Sand smelt ability to cope and recover from ocean’s elevated CO2 levels

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1. Introduction

The unprecedented amount of carbon dioxide being released by anthropogenic sources (Kerr, 2010) has surpassed the oceans’ capacity to absorb it, leading to changes in its chemistry in a process known as ocean acidification. With current rates of CO2 emissions, pH is expected to drop 0.3–0.4 units until the end of this century, from 8.1 to 7.8 or 7.7 (IPCC, 2014). These ocean pH changes, which are occurring at an unprecedented rate compared with similar events in the geological past, pose serious threats to marine life, which evolved over millions of years in a stable pH environment (Kerr, 2010).

Initially focused on corals and other calcified organisms due to its carbonate dependence (Orr et al., 2005; Hoegh-Guldberg et al., 2007), research into the effects of elevated CO2 has spread to other non-calcified marine organisms, such as fish. Early life stages might be particularly vulnerable (Melzner et al., 2009) given that physiological regulation mechanisms are poorly developed (Brauner, 2008). The available scientific findings report detrimental effects on larval growth (e.g. Baumann et al., 2012; Chambers et al., 2014; Silva et al., 2016; Rato et al., 2017), metabolism (e.g. Pimentel et al., 2015; Silva et al., 2016), sensorial perception (e.g. Dixon et al., 2010; Castro et al., 2017; Chung et al., 2014), and behaviour (e.g. Devine et al., 2012; Lopes et al., 2016). Moreover, exposure to increased CO2 levels may contribute to divert energy towards internal balance mechanisms, with increased metabolic costs (Silva et al., 2016), resulting in less energy available for other important tasks. As a result, these trade-offs may have major implications for survival, recruitment, and ultimately affect population replenishment and sustainability (Stiasny et al., 2016; Le

A B S T R A C T

Considered a major environmental concern, ocean acidification has induced a recent research boost into effects on marine biodiversity and possible ecological, physiological, and behavioural impacts. Although the majority of literature indicate negative effects of future acidification scenarios, most studies are conducted for just a few days or weeks, which may be insufficient to detect the capacity of an organism to adjust to environmental changes through phenotypic plasticity. Here, the effects and the capacity of sand smelt larvae Atherina presbyter to cope and recover (through a treatment combination strategy) from short (15 days) and long-term exposure (45 days) to increasing pCO2 levels (control: ~515 µatm, pH = 8.07; medium: ~940 µatm, pH = 7.84; high: ~1500 µatm, pH = 7.66) were measured, addressing larval development traits, behavioural lateralization, and biochemical biomarkers related with oxidative stress and damage, and energy metabolism and reserves. Although behavioural lateralization was not affected by high pCO2 exposure, morphometric changes, energetic costs, and oxidative stress damage were impacted differently through different exposure periods. Generally, short-time exposures led to different responses to either medium or high pCO2 levels (e.g. development, cellular metabolism, or damage), while on the long-term the response patterns tend to become similar between them, with both acidification scenarios inducing DNA damage and tending to lower growth rates. Additionally, when organisms were transferred to lower acidified condition, they were not able to recover from the mentioned DNA damage impacts.

Overall, results suggest that exposure to future ocean acidification scenarios can induce sublethal effects on early life-stages of fish, but effects are dependent on duration of exposure, and are likely not reversible. Furthermore, to improve our understanding on species sensitivity and adaptation strategies, results reinforce the need to use multiple biological endpoints when assessing the effects of ocean acidification on marine organisms.

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to assess the biochemical and morphological impacts in larvae induced by exposure to higher levels of energetic metabolism, from aerobic to anaerobic, respectively (Huggett et al., 2004), while removing these reactive metabolites (Abele and Puntarulo, 2004), a set of antioxidant enzymes, including catalase, have been widely used as biochemical, developmental, and behavioural level are addressed. Behavioural lateralization was chosen as the behavioural endpoint as a previous work with sand smelt larvae reported changes in lateralization under elevated pCO2 levels (Lopes et al., 2016). As the early assessment of climate change effects at the community and ecosystem level is particularly difficult, measurements at lower levels of ecological relevance have been extensively used to detect, and possibly extrapolate changes occurring at individual level (Lemos et al., 2010; Pan et al., 2015; Schunter et al., 2016). Biochemical biomarkers, mainly related with energy metabolism and oxidative stress, have been widely used as efficient tools to detect and monitor environmental variations and fitness costs (Huggett et al., 1992; Tonn et al., 2016; Silva et al., 2016; Kamyla et al., 2017). The exposure to environmental stressors may induce the formation of free radicals, contributing to the cellular vulnerability of organisms through lipid peroxidation (LPO) and/or DNA damages (Abele and Puntarulo, 2004), while isocitrate and lactate dehydrogenase are associated with shifts on energetic metabolism, from aerobic to anaerobic, respectively (Huggett et al., 1992).

Together with the biomarkers approach, lateralization was chosen to assess the biochemical and morphological impacts in Atherina presbyter larvae induced by exposure to high levels of pCO2, while gaining insight on the adaptation and recovery to ocean acidification, through a treatment combination strategy and extended exposure period.

2. Materials and methods

2.1. Ethics statement

The current study was performed under the guidelines of the Portuguese Veterinary Authority (DGV-Portugal, following FELASA category C recommendations) and the European directive 2010/63/UE for the protection of animals used for scientific purposes.

2.2. pCO2 treatments

Sand smelt were exposed to control pCO2 levels (pH 8.07, ~515 µatm), and two elevated pCO2 treatments: medium (pH 7.84, ~940 µatm) and high (pH 7.66, ~1500 µatm). Control treatment was selected based on previous pH measures at the sampling site (pH 8.05–8.07); the medium treatment was chosen based on pCO2 levels reported in the coastal areas where larvae of this species occur, where concentrations up to 1170 µatm have been reported under upwelling events (Cabeçadas and Oliveira, 2005); the highest pCO2 treatment was chosen as an extreme condition, which may be reached in upwelling systems if the worst IPCC scenarios are met by 2100.

The experiments were performed with artificial seawater resulting from the mixing of filtered freshwater with a salt mixture (TropicMarin*) and adjusted to a salinity of 34.5%. Except for the control treatment, where pH was directly influenced by ambient air, the two pCO2 treatments were achieved by CO2 injection. pH in the two pCO2 treatments was regulated by a pH computer (Tunze Aquarientechnik, Germany) connected to a pH probe connected to a 200-L sump. The pHbas (National Bureau of Standards Scale), in the three treatments, was daily cross-checked using a portable meter (SevenGo DuoPro, SG23), which was also used for daily measures of temperature and salinity. Due to diffusion pumps on each sump, oxygen levels were always kept above 90% saturation. Each sump was equipped with diverse filtration sets (ultraviolet, chemical, biological, and mechanical), and delivered a continuous supply of recirculated seawater into five 35-L aquariums at a flow-rate of ~600 mL min⁻¹. Aquariums were sealed on top with a clear glass lid to limit CO2 exchange with the atmosphere. Ammonia, nitrates, and nitrites were monitored weekly and kept below critical levels.

Total alkalinity (TA) was determined on a weekly basis, using automated Gran titrations, with certified reference material supplied by A. Dickson (Scripps Institutions of Oceanography, San Diego). pCO2 was calculated in CO2SYS (Pierrot et al., 2006) using in situ temperature, TA, and pH, the carbonic acid dissociation constants given by Millero et al. (2006) and the CO2 solubility coefficient of Weiss (1974) for each of the experimental treatments (Table 1).

2.3. Test organisms and experimental design

Sand smelt larvae were collected at the surface, in the very near-shore, at Portinho da Arrábida, Portugal (38°28′48″ N, 8°58′59″ W), using a 1 mm mesh hand net. Larvae were immediately transported to the laboratory and allowed to recover from handling effects for 2 days in 35-L tanks with recirculating seawater. Except for the two-days recovery period, larvae were daily fed ad libitum with Artemia nauplii and maintained under controlled temperature and salinity, and a summer light cycle of 14 h light:10 h dark. Individuals were then randomly assigned to a control (C), medium (M), or high (H) pCO2 treatment, and maintained in five replicate 35-L tanks, per treatment, throughout the experiment. Initial larval density was 12–15 larvae per tank (N = 73 in control treatment, N = 70 in mid pCO2 treatment, and N = 72 in high pCO2 treatment). After 30 days in treatment, larvae were either transferred between treatments or left in the original treatment for another 15 days, thus totalling 45 days in treatment (CC, CM, MH; MM, MC, MH; HH, HC, HM). No mortality was observed during the experiment, in any of the treatments. Larvae were randomly sampled at 15, 30, and 45 days.

Table 1

<table>
<thead>
<tr>
<th>pCO2 condition</th>
<th>pHbas (pH)</th>
<th>T (°C)</th>
<th>S (psu)</th>
<th>TA (µmol kg⁻¹)</th>
<th>pCO2 (µatm)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>8.07 ± 0.03</td>
<td>16.0 ± 0.3</td>
<td>34.7 ± 0.6</td>
<td>2253 ± 5</td>
<td>516 ± 43</td>
</tr>
<tr>
<td>Medium</td>
<td>7.84 ± 0.01</td>
<td>16.4 ± 0.3</td>
<td>34.9 ± 0.2</td>
<td>2245 ± 3</td>
<td>943 ± 29</td>
</tr>
<tr>
<td>High</td>
<td>7.66 ± 0.02</td>
<td>16.6 ± 0.2</td>
<td>34.6 ± 0.8</td>
<td>2247 ± 5</td>
<td>1504 ± 71</td>
</tr>
</tbody>
</table>
days, for morphometric and biochemical purposes, and sampled at 30 and 45 days for behavioural lateralization tests. Placing larvae immediately in a high pCO2 treatment, instead of adjusting to a gradual pH decrease, represent a toxicity challenge to the organisms and might limit data interpretation due to the possibility of a fast acclimation response to the surrounding environment. However, almost all experiments conducted to assess ocean acidification are acute experiments (Browman, 2016), which makes the present data comparable to most literature. Moreover, a recent review based on short- and long-term experiments revealed little evidence of acclimation to acidification for several species (Nagelkerken and Connell, 2015).

2.4. Behavioural lateralization

A total of 90 larvae – N = 12 per pCO2 treatment at 30 days and N = 6 per pCO2 condition at 45 days – were tested for behavioural lateralization (Table SI – Supplementary material). The test was conducted using a double t-chamber, adapted from Jutfelt et al. (2013), and Domenici et al. (2012) protocol. Briefly, each larva was transferred to the chamber and after 2 min a plastic rod was used to induce the animal to swim forward, without touching, until a left or right turning decision was made. The total turnings, after ten consecutive trial tests for each larva, were registered by direct observation. The relative lateralization index (L0), at the population level, was calculated using the formula (Bisazza et al., 1998):

\[
L_0 = \left( \frac{\text{Turns to the right} \times 100}{\text{Turns to the right} + \text{Turns to the left}} \right)
\]

A mean L0 near zero indicates that a given sample of the population is neither left- nor right-biased in its turning tendency, while extreme values of ± 100 indicate that fish turned right or left, respectively, on all 10 trials (Bisazza et al., 2000). The degree of lateralization at the individual level, irrespective of its left and right direction, was assessed at the individual level using the absolute lateralization index (L0):

\[
L_0 = |L_0|
\]

At the end of the experiment, fish were immediately placed on an ice-cold plastic keeper to slow down the metabolism, photographed under a dissecting stereo microscope for length measurements, using Image-J (v1.48; U. S. National Institutes of Health, Bethesda, Maryland), and frozen at ~80 °C until further processing.

2.5. Morphometric analysis

As described in Silva et al. (2016), the following morphometric traits were measured for each larva, as an indication of development (Jones and McCormick, 2002): standard length (SL), total length (TL), dorsal height (DH), caudal peduncle (CP), eye diameter (ED), and head length (HL) (Table SI – supplementary material).

2.6. Biomarker analysis

A total of 162 organisms – 15 replicate individuals per treatment at 15 and 30 days (C/M/H) and 8 replicate individuals per treatment at 45 days (CC, CM, CH; MM, MC, MH; HH, HC, HM), were weighed and homogenized by an electrical homogenizer, in a 1:30 proportion (m:v) of potassium-phosphate buffer (0.1 M, pH 7.4). After being separated into different microtubes, the homogenate was kept at ~80 °C until further analysis of: lipid peroxidation levels [LPO; in a 50:1 mixture with BHT (2.6-dieter-butyl-4-metylphenol) 4% in methanol to avoid tissue oxidation], DNA strand breaks quantification, and all the components of energy available (Ea), namely total protein, carbohydrate and lipid content, and also the activity of the electron transport system (ETS) as a measure of cellular respiration and energy consumption (Ec). To obtain the mitochondrial supernatant (MS) and the post-mitochondrial supernatant (PMS), the rest of the homogenate was separated into two microtubes that followed different centrifugations, respectively: 1) centrifuged for 5 min at 3000 g (4 °C), separated and stored at ~80 °C for posterior analysis of the activity of lactate dehydrogenase (LDH) and isocitrate dehydrogenase (IDH); 2) centrifuged for 20 min at 10,000 g (4 °C), separated and stored at ~80 °C for posterior analysis of the activity of catalase (CAT) and superoxide dismutase (SOD).

All parameters were analysed and determined in triplicates, with blanks (potassium-phosphate buffer 0.1 M, pH 7.4), using a Synergy H1 Hybrid Multi-Mode Microplate Reader (BioTek Instruments, Vermont, USA).

The protein concentration (PT) in MS and PMS fractions was determined following Bradford (1976) method, using bovine γ-globulin (BGG, Sigma-Aldrich, USA) as standard. Absorbance was read at 600 nm and results were expressed in mg of protein mL−1.

2.6.1. Oxidative stress biomarkers

Following McCord and Fridovich (1969), SOD activity was estimated using microplate and adaptations of Lima et al. (2007). This procedure is based on the reaction of generated superoxide radicals with cytochrome C, which reduction can be followed and measured spectrophotometrically at 550 nm for 10 min (25 °C). The presence of SOD decreases the reduction rate of cytochrome C, by competing for the superoxide radicals. Using a SOD standard of 1.5 U mL−1, the enzyme activity was expressed in U mg of protein−1, where 50% of cytochrome C reduction can be inhibited by 1 U of SOD.

CAT activity was determined based on the consumption of hydrogen peroxide (H2O2) as a substrate following Blairnone (1985). The substrate decrease was followed spectrophotometrically at 240 nm for 1 min (25 °C), using a molar extinction coefficient of 40 M−1 cm−1. Results were expressed in μmol min−1 mg of protein−1.

Following Ohkawa et al. (1979) and Bird and Draper (1984), the estimation of the content of thiobarbituric acid reactive substances (TBARS) allowed assessing LPO levels, quantified colorimetrically at 535 nm using a molar extinction coefficient of 1.56 × 105 M−1 cm−1. Results were expressed in nmol TBARS g ww−1.

DNA strand breaks analysis was based in the DNA alkaline precipitation assay (Olive, 1988) with adaptations from Lafontaine et al. (2000). The damaged DNA present in the supernatant links to Hoesch dye (1 μg mL−1 bis-benzimide, Sigma-Aldrich), allowing the estimation of damage levels by fluorescence, using an excitation/emission wavelength of 360/460 nm. Calf thymus DNA was used to perform the standard curves and results were expressed as μg of DNA mg ww−1.

2.6.2. Energy metabolism related biomarkers

The activity of LDH was measured following Vassault (1983) with adaptations of Diamantino et al. (2001). The process, based on the efficiency of LDH to convert pyruvate to lactate, allowed to detect a decrease in absorbance read at 340 nm for 5 min (25 °C), as a result of NADH oxidation. A molar extinction coefficient of 6.3 × 104 M−1 cm−1 was used, and results were expressed as nmol min−1 mg protein−1.

The determination of IDH followed Ellis and Goldberg (1971) with the adaptations of Lima et al. (2007), was based in the decarboxylation of isocitrate by this enzyme. The increasing levels of NADPH, resulting from NADP+ conversion, allowed the determination of IDH activity measured at 340 nm for 3 min (25 °C). Results were expressed as nmol min−1 mg protein−1, with a molar extinction coefficient of 6.22 × 103 M−1 cm−1.

Following De Coen and Janssen (1997, 2003), total content of protein, lipids and carbohydrates were measured as energy available (Ea) and expressed as MJ g−1 ww−1. All energy reserves were determined and transformed into its energetic equivalent (39.5 kJ g−1 lipid, 24 kJ g−1 protein, 17.5 kJ g−2 glycerogen) (Gnaiger, 1983). Energy consumption (Ec), given by the electron transport system (ETS) activity (King and Packard, 1975), was calculated by formazan generation, which caused an increase in absorbance at 490 nm for 3 min (25 °C). Based on the stoichiometric relation that for every 2 mmol of formazan...
formed 1 mmol of O₂ is consumed (De Coen and Janssen, 1997), the oxygen consumption rate was calculated and then transformed into caloric values using oxyenthalpic equivalents of 484 kJ mol⁻¹ O₂ (Gnaiger, 1983). Results were expressed as mJ h⁻¹ mg ww⁻¹.

2.7. Data analysis

The effect of pCO₂ treatment (control, medium, and high) on behavioural lateralization and biochemical endpoints was addressed by one-way analysis of variance (ANOVA), followed by Tukey post-hoc test for multiple comparisons between treatments, within each timepoint (Zar, 2010). When normality and homocedasticity were not validated, Kruskal-Wallis was applied (Zar, 2010) followed by Least Significant Difference or Games-Howell post-hoc test for multiple comparisons between treatments. Only the effects within each timepoint were considered given the different developmental stages of the larvae over time. The influence of pCO₂ treatment (control, medium, and high) on morphometric traits was analysed using a multivariate analysis of variance (MANOVA). In addition, a Principal Component Analysis (PCA; Ter Braak, 1986) was performed to evaluate the pattern of distribution and correlation based on all endpoints (biomarkers, morphometry, and behaviour) and treatments. When applicable, data were standardized and log (x + 1) transformed (Legendre and Legendre, 1979). All univariate statistical tests were performed with IBM SPSS Statistics 23. PCA was performed with CANOCO version 4.5 package (Ter Braak and Smilauer, 1998). Results were presented as mean ± standard-deviation (SD) and significance level was set at P < 0.05.

3. Results

3.1. Lateralization

Absolute and relative lateralization did not differ significantly among treatments at 30 (F(2,239) = 2.882; P = 0.070, and F(2,239) = 0.939; P = 0.401, respectively; Table SII – supplementary material) and 45 days of exposure (F(2,135) = 0.637; P = 0.542, and F(2,135) = 0.060; P = 0.942, respectively; Table SII – supplementary material). Similarly, no differences in L₁ and I₂ were detectable when also considering the transference between treatments at 45 days (CC, CM, CH; MM, MC, MH; HH, HC, HM) (F(8,435) = 0.989; P = 0.457 and F(8,435) = 1.448; P = 0.203, respectively).

3.2. Morphometry

The results obtained, and here depicted through a MANOVA analysis, show that none of the morphometric traits were affected by pCO₂ treatment at any timepoint. The exception occurred only for HL, which was significantly different between medium and high pCO₂ treatment at 15 days of exposure (F(2,42) = 7.802; P = 0.001; Table SII – supplementary material). Nonetheless, the PCA results for the 15 day of exposure demonstrated a trend for higher lengths in high pCO₂ treatment (dark blue grouping) and smaller larvae in control and medium pCO₂ treatments (light blue grouping), as well as a separation of responses between treatments (Fig. 1a). Thus, increasing pCO₂ levels tends to increase mostly the morphometric traits ED, TL, SL, DH, and HL, also emphasizing the weaker correlation between HL and the remaining morphometric traits, which reinforces MANOVA results regarding acidification effects on this variable – lower HL in medium pCO₂ treatment and higher HL in the highest treatment.

At 30 days of exposure, the treatment separation tendency was maintained, and all morphometric traits became more correlated and more positively associated with the high pCO₂ treatment (Fig. 1b). After 45 days of continuum exposure to the same pCO₂ conditions, although no significant differences were detected, and the treatment separation tendency was maintained, a reverse pattern was revealed for all morphometric traits, with exception of CP (Fig. 1c). CP remained highly related with high and medium pCO₂ treatments, while the strong correlation of the remaining morphometric vectors was at this point more associated to control pCO₂ treatment.

Considering the transfer between treatments (CC, CM, CH; MM, MC, MH; HH, HC, HM), at 45 days no statistical significant differences were detected although the tendency for higher morphometry lengths under more acidified conditions, particularly CP, SL, and ED, was observed once more (Fig. 1d).

3.3. Biomarker analysis

The biomarker analysis revealed different responses over time considering oxidative stress and energy metabolism related parameters (Fig. 2; Table SII; Fig. SII and SIII – supplementary material).

At 15 days of exposure, enzymatic activities of IDH and LDH showed statistically significant differences, with higher activities detected in medium pCO₂ treatment for IDH (F(2,42) = 49.941; P = 0.000; Table SIII and Fig. SII – supplementary material) as well as for LDH (Least Significant Difference; \( \chi^2_{2} = 283.376; P = 0.000; \) Table SIII and Fig. SII – supplementary material). Overall Ea also presented statistically significant differences (F(2,42) = 3.542; P = 0.038; Table SIII and Fig. SII – supplementary material). This scenario was probably influenced by the LP content, which showed the same response pattern (F(2,42) = 4.749; P = 0.014; Table SII and Fig. SII – supplementary material), meaning higher values in the medium pCO₂ treatment than in control both for LP (Tukey; P = 0.014; Table SIII and Fig. SII – supplementary material) and Ea (Tukey; P = 0.031; Table SIII and Fig. SII – supplementary material). Regarding oxidative stress parameters, no statistical differences were observed for any measured parameters at 15 days of exposure. The only exception occurred for LPO, where the levels under the high pCO₂ treatment were significantly lower than in control (Least Significant Difference; \( \chi^2_{2} = 17.139; P = 0.030; \) Table SIII and Fig. SII – supplementary material). These responses can be easily depicted from the PCA analysis (Fig. 1a), where a strong association between the previously mentioned parameters LDH, IDH, LP, and Ea, and the medium pCO₂ treatment is observed (light blue grouping) and determinant to explain 71.6% of the total variance expressed. Apart from the increased morphometric traits already mentioned (Section 3.2), at this time point the high pCO₂ treatment was also associated with higher Ec values and lower carbohydrates (CBH) levels were detected in larvae from high pCO₂ treatment (Least Significant Difference; \( \chi^2_{2} = 61.658; P = 0.003; \) Table SIII and Fig. SII – supplementary material), being this biomarker highly determinant to explain 65.7% of the total variance expressed in the PCA (Fig. 1b; dark blue grouping).

At 30 days of exposure, a reverse pattern was detected for LPO, with higher levels in high pCO₂ treatment (F(2,42) = 5.588; P = 0.005; Table SII and Fig. SII – supplementary material), while DNA damage levels decreased from control to high treatment (Tukey; P = 0.022; Table SII and Fig. SII – supplementary material). Energy consumption levels were higher in organisms exposed to medium pCO₂ treatment (F(2,42) = 5.249; P = 0.009; Table SIV and Fig. SII – supplementary material), and lower carbohydrates (CBH) levels were detected in larvae from high pCO₂ treatment (Least Significant Difference; \( \chi^2_{2} = 61.658; P = 0.003; \) Table SIII and Fig. SII – supplementary material), being this biomarker highly determinant to explain 65.7% of the total variance expressed in the PCA (Fig. 1b; dark blue grouping). Along with the decreased CBH, and lower DNA damage levels, the high pCO₂ treatment was found highly associated with increased morphometric traits, high LDH, SOD, and IDH activities, as well as with the before mentioned high LPO levels (Fig. 1b). The 30 days exposure to medium pCO₂ treatment was characterized mainly by higher levels of PT and Ec, together with lower levels of LP (Fig. 1b; light blue grouping).

At the longest timepoint, the DNA damage tendency observed at 30 days was reverted at 45 days of exposure, with larvae presenting superior damage levels with both acidification treatments (Tukey; P = 0.000; Table SIII and Fig. SII – supplementary material). A reverse response pattern was also observed for Ec, where the oxygen consumption rates were higher in larvae from the high pCO₂ treatment (dark blue grouping) when compared to the medium treatment (Tukey;
The association between higher DNA damage under acidification conditions is also clear from the PCA analysis of 45 days (Fig. 1c), with higher CP values, and lower SOD, LA and most other morphometric traits, characteristics of organisms under acidification treatments and determinant to explain 85.4% of the total variance expressed (Fig. 1c).

Regarding the 45 days after the treatment transference (CC, CM, CH; MM, MC, MH; HH, HC, HM), the parameters that most contributed to explain the variability of data were DNA damage, Ec, LA, PT, and SOD (Fig. 1d), although only few significant differences were detected for the measured biomarkers, namely for Ec, SOD, and DNA damage (Fig. 2; Table SII and SIV – supplementary material). In particular the DNA damage endpoint highlights the apparent irreversibility of effects detected for organisms that had been previously exposed to either medium or high pCO2 treatments and then transferred to lower acidification conditions (same levels from MM to MC, and from HH to HC and HM; Fig. 1d and Fig. 2c). However, when organisms under control conditions were transferred to medium or high treatments for the same period of time, DNA damage levels increased abruptly (from CC to CM and CH; Fig. 1d and Fig. 2c), greatly contributing for the 53.7% of the PCA total variance (Fig. 1d). Interestingly, and contrary to the previous cases, here all the remaining vectors presented lower dimensions, being less relevant to contribute to the overall pattern.

4. Discussion and Conclusion

During the past decade, behaviour modification has been widely used as an indicator of ocean acidification stress, not only due to its inherent plasticity but also to its direct relation with physiological processes (Nagelkerken and Munday, 2016). Detrimental effects on fish have been related with neuronal functions impairment in acidosis scenarios (Hamilton et al., 2014), meaning that behavioural changes may reflect physiological impacts and compensatory mechanisms (Chivers et al., 2014; Heuer and Grosell, 2014). Studying responses to ocean acidification at different levels of biological organization, from enzymatic activities to behaviour and organism development, has been highlighting changes in energy allocation and dramatic compensations at cellular level, even in cases underlying an apparent resilience (Le Quesne and Pinnegar, 2012; Pan et al., 2015). Despite the importance
of assessing species-by-species impacts and adaptation capacity to ocean acidification, little has been carried out to broaden and link these different areas, hampering our understanding on identifying processes and mechanisms subjected to selective pressures.

Lateralization has been reported as a sensitive behavioural endpoint in some organisms exposed to ocean acidification, which impact was already confirmed in Atherina presbyter larvae (Lopes et al., 2016), who reported decreased lateralization index after 7 and 21 days of exposure to high pCO2 levels (~2080 μatm, pH = 7.61). However, in the present study, the difference was not statistically significant (Table SII). The different results might be attributed to differences in experimental designs – in the present study the highest pCO2 condition was still below the extreme pCO2 level used by Lopes et al. (2016) (~1500 μatm vs. ~2080 μatm), as well as the longer-exposure period of 30 and 45 days. Sand smelt larvae may have acclimated to the tested pCO2 conditions, and/or the highest tested pCO2 level was insufficient to induce changes at the lateralization level.

Considering previous results obtained by Silva et al. (2016), the general morphological analysis from this study revealed the same trend for larger sizes for larvae in high pCO2 conditions and smaller sizes for larvae in medium and/or control treatment, after 15 and 30 days of exposure (Fig. 1a and b, respectively). Corroborating this trend, a treatment effect was detected for HL between medium and high pCO2 treatment, after 15 days of exposure, revealing larger head lengths with increasing pCO2 condition (Fig. 1a). Despite the lack of statistically significant differences among treatments for the other morphometric traits, a reversible tendency was observed after 45 days of exposure to the same pCO2 levels (Fig. 1c). With the exception of CP - maintaining the same trend - all morphometric traits presented a tendency to increase in control conditions, with medium and high pCO2 treatments inducing an opposite effect. Although ocean acidification impacts are often species specific, reduced growth sizes associated with the exposure to high levels of pCO2 have been reported and related mainly with decreased energy levels available for tissue synthesis and other important functions, given that physiological adaptation mechanisms rely on expensive metabolic costs (Baumann et al., 2012; Hamilton et al., 2017). Other studies, on the contrary, have provided evidence of increased growth or a lack of effect under increased environmental pCO2 (Munday et al., 2009b, 2011; Hurst et al., 2012; Perry et al., 2015), which has been associated with more efficient capacity of acidification.
base regulation systems (Brauner, 2008; Melzner et al., 2009). Larger sizes induced by acidification scenarios may be related with changes in energy allocation, rather than increased food intake (Hurst et al., 2012), although elevated pCO2 and low pH may enhance feeding activity through olfactory, gustatory, and consequently appetite stimulations (Munday et al., 2009a) – which were not addressed in this study. Overall, development seems to be differently affected by pCO2 exposures and exposure period. Previously, elsewhere, Silva et al. (2016) reported higher growth in high-acidified conditions – similar to the present work. Nevertheless, extended exposure periods reveal different growth trends, which highlight the importance for additional caution into extrapolating short-term experiments results and the need to address the effects in different life-stages and with longer and more realistic exposures.

Given the tendency for a decrease of lipid peroxidation with increasing acidification values, although with higher lipid content, increase in metabolism and a tendency for smaller body sizes in medium pCO2 treatment after 15 days of exposure, the present results seem to corroborate the assumptions of a change in energy allocation to avoid oxidative stress damage (Fig. 1a, SI and SII). Similar results were observed in Silva et al. (2016), particularly for LPO, which levels decrease with the increase of acidity. Although LPO formation is associated with the increase of oxidative stress (Gutteridge, 1995), even in acidified scenarios with different taxa (Conradi et al., 2016; Freitas et al., 2016a, 2016b), with a 15 days exposure period, the results obtained for Atherina presbyter revealed an opposite pattern with decreasing levels of LPO (Silva et al., 2016). Recent studies have identified a strong effect of elevated pCO2 on the overall fatty acid composition of fish larvae (Frommel et al., 2011; Díaz-Gil et al., 2015), bivalves (e.g., Timmins-Schiffmann et al., 2014), and gastropods (e.g., Valles-Regino et al., 2015). Timmins-Schiffmann et al. (2014) reported reduced levels of highly unsaturated fatty acids (components of the cell membrane sensitive to oxidation) in Crassostrea gigas exposed to acidified conditions, suggesting a contribution and possible explanation to lower levels of LPO. Díaz-Gil et al. (2015) identified the reduction in two essential unsaturated fatty acids (EPA, and ARA) associated with larval escape performance on red drum larvae (Sciaenops ocellatus), under increased pCO2 exposures. These authors also suggest that alterations in fatty acid storage and proportion under acidified scenarios may also influence the development performance (Díaz-Gil et al., 2015). Furthermore, in a study with Atlantic cod larvae (Gadus morhua) exposed to acidification scenarios, increasing lipid content, as well as fatty acid variation, namely on essential fatty acids, resulted in organism vulnerability and tissue damage with consequent significant impacts in physiological functions (Frommel et al., 2011). These changes may challenge growth, survivorship and viability of species under global changes and thus, research on this topic following the present study, including a fatty acid profile analysis, should be further addressed.

After 30 days of exposure to the same pCO2 treatments, an abrupt increase in Ec and therefore in cellular oxygen consumption, was observed for organisms in the medium pCO2 treatment (Fig. 1b and SI), while carbohydrates levels remained elevated and lipids decreased, possibly indicating a higher metabolic demand together with an increased energy intake, although feeding rates were not addressed. On the other hand, an abrupt decrease of carbohydrates levels was detected for larvae from high treatment, along with a trend for IDH and LDH increase (despite no alterations in Ec levels). Given the tendency for smaller sizes, organisms from medium treatment might be allocating their energy reserves to maintain physiological and important functions, as well as developing their antioxidant defence strategies (Pimentel et al., 2015), while organisms from high treatment might be allocating energy mostly to growth and other metabolic pathways. Also, a relevant increase in LPO levels was detected for larvae from high treatment at this stage (Fig. 1b and SI), when compared with the same treatment at 15 days of exposure. This may be due to more energy invested for growth and less for oxidative defence strategies in these early stages, which can be corroborated by the lower levels of CAT and SOD on organisms from high treatment at 15 days of exposure (Fig. 1a).

Although the tendency for higher growth in high pCO2 treatment was maintained for 15 and 30 days of exposure, after 45 days a reverse pattern was detected for morphometric endpoints, presenting a decrease in growth for both acidified conditions (Fig. 1c). Higher levels of Ec are associated with less energy available to grow. This pattern can be observed in larvae from high pCO2 treatment at this stage but also in larvae from medium pCO2 treatment at 30 days of exposure (Fig. 1b and SI), both with increasing levels of energy consumption and consequent lower growth. Nevertheless, the opposite cannot be excluded in either situation as ontogeny stage affects the development of the antioxidant response system (Pimentel et al., 2015), and thus less energy may be needed to fulfil homeostasis maintenance tasks. However, possible disruptions derived from an extended period of exposure (45 days) to acidification may have also occurred, explaining the continuous higher levels of DNA damage in organisms from medium and high treatment at this stage (Fig. SII). Therefore, it may be hypothesized that a continued exposure to ocean acidification scenarios may induce vulnerable and fragility conditions at the individual level with overall impacts that are still yet to fully unravel.

Another goal of this study was to explore if any potential effect of ocean acidification on sand smelt larvae, exposed for 30 days, was reversible after placing larvae back into control conditions or transferring them between the different treatments, for an extra period of 15 days. Results revealed differences only at the biochemical level. Overall, organisms maintained at the same pCO2 conditions for 30 days presented only few significant differences once moved to a different treatment for an extra 15 days (Fig. 1d). Organisms reared in control, when transferred to acidified conditions, had higher levels of DNA damage, whereas organisms prior exposed to 30 days in acidified conditions did not recover from DNA damage when transferred to less acidified scenarios – within the present timeframe - indicating potential irreversible effects even when those individuals were placed back into control conditions (Fig. 2c). These impacts indicate the potential jeopardy of these scenarios even when site/organism restoring is possible, in endpoints highly related to population and ecosystem success (Lewis and Galloway, 2009). To date, although studies on ocean acidification effects devoted to adaptation and recovery analysis can be characterized as a growing research field (Munday et al., 2010; Chivers et al., 2014; Hamilton, 2014; Hasler et al., 2016), more focus is still required to assess the potential of marine species to adapt to future ocean conditions, because adaptation will/have been undoubtedly occur/occurring, but its analysis is still scarce. Recovering capacity after acidification exposure has been reported in a few fish species, such as the Californian rockfish (Sebastes diploproa), which anxiety levels were reversible (Hamilton, 2014), damselfish (Pomacentrus wardi) that showed restored predator avoidance (Munday et al., 2010; Chivers et al., 2014), and largemouth bass (Micropterus salmoides), which movement behaviour was recovered (Hasler et al., 2016). While the long-term costs of adaptation are still unknown, a wide variation of responses among individuals and treatment groups, as well as its different times to recover to normal values, provide an important target of different adaptation abilities and consequent natural selection (Heuer and Grosell, 2014; Rossi et al., 2015). Thus, the determination of possible trade-offs related with acclimation and adaptation costs, such as altered gene expression and metabolic demand, is crucial to understand if a reduced overall fitness and less ability to cope with this environmental stressor may be related with higher tolerance levels to increased acidity and how this may affect overall dynamics and the ecosystem.

Dramatic changes are expected to occur due to global change impacts. Understanding species sensitivity together with adaptation strategies to ocean acidification is of utmost importance to maximize our knowledge of how ocean chemistry changes may affect species, populations and the entire ecosystems. Therefore, linking integrative knowledge across a diversity of approaches and disciplines must be the
basis to reach a better understanding of the interaction between ecological impacts and evolutionary changes due to ocean acidification.

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Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at http://dx.doi.org/10.1016/j.ecoenv.2018.02.011.

References


