tanks

Transplantation experiments



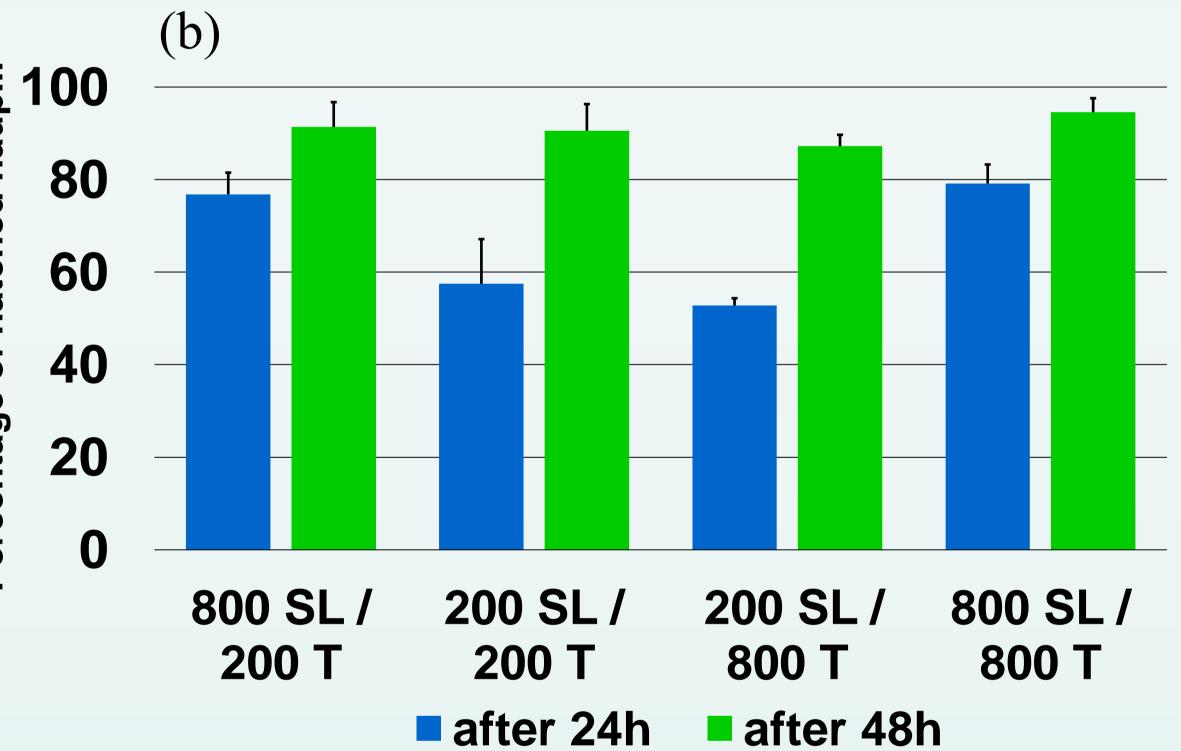


Fig.2 (a,b) Transplantation experiments results. The x-axis shows the experimental treatments; (800 SL) selection line manipulated with 800 μatm CO₂, (200 SL) selection line manipulated with 200 μatm CO₂, (800 T) treatment with 800 μatm CO₂ manipulated water and food, (200 T) treatment with 800 μatm CO₂ manipulated water and food, (200 T) treatment with 200 µatm CO₂ manipulated water and food. (a) Diagram of the developmental stage distribution. The left y-axis shows the percentage of the respective stages. The legend shows the developmental stages; (Nauplii) nauplii stages 1-6, (C1 - C3) copepodite stages 1-3, (C4 - C6) copepodite stages 4-6. The ANOVA showed only a significant treatment effect (p < 0.05) but none for treatment effect (p < 0.05), but none for the selection line nor for an interaction (Data pooled from N = 3 replicates). (b) Diagram of the percentage of hatched nauplii. The left y-axis shows the percentage of the hatched nauplii. The legend shows the two sampling points; The ANOVA showed only a significant selection line effect (p < 0.05), but none for the treatment nor for an interaction (Data pooled from N = 3 replicates).

Set up

transplantation experiments were conducted (4 food. Developmental rate, egg hatching rate were measured.

Results

To test if the copepods adapted to their selection line A. tonsa developed much slower under high CO₂ conditions but conditions within 6 month / 12 generations there was no difference between the SLs and thus no detectable adaptations to CO₂ regarding the developmental speed. treatments, 3 replicates). The copepods from both CO₂ Furthermore there was no detectable difference between the high SLs were cross incubated with different water pCO₂ and low CO₂ treatments and selection lines according the amount of concentrations in connection with the corresponding produced eggs. Additionally the hatching rate did not vary between production and the CO₂ treatments whereby the nauplii from the high CO₂ selection lines hatched significantly faster in the first 24 hours.

On going studies and future plans

Mesozooplankton / Metabarcoding Gene expression

in species composition and genotype experimental populations frequency between (suffering under ocean acidification compared present-day CO₂ conditions) were detected metabarcoding. Samples were taken during a mesocosm experiment in the Gullmarsfjord in Sweden 2012.

We plan to carry out an lab experiment with A. tonsa to investigate gene expression pattern in various CO2 environments. From previous experiments we already known that copepods suffer most under a low food quality due to a changed stoichiometry of the algae. Under high CO₂ conditions (800 µatm pCO₂) they spend most of their energy in reproduction and therefore develop slower. We therefore focus on genes which are involved in growth processes, moulting and reproduction.





