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## **Copepod oxygen consumption along a salinity gradient** 1 Tanguy Soulié<sup>1,2,\*</sup>, Jonna Engström-Öst<sup>3,\*</sup>, Olivier Glippa<sup>3</sup> 2 3 4 <sup>1</sup>Département de Génie Biologique, Institut National des Sciences Appliquées de Toulouse, Toulouse, France 5 <sup>2</sup>MARBEC (MARine Biodiversity, Exploitation and Conservation), Univ Montpellier, CNRS, 6 7 Ifremer, IRD, Montpellier, France <sup>3</sup>Novia University of Applied Sciences, Raseborgsvägen 9, FI-10600 Ekenäs, Finland 8 9 10 \*Corresponding authors: Tanguy Soulié, tanguy.soulie@gmail.com Jonna Engström-Öst, jonna.engstrom-ost@novia.fi 11 12

#### 13 Abstract

Climate change induced salinity decrease is currently occurring in many estuarine coastal 14 15 zones, due to increased outflow of freshwater. This freshening can be a problem for brackishwater animals, already living on the edge of their salinity tolerance. In the current study, we 16 measured oxygen consumption of common copepod Eurytemora affinis (hereafter E. affinis) 17 along a natural salinity gradient in the western Gulf of Finland. The salinity varied between 3 18 in the inner bay and 7 in the offshore area along the gradient, and pH varied between 7.05 and 19 7.86. Our results show that respiration increased with decreasing salinity, as expected for a 20 genus more commonly found in estuarine/saline waters, even if it has colonised brackish waters. 21

Our results suggest that future decreasing salinity could enhance respiration rate, and so energy requirements, of large-bodied zooplankton in estuarine areas such as the Baltic Sea and could lead to lower food quality availability for coastal planktivorous fish, such as herring and sprat.

25 Key words: oxygen consumption, coastal zone, salinity, zooplankton, climate change

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#### 27 **1. Introduction**

Surface water salinity is slowly decreasing in many coastal estuarine areas due to climate 28 change related processes (Kuosa et al., 2017; Nielsen et al. 2020), such as increased 29 precipitation causing more freshwater run-off (Graham, 2004). Certain basins, such as the Gulf 30 of Finland and the Gulf of Bothnia in the brackish Baltic Sea could therefore be subjected to 31 profound changes in hydrography (Meier et al., 2012; Almén et al. 2017), with resulting 32 consequences to biodiversity and plankton community composition (Ojaveer et al., 2010; 33 34 Kuosa et al. 2017; Mäkinen et al. 2017). Salinity changes can have great impact on aquatic invertebrates especially in brackish-water areas. According to Mäkinen et al. (2017), the 35 freshening of the seawater is expected to benefit small-bodied brackish species at the expense 36 37 of marine large-bodied copepods.

Already in 1934, Remane (cited in Whitfield et al., 2012) showed that the largest abundance of brackish-water species was found in areas with salinity level between 5 and 7. *Eurytemora affinis*, the species in focus in the current work, has been thoroughly studied in relation to salinity. Invasion ecological studies and biogeographical work using *E. affinis* as a test animal demonstrate its high ability to invade freshwater systems from a saline environment (Lee 1999, Lee et al. 2012, 2015). Notably, high food concentration significantly raised low-salinity tolerance in *E. affinis* (Lee et al. 2016). Embryonic development time, inter clutch time and clutch size were fairly stable in salinities ranging from 5 to 15 (Devreker et al., 2009), and egg
hatching success was highest between salinities 5 and 20 (Kuismanen et al. 2020). However,
limited knowledge is available on animal respiration in the brackish-water environment, and
how oxygen consumption will change with fluctuating salinity (Feely et al. 2010).

Seasonal hydrography can show large fluctuations, especially in eutrophicated coastal zones, 49 due to primary production, respiration, oxygen deficiency, ocean acidification and precipitation 50 (Waldbusser and Salisbury, 2014, Humborg et al. 2019). Community respiration (i.e., carbon 51 dioxide release) is an important factor that will contribute to coastal hydrography variations 52 53 (Sunda and Cai, 2012). The model by Sunda and Cai (2012) predicts a larger effect of respiration on the carbonate chemistry processes in the Baltic Sea than in other areas (here: the 54 55 Gulf of Mexico) due to lower average salinity and temperature. They also show that the greatest 56 respiratory increases in  $pCO_2$ , associated with parallel decreases in pH, occurred in the lowest salinity (here 4), consistently over a large range of temperatures. The model was run at several 57 temperatures and showed similar effects for all. 58

*E. affinis* is a frequently studied calanoid copepod. It is considered euryhaline, as it can tolerate
and live in a wide range of salinities (Lee et al., 2003). *E. affinis* is an important grazer on
microalgae in the planktonic food-web and constitutes a central food source for planktivores,
such as herring, sprat, three-spined sticklebacks, and mysid shrimps in the Baltic Sea
(Viherluoto et al., 2000; Peltonen et al., 2004).

*E. affinis* populations are decreasing in the Baltic Sea, which is suggested to be a combined
consequence of eutrophication, climate change and over-fishing (Suikkanen et al., 2013;
Mäkinen et al., 2017). In this regard, the aim of the current study was to measure respiration
rates, i.e., oxygen consumption (Gyllenberg & Lundqvist 1979, Li & Gao 2012) in the common
crustacean copepod *E. affinis*. The study was performed along a salinity gradient in a brackish-

water area. We sampled copepods at four different sites of different salinity in the western Gulf 69 70 of Finland and transferred them to respiration chambers in the laboratory. Our main hypothesis was that salinity affects the respiration rate of *E. affinis* (here female egg-carrying individuals) 71 negatively, because of several vital rates being negatively affected by salinity <5 (Devreker et 72 al. 2009, Kuismanen et al. 2020). The salinity gradient used in the present study stretched from 73 3 in an inner coastal bay to 7 in a pelagic offshore area. The salinity gradient is located at the 74 75 entrance to the Gulf of Finland, which is known to be a site of strong hydrographic variability affected by strong south-westerly winds, upwellings from the deep Baltic proper, and 76 freshwater inputs from River Svartån (Alenius et al., 1998). 77

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#### 79 2. Material and methods

#### 80 <u>2.1 Sampling</u>

We collected copepods along a brackish salinity gradient to reveal differences in oxygen 81 consumption of the euryhaline species Eurytemora affinis. Sampling of seawater and 82 zooplankton took place at four sites along the gradient, from June 19<sup>th</sup>, 2017, to August 15<sup>th</sup>, 83 2017. The sampling sites were Pojoviken Bay (60°02'7.10"N, 23°31'6.50"E), Rävnäs 84 (60°0'6.238''N, 23°27'4.826''E), Storfjärden (59°52'5.6''N, 23°15'1.4''E) and Längden 85 (59°46'4.64" N, 23°15'6.76"E), reaching from a shallow (10 m) less-saline site (Pojoviken 86 Bay) to a deep (60 m) more-saline pelagic site Längden off Tvärminne archipelago (Fig. 1A.). 87 Those sites were chosen as they differ mostly by their salinity due to their different distance to 88 the shore, even if factors other than salinity could greatly differ sporadically over a year. The 89 water from these sites was used for the respiration experiments. Sampling took place once or 90 twice per week during the summer growth season. During each sampling occasion we took a 91 Conductivity, Temperature, Density (CTD) profile (Mini CTD, www.valeport.co.uk) for 92

salinity and temperature between the surface and the depth (Table 1). Note that at Längden,
CTD was not deployed to the bottom (60 m depth) but only at 53m due to technical constraints.
As copepods migrate from being in the bottom during the day to being at the surface at night,
average salinity and temperature from bottom to surface were calculated and used in the
analyses, and are presented in Table 1.



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Figure 1. Location of sampling sites (A.) and experimental set-up (B.) In A., P. B.: Pojoviken 99 Bay, Rävn.: Rävnäs, Storf.: Storfjärden, and Längd: Längden. In B., an oxygen micro-optode 100 (1) was put in 4mL glass chambers (2) equipped with a magnet (3) and a mesh (4). The 101 respiration chambers were placed onto a rack (5) immersed in a temperature-constant water 102 bath (6), which was connected to the stirring controller device (7), and to the laptop (8). The 103 bath water temperature was continuously monitored (9). The oxygen micro-optode was 104 connected to the laptop, which recorded the measurements every second and displayed the 105 respiration rate in real-time. 106

Seawater was sampled from the targeted depth (bottom water) using a 2 L Limnos water 107 108 sampler. We sampled water in 250 mL airtight glass bottles in triplicate for pH (Table 1), to be 109 determined at room temperature in the laboratory later the same day. Field samples were adjusted to room temperature and measured with a WTW Inolab 720 pH-Meter, calibrated with 110 111 buffers pH 7 and 10. For chlorophyll a analysis, seawater (200 mL) was filtered on a 25 mm glass-fibre filter (GF/F Whatman). Subsequent to 5 mL ethanol extraction overnight, samples 112 113 were measured in triplicate in a 96-well plate, using a spectrophotometer (Varian Cary Eclipse Fluorescence Spectrophotometer) at 664 nm incident ray. For the calibration of the 114 spectrophotometer, three standards (1.04, 10.4 and 104 µg L<sup>-1</sup>) were used. The blank was 115 116 calibrated with pure ethanol (96%). Dissolved oxygen (DO) concentration and saturation were monitored at the depth where zooplankton was collected using an YSI DO meter 117 (www.ysi.com). 118

Zooplankton were collected using a 200 µm plankton net with cod end, with tows from the 119 120 bottom to the surface. This was done as zooplankton are distributed deeper in the water column 121 during the day. At the deep site Längden, while the maximum depth is around 60m, the net was lowered to 30 m only due to technical constraints. After each tow, zooplankton was transferred 122 to a cooler containing seawater from bottom water from each site and icepacks. The animals 123 were transported to the laboratory where they were stored and acclimated at 12°C for up to 4 124 125 hours until further experiments. The laboratory temperature was set as 12°C on the first 126 sampling day and kept constant for all the experiments (Rumed P530 Climatic Chamber, www.rumed.de). The salinity and the oxygen content in the cooler were measured prior to 127 experiments. 128

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#### 130 <u>2.2 Copepod handling and respiration measurements</u>

The water used for the respiration experiment was collected in 8 L containers, transported in 131 132 coolers and transferred to a climate chamber at arrival. To measure the oxygen consumption rates from four different sites along the salinity gradient (salinities 3-7), two oxygen 133 microsensors (optodes) were used (Fig. 1B.). Approximately 12 female copepods were 134 randomly selected, sorted on ice and acclimated for ~1-2 h prior to the start of the measurement 135 incubation in a 400 ml glass beaker in 12°C. Ten ovigerous females (egg numbers not recorded) 136 in good condition were transferred from the beaker to the measurement unit (= 4 ml glass) 137 cylinder chamber) filled with filtered seawater (see above) and acclimated for another 15 min, 138 before respiration rates were monitored for 2h. The chamber was equipped with a glass ring 139 140 and a 200 µm mesh circle and a glass-coated magnet, to keep the water properly circulated in 141 the chamber. The mesh was used to prevent the copepods from being injured by the magnet.

Respiration rates were monitored at  $12^{\circ} \pm 0.1^{\circ}C$  (mean  $\pm$  SD) using two micro-respiration 142 multimeters (four-channel multimeter, Unisense A/S). The setup enables a continuous and 143 precise follow-up of dissolved oxygen concentration in the chamber (1 measurement  $s_{-1}$ ). The 144 0% DO concentration calibration was done using a 0.1M NaOH solution (50 mL) in which 1 g 145 sodium ascorbate was added. The solution was left to incubate for 2h in order to reach the 0% 146 147 saturation point. The 100% DO calibration was achieved by vigorous oxygenation of the water. The micro-chambers were placed on a plastic rack in a temperature-controlled water bath (12 148 149  $\pm 0.1^{\circ}$ C). Stirring (500 RPM) was achieved using a glass-coated magnet placed in the chamber, 150 used to prevent the formation of an oxygen gradient in the chamber. The experiments were run in a dark climate chamber, for at least 2h. Real-time data were recorded on a laptop via the 151 Unisense® rate software. The connection between the oxygen sensors and the laptop was done 152 153 with a Unisense® MicroOptode Meter. A precise description of the experimental device is presented in Fig. 1B. 154

Depending on the abundance of E. affinis adult female specimens found in the collected 155 156 samples, between 2 and 4 replicates (10 animals repl-) were done for each sampling time. For each sample, one or two control runs without animals were conducted with filtered seawater to 157 compensate for potential microbial respiration, and to ensure that the measured respiration was 158 due to the incubated animals. A non-parametric Kruskal-Wallis test was performed between O<sub>2</sub> 159 concentration at start and after 2h of incubation to make sure that  $O_2$  concentration showed no 160 161 significant decrease over the allotted time period. Filtered seawater was produced from fieldcollected water using 200 µm plankton mesh, and then vacuum-filtered using GF/C glass 162 microfibre filters 1.2  $\mu$ m ( $\bigotimes$  47 mm Whatman®) and stored at +12°C. 163

Individual dry weight was measured for each respiration measurement. To do so, after each 164 respiration experiment, copepods were extracted from the respiration chamber using a Pasteur 165 pipette and put on a Petri dish. Then, animals (10) were picked using forceps and put in foil 166 cups, which were beforehand weighed using a precision scale (Metler Toledo). Foil cups were 167 then put to dry at 60°C in a oven for 24 hours. Then, foil cups were weighed again, and 168 individual dry weight was calculated as the difference of weight between the foil cup when it 169 was empty and with copepods, divided by the number of weighed animals (10). Respiration 170 171 rates were standardized by individual dry weights.

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#### 173 <u>2.3. Statistical analysis</u>

All data residuals were checked for deviations in variances and normal distribution using the Shapiro-Wilk and Levene tests, and analysed using common least square linear regression. The environmental data were checked for collinearity using the Variance inflation factor (VIF), and for autocorrelation using the Durbin-Watson estimate. The VIF represents the ratio of the variance of a model including multiple factors to the variance of the model if it only includes a

single independent parameter. Hence, a high value for the VIF indicates that the tested 179 180 parameter is highly collinear with the other factors included in the model. VIF was calculated with salinity as the single independent variable and environmental parameters (pH, salinity, 181 water temperature, Chl a) for the multifactor model, and a value of 1.823 was obtained, 182 suggesting that salinity is poorly collinear with the other tested environmental factors. The 183 Durbin-Watson test was performed for the residuals of a linear regression model with 184 respiration rate against salinity, and a value of 1.076 was obtained, which is indicative of low 185 positive linear autocorrelation. The dependent response variable oxygen consumption was 186 analysed against the environmental variable in focus (salinity). In addition, an analysis of 187 188 variance (ANOVA) followed by a post-hoc Tukey Honest Significant Difference (Tukey HSD) 189 were used to assess differences between respiration rates among sampling stations. All the analyses were performed using the free software R, version 3.4.3 (R Core Team, 2013). 190

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#### 192 **3. Results**

#### 193 <u>3.1. Seasonal environmental conditions</u>

194 The average salinity from bottom to surface ranged from 3 in the inner archipelago (Pojoviken Bay) to 7.1 in the deep offshore areas (Längden) (Table 1, Fig. 2). The average temperature 195 196 from bottom to surface ranged from 4.1°C to 11.2°C, of which the higher temperatures were recorded in inshore areas (Pojoviken Bay and Rävnäs) during late summer (July - August). 197 Dissolved oxygen (DO) concentration below the thermocline (10 m) was monitored to ensure 198 that the zooplankton in the sampling area were not suffering from hypoxia. DO varied between 199 200 4.6 mg l<sup>-1</sup> in the inshore areas in the middle of the summer to 11 mg l<sup>-1</sup> at Storfjärden. DO was in general highest in early summer and decreased towards the end of the summer. The pH below 201 the thermocline (10 m) peaked at the Storfjärden monitoring station, and the lowest recording 202

203 occurred in the inner archipelago at Rävnäs and Pojoviken Bay (**Table 1**), varying between 7.1 204 and 7.8, and no general trend was detected in the data. Chlorophyll *a* (Chl *a*) concentration 205 varied from 1.3  $\mu$ g l<sup>+</sup> in July at Storfjärden to 5.2  $\mu$ g l<sup>+</sup>at Rävnäs in August, and was generally 206 higher in Rävnäs compared to the other sampling sites (**Table 1**). Finally, the individual dry 207 weight of the sampled copepods varied between 0.001 mg (Längden, Storfjärden) to 0.004 mg 208 (Rävnäs, Storfjärden), with generally higher values in Rävnäs than in other stations (**Table 1**).



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- Fig. 2. Temperature (A) and salinity (B) profiles of the sampling sites Längden (July 10<sup>th</sup>), Pojoviken
  Bay (P. B., July 24<sup>th</sup>), Rävnäs (July 31<sup>st</sup>), Storfjärden (Storf., July 3<sup>rd</sup>). Only one profile per site is
  represented in the figure.

222 Table 1. Hydrographical and biological conditions measured between June and August at the sampling

sites along the environmental gradient used for respiration measurements. Water temperature, salinity

and DO correspond to average values from bottom to surface.

Sampling station	Depth (m)	Date (first to last sample)	Water temperature (°C)	рН	Salinity	Chl a (µg L <sup>-1</sup> )	DO saturation (%)	Copepod dry weight (mg)
Längden	59 – 60	July 10 <sup>th</sup> – August 10 <sup>th</sup>	4.1 - 5.3	7.3 – 7.36	7 – 7.1	1.9 – 2.8	58.5 - 65	0.001- 0.002
Pojoviken Bay	12 - 14	June 29 <sup>th</sup> – July 24 <sup>th</sup>	5.4 - 7.9	7.12 – 7.23	3 - 3.5	2-3.4	49.5 - 50.8	0.002- 0.003
Rävnäs	9 - 10	July 31 <sup>st</sup> – August 15 <sup>th</sup>	9.8 - 11.2	7.05 – 7.2	4.2 – 4.6	4.9 – 5.2	38.2 - 54.5	0.002- 0.004
Storfjärden	32 - 33	June 19 <sup>th</sup> – August 3 <sup>rd</sup>	5.4 - 9.3	7.46 – 7.86	6.1 – 6.7	1.3 – 2.7	69.8 - 88.7	0.001- 0.004

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### 227 <u>3.2. Respiration rates</u>

Respiration rates of *E. affinis* varied between 0.3 and 5.9  $\mu$ mol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup>. When normalized 228 by body weight, they varied from 0.1 to 2.4  $\mu$ mol L<sup>-1</sup> ind<sup>-1</sup>  $\mu$ g<sup>-1</sup> h<sup>-1</sup>, and rates increased 229 significantly with decreasing salinity (Fig 3A). They were significantly different depending on 230 the sampling station (ANOVA,  $p = 3.1 \times 10^{-3}$ , Fig. 3B.). More precisely, respiration rates were 231 significantly higher at Pojoviken Bay (salinity 3-3.5) compared to Längden (salinity 7-7.1) and 232 Storfjärden (salinity 6.1-6.7), while they were not significantly different at Längden, Rävnäs 233 234 (salinity 4.2-4.6) and Storfjarden (Tukey HSD, Fig 3B.). The controls were performed similarly as the treatment, but without animals, and showed no significant changes in oxygen 235 concentration (Kruskal-Wallis, p = 0.7453). 236





Fig 3. A. Respiration rates standardized by body weight against salinity and **B.** Respiration rates 239 standardized by body weight of Eurytemora affinis ovigerous females at the four different 240 sampling stations (Längd.: Längden, P. B.: Pojoviken Bay, Rävn.: Rävnäs, Storf.: Storfjärden). 241 242 In A., the line and shaded area represent the least square linear relationship and its 95% confidence interval, respectively. The equation,  $R^2$  and p-value of the linear relationship are 243 indicated on the upper left corner of the panel. In **B**., for each box, the lower quartile, median 244 and upper quartile values are displayed with horizontal lines. The result of an ANOVA test 245 comparing respiration rates among stations is indicated on the upper left corner of the panel. 246 Letters represent significance groups obtained with a Tukey HSD post-hoc test comparing 247 respiration rates among sampling stations. 248

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#### 251 4. Discussion

The goal of the present work was to experimentally assess how the respiration rate of E. affinis 252 changes along a natural salinity gradient in order to better our understanding about the potential 253 effects of future climate-change induced salinity decrease on a key copepod species of the Baltic 254 Sea. We hypothesized that, due to the fact that *E. affinis* is more commonly found in brackish 255 256 or saline waters, its respiration rate would be higher in low salinities. We incubated *E. affinis* copepods from the Baltic Sea in water collected along a brackish-water salinity gradient and 257 indeed found that their oxygen consumption increased significantly with decreasing salinity. 258 Respiration rates ranged from 0.3 to 5.9  $\mu$ mol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup>, which is consistent with other studies 259

in which Eurytemora respiration rate was assessed. Indeed, Barthel (1983) obtained a mean 260 respiration rate of 10.8 µmol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup> for *E. affinis* at 15°C and a salinity of 12, while Roddie 261 et al. (1984) found respiration rates ranging from 0.5 to 2.4 µmol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup> at 10°C and a 262 salinity ranging from 3 to 10. In addition, Raymont (1959) found a mean respiration rate of 4.4 263 µmol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup> for *E. herdmani* at 15°C. Conversely, Gyllenberg and Lundqvist (1978) found 264 lower respiration rates for *E. hirundoides*, ranging from 0.15 to 0.99 µmol L<sup>-1</sup> ind<sup>-1</sup> h<sup>-1</sup>, at a 265 266 temperature of 4°C. This discrepancy could be due to differences in temperature between experiments, as copepods respiration rate increases with temperature (Heine et al. 2019). This 267 could also be due to methodological reasons, such as the acclimation and exposure time during 268 269 the experiments. Finally, physiological reasons might explain these discrepancies, either due to 270 the initial physiological state of the organisms or to different physiological responses to stress factors. 271

*Eurytemora* is an estuarine genus that has colonized the freshwater environment repeatedly over 272 time, and notably through locally adapted populations (Lee, 1999). The genus inhabits presently 273 274 a wide range of salinities, being most common in salinities between 5 and 15 (Devreker et al., 2009), and which acclimation in different salinities is widely studied (Lee 1999; Lee and 275 Petersen, 2003, Xuereb et al., 2012). The Baltic Eurytemora clade originally arrived from the 276 Caspian Sea (Sukhikh et al., 2013). In our study, the fact that the respiration rate of E. affinis 277 278 was lower in higher salinity conditions might be related to this saline origin, even if it is known 279 that Eurytemora has colonized brackish waters and has adapted to low saline waters by increasing its metabolism (Lee 1999). Eurytemora can tolerate freshwater due to the activity of 280 specialised organs called 'Crusalis organs', which handle ion transport and are structures found 281 282 for the first time in crustaceans (Johnson et al., 2014). However, this ability of Eurytemora to tolerate low salinities leads to an increased requirement in energy, which results in an 283 enhancement of the respiration rate. Consequently, several published works show that low 284

salinity can be stressful for *Eurytemora* (Lee, 1999, Xuereb et al., 2012, Kuismanen et al.,
286 2020).

The salinity gradient studied in the present work is natural and is affected by several 287 288 environmental factors in the coastal zone, partly by freshwater inflow via River Svartån in the northern part, or by upwelling from the main Baltic basin in the offshore area. Also, winds 289 prevailing in the area affect the strength of the salinity gradient (Vuorinen et al., 1998 and 290 references therein). Many studies predict decreasing salinity in the area due to increasing 291 precipitation and resulting run-off (Meier et al., 2012). In our study, salinity could be one of the 292 293 main factors explaining the increase in respiration, as a negative relationship was found between salinity and respiration rate. This result is congruent with Kuismanen et al. (2020) who 294 295 measured egg production, egg hatching success and survival of *Eurytemora* in salinities ranging 296 between 0 and 25 and showed that salinity between 10 and 15 seemed to be the optimal salinity 297 range for Baltic Eurytemora. When the need for energy increases, respiration increases, and stressful conditions induce an elevated energy need (Mauchline 1998, Whiteley, 2011; Li and 298 299 Gao, 2012). Therefore, the enhanced respiration rate of *E. affinis* found in low salinities in the present work suggests an increased food intake of the species to compensate for the extra energy 300 demand, which could have important consequences for the entire food web. 301

In addition to the salinity decrease, other environmental factors could have explained the 302 reported enhancement of respiration. Temperature plays an important role in metabolism. 303 Hence, as the temperatures varied between 4.1 and 11.9°C over the summer at the different 304 sites, we acclimated the animals to the laboratory temperature (12 °C) in order to mitigate the 305 306 potential effect of temperature on the respiration rate, even if the temperature difference between the sampling sites and the incubation in the laboratory may have affected the measured 307 respiration rate. Nevertheless, the water temperature is predicted to greatly increase in the next 308 century due to climate change (IPCC 2018). Hence, warming might affect Eurytemora 309

respiration rates in the Baltic Sea, and the effects of warming may play a synergistic role with decreasing salinity, as it is known that it generally increases plankton respiration rate (López-Urrutia et al., 2006; O'Connor et al., 2009). Moreover, Xuereb et al. (2012) found a synergic effect of temperature and salinity on the gene expression of *Eurytemora* originating from the Seine Estuary, suggesting increased stress for the copepods in warming water and decreasing salinity.

The pH is another important factor modulating copepod respiration rate: in the Mediterranean 316 Sea, acidification had a negative effect on some physiological traits of Acartia clausi 317 318 (Zervoudaki et al. 2014). In the present study, low pH was generally associated with low salinity, and it is not possible to disentangle the effects of pH and salinity on the respiration 319 320 rates. Physiological conditions, age, biomass, UV radiations and food conditions could also 321 play an important role in respiration rate (Ma et al. 2013). In order to mitigate the effect of physiological and food conditions, only ovigerous females were considered, and they were 322 starved shortly before the experiments. It should however be noted that ovigerous females are 323 324 known to respire more than females without egg sacs (Svetlichny et al., 2017).

To conclude, we showed that the respiration rate of *E. affinis* responded as expected; rates increased in lower salinity along a salinity gradient. Even though it is not possible to identify salinity as the only factor explaining the enhanced respiration rate, our results imply that the predicted freshening of the Baltic Sea with climate change could have important consequences on the metabolism of copepods, and *in fine* on the entire ecosystem. In this context of global change, future studies should investigate the interactive effects of multiple environmental factors, such as temperature and salinity, on common copepod respiration in the coastal zone.

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#### **6. Declaration of interest statement**

337 The authors declare no conflict of interest.

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