RESEARCH ARTICLE



Gonadal antioxidant responses to seawater acidification and hypoxia in the marine mussel *Mytilus coruscus*

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Abstract

This study investigated the combined effects of seawater acidification and hypoxia on the antioxidant response in gonads of the thick shell mussel *Mytilus coruscus*. Mussels were collected along the Shengsi Island, East China Sea, where oxygen and pH fluctuations frequently occur in summer. Mussels were exposed to three pH (8.1, 7.7, and 7.3) and two dissolved oxygen (DO) levels (6 and 2 mg L⁻¹) for 21 days followed by a 10-day recovery period (pH 8.1 and DO 6 mg L⁻¹). Gonad surface area (GSA) and activity of superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GPX), glutathione (GSH), glutathione S-transferase (GST), and malondialdehyde (MDA) in gonad were measured at days 21 and 31. Complex and enzyme-specific responses were observed after the 21-day exposure period. Overall, PCA analysis revealed a stronger effect of pH than DO. Integrated biomarker response (IBR) analysis demonstrated that low pH and DO decreased mussel's antioxidant system and increased oxidative damage with potential consequences for gonad development. Mussels exposed to low pH and DO were only partly able to recover a normal enzymatic activity after 10-day recovery period. This suggests that mussels exposed to short-term pH and DO fluctuations event in the field may suffer lasting negative impacts.

Keywords Acidification · Hypoxia · Gonad · Antioxidant response · Mytilus coruscus

Introduction

Ocean acidification (OA) refers to the perturbation of the seawater carbonate chemistry when excess carbon dioxide from the atmosphere is absorbed by the ocean (Birchenough et al. 2015; Orr et al. 2005; Feely et al. 2004). The atmospheric CO_2 concentration has increased from 280 to 400 ppm since the beginning of the industrial revolution. Projections suggest that

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it may reach up to 1200 ppm by the end of this century leading to an average 0.4 pH units decrease in surface ocean (Caldeira and Wickett 2003). A large body of evidence is showing that these changes have the potential to cause dramatic consequences on marine organisms and ecosystem (Noor and Das 2019; Jin et al. 2020). Among marine animals, bivalves are particularly sensitive to OA (Tan and Zheng 2020) with impacts on survival, calcification, growth, and reproduction.

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Exposure to OA leads to increased energy costs to maintain acid-base regulation and compensate for calcium carbonate dissolution (Anthony et al. 2008; Doney et al. 2009; Nakamura et al. 2011; Orr et al. 2005).

Hypoxia, a decrease in oxygen concentration, is another global driver in coastal waters (Deutsch et al. 2011; Zhang et al. 2010; Schmidtko et al. 2017; Breitburg et al. 2018), which is mainly caused by global warming and eutrophication (Breitburg et al. 2018). Over 400 hypoxic zones were documented in the coastal zone since 2008 for a total of 245000 km² area. For example, hypoxic zones have been observed in the East China Sea since the 1950s (Diaz and Rosenberg 2008; Li et al. 2011; Chen et al. 2020). Large amount of nutrients released from the Yangtze River increased the primary productivity in the East China Sea, which consequently leads to eutrophication and hypoxia. These hypoxic zones could be greater than 15000 km² (Chen et al. 2020).

Previous studies on mussels have showed that expression of oxidative stress-related genes and activity of antioxidant enzyme activities were affected by hypoxia (e.g. Woo et al. 2013; Sui et al. 2017; Nogueira et al. 2017, Huang et al. 2018a, 2018b). These are linked to increased production of cellular reactive oxygen species (ROS) (e.g. Ekau et al. 2010; Levin et al. 2009), and a potential risk of oxidative damage (Hermes-Lima et al. 2015).

Exposure to OA and hypoxia can inhibit gonadal development in marine invertebrates (Parker et al. 2018; Mos et al. 2016; Kurihara et al. 2013; Uthicke et al. 2014; Aguirre-Velarde et al. 2019), as a consequence of shifts in energy budget (Sui et al. 2016), subsequently affecting reproduction and population dynamics. However, the combined effects of hypoxia and acidification are still poorly understood. Marine mussels *M. coruscus* were collected from the Shengsi Island, East China Sea, where they regularly experience low levels of dissolved oxygen and pH during the wet season (Sui et al. 2017). To better understand the combined effect of OA and hypoxia, they were exposed to decreased pH and DO using a full factorial design. We hypothesized that OA and hypoxia will negatively impact their gonad antioxidant responses and their reproduction.

Materials and methods

Animals

Thick shell mussels *M. coruscus* (80 ± 10 mm shell length, 160 ± 20 mg dry tissue weight) were sampled from the Shengsi island, East China Sea ($30^{\circ} 33' 00.945''$ N, $121^{\circ} 49'$ 59.757" E) and immediately transferred to the laboratory. Undamaged mussels were selected and acclimated for 2 weeks in 500L aquaria containing aerated seawater with salinity (25 psu), temperature (25° C), oxygen concentration

(6 mg L⁻¹), and pH value (8.1) The light regime was 12 h light/12h dark and mussels were fed with the microalgae *Chlorella* spp. every 12 h (25,000 cells mL⁻¹).

Experimental design

Mussels were exposed to pH (three levels: 8.1, 7.7, and 7.3) and DO (two levels: 6 and 2 mg L^{-1}). Each treatment had three replicates (tanks) containing 30 mussels. Target pH 8.1 was selected as the present average at the sampling site; pH 7.7 corresponds to the extreme present variability and the extreme pH projected for the year 2100 and the minimum of present natural variability at the sampling site (Li et al. 2014), and pH 7.3 is the expected pH by 2300 (Caldeira and Wickett 2005). DO 6 mg L^{-1} is within the present range of variability at the sampling site while 2 mg L^{-1} corresponds to extreme minimum of documented DO in the East China Sea $(2-3 \text{ mg L}^{-1} \text{ in})$ summer; Chen et al. 2007a, 2007b). pH was maintained by bubbling pure CO₂. The flow rate was controlled by pH feedback STAT systems (DAQ-M) operated by the CapCTRL software (Loligo Systems Inc, Tjele, Denmark). DO was maintained by injecting N2 and air via an O2 regulator (Loligo Systems Aps, Tjele, Denmark). Daily, salinity was measured daily using a multiparameter instrument (model 5200A, YSI, USA) and total alkalinity (A_T) was determined by titration with methyl red-methylene blue. pCO₂, saturation states of calcite (Ω ca), aragonite (Ω ar), and dissolved inorganic carbon (DIC) were calculated using the CO₂SYS software. A summary of the carbonate chemistry is presented in Table 1. Mussels were maintained for 21 days under the treatment conditions and then transferred back to the same condition than during acclimation (salinity 25 psu, temperature 25 °C, DO 6 mg L^{-1} , and pH 8.1) for a 10-day recovery period.

Tissue collection and preparation

Gonads were collected at days 21 and 31. As female accounted for more than 80% of the mussel population used for the experiment, only females were used for analyses. For each treatment, six females were obtained and then dissected. Gonads were carefully excised, surface dried with tissue paper, thoroughly washed with phosphate buffer (50 mM; pH 7.4), and kept on ice. The gonads were photographed and the surface area (GSA in mm²) was measured using the software ImageJ. Aliquots from each tissue pool were immediately frozen in liquid nitrogen and stored at -80 °C for further analysis. The gonads were thawed on ice and homogenized (1:4, w-v) in 0.1 M Tris-HCl buffer (pH 7.5) containing 0.15 M KCl, 0.5 M sucrose, 1 mM EDTA, 1 mM Dithiothreitol (DTT, Sigma), and 40 μ g mL⁻¹ Aprotinin (Sigma). Homogenization was performed at 4 °C using 12-15 strokes of a motor driven Teflon Potter-Elvehjem homogenizer. Homogenized samples were sonicated for 2 min at 0 °C with

Treatments pH*DO	DO (mg L^{-1})	рН	Salinity (psu)	Temperature (°C)	A_{T} (µmol Kg ⁻¹)	DIC (µmol Kg ⁻¹)	pCO ₂ (matm)	Ωca	Ωar
8.1*6.0	6±0.05	8.11±0.01	25.1±0.1	25.1±0.1	2308±9	2044±9	366±9	5.26±0.08	3.36±0.05
7.7*6.0	6±0.03	7.71±0.02	24.9±0.6	25.2±0.1	2291±8	2200±16	1060±58	2.35±0.10	1.50±0.07
7.3*6.0	6±0.02	7.31±0.01	25.0±0.1	25.1±0.1	2209±6	2238±6	2697±31	0.96±0.01	0.61±0.01
8.1*2.0	2±0.01	8.11±0.02	24.9±0.0	24.9±0.1	2310±7	2050±9	370±19	5.21±0.19	3.32±0.12
7.7*2.0	2±0.05	7.70±0.01	25.1±0.1	25.0±0.2	2288±7	2198±11	1062±37	2.33±0.06	$1.49{\pm}0.04$
7.3*2.0	2±0.04	7.30±0.01	24.8±0.1	25.0±0.2	2208±5	2240±6	2730±65	0.94±0.02	$0.60{\pm}0.01$

a Braun Labsonic U sonifier at 50% duty cycles and then centrifuged at 12000g for 45 min at 4 °C. Supernatants were collected for biochemical analysis.

Antioxidant assays

Commercial kits from the Nanjing Jiancheng Bioengineering Institute (Nanjing, China) were used for antioxidant assays. Optical density values were measured using a microplate reader (Flexstation® 3, Molecular Devices, California, USA). The protein content of enzyme crude extract was determined using Coomassie Brilliant Blue (G-250) method (Bradford 1976). The protein content was used to normalize enzyme activities.

Superoxide dismutase assay

The nitro blue tetrazolium (NBT) method was used to measure superoxide dismutase (SOD) activity according to Sun et al. (1988). NBT was reduced to blue forms by O_2^- generated by the xanthine/xanthine oxidase system, which has a strong absorbance at 560 nm. One unit (U) of SOD is defined as the amount that inhibits the rate of NBT reduction by 50%.

Catalase assay

Catalase (CAT) activity was measured using a spectrophotometric assay of hydrogen peroxide based on the formation of its stable complex with ammonium molybdate at 405 nm (Góth 1991). In brief, a H_2O_2 degradation reaction catalyzed by CAT was terminated by adding ammonium molybdate, and the intensity of a yellow complex formed by molybdate and H_2O_2 at 405 nm was measured. One U of CAT activity is defined as the degradation of 1 µmol H_2O_2 per second per mg of protein.

Glutathione assay

GSH level was measured according to the method of Ringwood et al. (1999) by reading the optical density of the

yellow substance formed when 5,5'-dithio-2-nitrobenzoic acid was reduced by glutathione (GSH) at 412 nm. GSH content in the extract was determined as nmol mg-1 protein by a standard curve generated with GSH at diverse concentrations.

Glutathione peroxidase assay

Activity of glutathione peroxidase (GPX) was measured by quantifying the rate of oxidating reduced glutathione (GSH) to oxidized glutathione (GSSG) induced by H_2O_2 (Xia and Zhou 1987). The method for measuring the amount of GSH is described above. One U of GPX is defined as the amount that reduces the level of GSH by 1μ mol L⁻¹ in 1 min per mg of protein.

Malondialdehyde assay

Malondialdehyde (MDA) was assessed by measuring the thiobarbituric acid reactive substances (TBARS). MDA in the lipid peroxide degradation product can be condensed with thiobarbituric acid (ZBA) to form a red product with a maximum absorption peak at 532nm. The TBARS formed was measured in a microplate reader at 532 nm and quantified as malondialdehyde equivalents using1,1,3,3-tetramethoxypropane as the standard (Ohkawa et al. 1979). MDA content in the extract is expressed as nmol TBARS mg⁻¹ protein.

Glutathione S-transferase assay

The glutathione S-transferase (GST) activity was determined according to Habig et al. (1974), using 2 mM 1-chloro-2,4-dinitrobenzene (CDNB) and 2 mM reduced glutathione in 0.1 M potassium phosphate buffer, pH 7.0. The absorbance was monitored for 2 min at 30 °C at 340 nm for a better detection in the microplate reader. One U of GST activity is the amount which catalyzes the conjugation of 1mM of substrate per minute.

Integrative biomarker index

Six biomarkers (SOD, CAT, GPX, MDA, GST, GSH) were integrated in a IBR index according to Beliaef and Burgeot (2002) as: (1) calculation of mean and standard deviation for each biomarker; (2) standardization for each sample: Y = (X - m)/s, where Y = standardized biomarker response, X = response value of each biomarker, m = mean value of the biomarker, and s = standard deviation of the biomarker; (3) Z was then calculated as Z = Y or Z = -Y for biomarker induced or inhibited; (4) score for the biomarker was computed as S = Z+ |min|, where $S \ge 0$ and |min|= absolute minimum value of Yfor each biomarker; (5) scores were visualized in a star plot and correspond to radial coordinates. IBR was finally calculated as:

IBR = Sin
$$\alpha$$
 [($S_i \times S_{i+1}$)/2 + ($S_{i+1} \times S_{i+2}$)/2 + ... + ($S_{n-1} \times S_n$)/2]

where α = angle formed by each two consecutive scores, and *n* = the number of biomarkers.

Statistical analysis

All statistical analyses were performed using SPSS 18.0. Shapiro-Wilk test was used to test the normality of the data, and Levene's test was used to test the homogeneity of the variance prior to statistical analysis. Effects of pH, hypoxia, and their interactions on SOD, CAT, GPX, GSH, GST MDA, and GSA were analyzed by two-way analysis of variance (ANOVA). When relevant, one-way ANOVA, Tukey's HSD post hoc test, and Student's *t* test were used to resolve the effect of individual parameters. Principal component analysis (PCA) was conducted using Origin 2018. Results were expressed as mean \pm SEM and a threshold of *p* <0.05 for significant difference.

Results

Gonad surface area

pH, DO, and their interactions had a significant effect on gonad surface area (GSA) after the 21-day exposure period (Table 2) while only pH and the interaction between pH and DO had significant effects after the 10-day recovery period (Table 3). Overall, exposure to low pH led to a decrease in GSA. Low DO significantly decreased GSA under pH 8.1 after exposure and under pH 7.3 after recovery (Fig. 1).

Antioxidant assays

After the exposure period, SOD activity was significantly impacted by DO (Table 2) leading to an increased activity under **Table 2**Summary of two-way ANOVA testing effects of pH and dis-
solved oxygen (DO) on SOD, CAT, GPX, MDA, GST, GSH, and GSA
in the gonad of *M. coruscus* after 21-day exposure period

Sources		pН	DO	$\mathrm{pH} \times \mathrm{DO}$
	df	2	1	2
SOD	MS	1152.725	4055.217	255.146
	F	2.627	9.241	0.581
	Р	0.089	0.005	0.565
CAT	MS	0.761	0.501	0.061
	F	9.924	6.530	0.802
	P	< 0.001	0.016	0.458
GPX	MS	111.234	108.116	87.260
	F	32.802	31.883	25.733
	P	< 0.001	< 0.001	< 0.001
GSH	MS	97.457	52.493	11.250
	F	9.491	5.112	1.096
	Р	< 0.001	0.031	0.347
GST	MS	1218.627	37.319	55.764
	F	142.138	4.353	6.504
	Р	< 0.001	0.046	0.005
MDA	MS	101.857	0.595	14.596
	F	20.267	0.118	2.904
	Р	< 0.001	0.733	0.070
GSA	MS	19120.090	7327.132	3660.906
	F	21.510	8.243	4.118
	Р	<0.001	0.007	0.026

low DO (Fig. 2). No significant effect was observed for SOD activity after the recovery period (Table 3).

After the 21-day exposure period, pH and DO had a significant effect on CAT activity, but not their interaction (Table 2). Both low pH and low DO led to a decrease in CAT activity (Fig. 3). However, no significant effect was observed after the 10-day recovery period (Table 3).

GPX activity was significantly affected by pH, DO, and their interaction both after the exposure (Table 2) and recovery periods (Table 3). When significant, pH and DO led to an increased GPX activity. After the exposure period, at DO 6 mg L^{-1} , GPX activity was significantly increased at pH 7.7 compared to the other two pHs. At DO 2 mg L^{-1} , GPX activity increased with decreasing pH (Fig. 2.). Exposure to low DO significantly increased GPX activity at pH 8.1 and 7.3 but not at pH 7.7. After recovery, low DO significantly increased GPX activity under pH 7.7 and 7.3 (Fig. 4).

After the exposure period, both pH and DO had a significant effect on GSH activity but not their interaction (Table 2). Exposure to low pH and DO led to an increased GSH activity (Fig. 5). The tested parameters had no significant effect on GSH activity after a ten-day recovery (Table 3).

Table 3Summary of two-way ANOVA testing effects of pH and dis-solved oxygen (DO) on SOD, CAT, GPX, MDA, GST, GSH, and GSAin the gonad of *M. coruscus* after 10-day recovery period

Sources		рН	DO	pH x DO
	df	2	1	2
SOD	MS	301.336	415.817	132.890
	F	0.587	0.810	0.259
	Р	0.562	0.375	0.774
CAT	MS	< 0.001	0.002	< 0.001
	F	0.088	1.231	0.090
	Р	0.916	0.276	0.914
GPX	MS	88.815	399.039	106.971
	F	17.630	79.209	21.234
	Р	< 0.001	< 0.001	< 0.001
GSH	MS	0.855	0.038	8.643
	F	0.064	0.003	0.003
	Р	0.938	0.958	0.531
GST	MS	1476.412	16.854	24.704
	F	52.513	0.599	0.879
	Р	< 0.001	0.445	0.426
MDA	MS	1.109	0.051	0.505
	F	0.258	0.012	0.118
	Р	0.774	0.914	0.889
GSA	MS	14099.556	1345.728	2799.864
	F	23.343	2.228	4.635
	Р	< 0.001	0.146	0.018

After the exposure period, pH, DO, and their interaction had a significant effect on GST activity (Table 2). When exposed to DO 6 mg L^{-1} , GST activity decreased with decreasing pH. Under DO 2 mg L^{-1} ; GST activity was significantly lower at pH 7.3 compared to the other two tested pHs.



Fig. 1 Gonad surface area (GSA) of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L⁻¹) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period. Letters indicate significant effects (p<0.05; post hoc test) between pH treatments within DO level and asterisks significant effects (p<0.05; post hoc test) between DO treatments within pH level



Fig. 2 Superoxide dismutase (SOD) activity in the gonad of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period

Exposure to low DO only significantly impacted by GST activity at pH 8.1 (Fig. 6). After the recovery period, only pH had a significant effect on GST activity (Table 3) leading to a reduced activity at pH 7.3 compared to other two pHs (Fig. 6). The interaction between DO and pH was found significant only after exposure (Table 2).

Only pH had a significant effect on MDA level after the 21day exposure period (Table 2). The MDA level was significantly lower at pH 8.1 compared to the two other pHs (Fig. 7). After the recovery period, no significant differences in MDA levels were observed (Table 3).

Integrated biomarker response

Standardized biomarkers (SOD, CAT, GPX, MDA, GST, GSH) were plotted in IBR star plots (Fig. 8). Overall, the IBR



Fig. 3 Catalase (CAT) activity in the gonad of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period



Fig. 4 Glutathione peroxidase (GPX) activity in the gonad of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L⁻¹) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period. Letters indicate significant effects (p<0.05; post hoc test) between pH treatments within DO level and asterisks significant effects (p<0.05; post hoc test) between DO treatments within pH level

values tend to decrease with decreasing pH and DO with the exception of the pH 7.3 and DO 2 mg L^{-1} treatment (Fig. 8).

Principal component analysis

Principal component analysis (PCA) showed that PC1 expressed 64.94% and PC2 13.08% of total variance after the 21-day exposure (Fig. 9). PC1 allowed to discriminate between pH 8.1 and decreased pHs and was mostly influenced by CAT, GST, GPX, GSH, MDA, and GSA. PC2 was mostly influenced by SOD activity (Fig. 10).



Fig. 5 Glutathione (GSH) activity in the gonal of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period



Fig. 6 Glutathione S-transferase (GST) activity in the gonad of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L⁻¹) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period. Letters indicate significant effects (p<0.05; post hoc test) between pH treatments within DO level and asterisks significant effects (p<0.05; post hoc test) between DO treatments within pH level

Discussion

The purpose of this study was to clarify the antioxidant response in the gonad of the marine mussel *M. coruscus* under ocean acidification and hypoxia and its connection with gonad development.

Exposure to environmental stressors modulates the dynamic balance between the generation and removal of reactive oxygen species (ROS). ROS can be removed by antioxidant enzymes mostly on the polyunsaturated fatty acids in the plasma membrane phospholipids. As a consequence, modification in the level of MDA, the product of lipid peroxidation, is an indicator of changes in ROS dynamics (Diguiseppi et al. 1984).



Fig. 7 Malondialdehyde (MDA) activity in the gonad of *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) and pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period and after a 10-day recovery period









Fig. 9 Integrative biomarker response index (IBR/n) in *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) combined with pH levels (8.1, 7.7, and 7.3) (A) and after recovery (B)

Fig. 8 IBR star plots of all tested biomarkers in *M. coruscus* exposed to different DO concentrations (6 and 2 mg L^{-1}) combined with pH levels (8.1, 7.7, and 7.3) after a 21-day exposure period (A) and after a 10-day recovery period (B)

The relative contribution of each antioxidant enzyme to protect against oxidative stress is not well known. In this study, the effect of pH, DO, and their interaction was enzyme-specific after a 21-day exposure. When significant, low pH and DO led to a decrease in SOD, CAT, and GST activity and an increase in GPX and GSH activity and MDA level.

As a vital scavenger of H_2O_2 , CAT is considered a second line of antioxidant defense (Chelikani et al. 2004). Although CAT activity often increases as a result of the increased H_2O_2 under oxidative stress (Hermes-Lima 2004), multiple stressors may change this trend (Matozzo et al. 2013). In our study, CAT activity was significantly reduced by exposure to low pH and DO. Woo et al. (2013) also found that CAT activity decreased when mussels *Mytilus galloprovincialis* were exposed to hypoxia.

GPX not only participates in the conversion of H_2O_2 to water and molecular oxygen but also plays a vital role in the use of GSH as a reducing agent to convert other lipids to nontoxic products (Sies et al. 1997). In our study, GPX activity was significantly increased under certain combinations of low pH and low DO. Lima et al. (2019) showed that the GPX activity in oyster *Crassostrea gasar* was not impacted by an exposure to a combination of low pH and phenanthrene for 96 h. Johannsson et al. (2018) found GPX activity increased in the brain and gills of Characid fish *Cyphocharax abramoides* during hypoxia. Although both GPX and CAT could catalyze the decomposition of H_2O_2 , their different changes suggested that GPX is more capable of scavenging free radicals than



Fig. 10 Biplot originating from principal component analysis integrating all measured variables (SOD, CAT, GPX, GST, GSH, MDA, GSA) in six different treatments (\blacksquare DO 6 mg L⁻¹ ×pH 8.1, \square DO 2 mg L⁻¹ ×pH 8.1, \blacklozenge DO 6 mg L⁻¹ ×pH 7.7, \diamond 2 mg L⁻¹ ×pH 7.7, \blacklozenge DO 6 mg L⁻¹ ×pH 7.3, \diamond 2 mg L⁻¹ ×pH 7.3)

CAT, which is consistent with previous research (Dorval et al. 2003). Besides acting as a substrate for GPX and glutathione reductase, GSH is also a major thiol compound that acts as a protective agent for a variety of toxic substances through thiol groups (Habig et al. 1974; Moreno et al. 2005). In our experiment, the GSH activity was increased by low pH and DO, showing a similar trend with GPX. Some studies have shown that OA and hypoxia or the combination with other stressors either decrease or increase GPX activity, depending on the species and specific tissue (Huang et al. 2018a, 2018b; Khan and Ringwood 2016). In addition to being an important regulator of detoxifying oxygen free radicals, glutathione has also been reported to be associated with reproductive success in oysters. Ringwood and Conners (2000) found that parental depletion of GSH may increase the susceptibility of embryos of the oyster Crassostrea virginica to metal toxicity. Hence, another explanation for the increase in GSH is to maintain the health state of the gonads.

GST plays different functions in the process of chemical detoxification, through which it greatly increases the basis of sulfhydryl reaction metabolites and reduces oxygen species (Aniya et al. 1993; Sheehan et al. 2001). In our experiment, the GST activity was significantly decreased under low pH and DO, indicating a disruption in the balance of GST, supporting similar observations by Lima et al. (2019).

MDA level is often measured in combination with antioxidant enzymes. The activity of antioxidant enzymes indirectly reflects the ability of the organism to scavenge oxygen free radicals, while the level of MDA indirectly reflects the severity of the attack of the organism cells by free radicals. In our results, MDA level was significantly increased under low pHs, suggesting oxidative damage.

At the end of the recovery period, enzyme activities were partially restored with no significant effects of pH, DO, or their interaction for SOD, CAT, GSH, and MDA. GST activity was only partially restored but was still significantly impacted by the exposure to low pH during the first 21 days. Exposure to low pH and DO had still significant effects on GPX after the recovery period. The consequences of these longer lasting impacts are undetermined.

Different enzymes have different sensitivity to environmental stressors. To better capture the overall stress response, the analysis of several enzymes and even other biomarkers can be combined. The IBR index is such a powerful tool. It has been widely used in many studies to assess stress responses and ecological risks (Cao et al. 2019; Damiens et al. 2007; Xie et al. 2016). In this study, exposure to low pH and DO led to a decrease in IBR/n with DO 2 mg L⁻¹ and pH 7.3 showing the highest IBR/n value, reflecting this group was the most highly impacted one. Similarly, the lowest values of GSA were observed at lowest pH and DO. When exposed to stressing conditions, mussels exhibit physiological trade-offs. Under extreme environmental pressures, they may divert energy from reproduction to costly physiological defenses (Petes et al. 2008; Béguel et al. 2013). Our results suggest that the combined stress of ocean acidification and hypoxia negatively impact GSA as a consequence of such shift in energy allocation with potential long-lasting consequences on their fitness.

Conclusion

Our results suggest that exposure to acidification and hypoxia modulates mussel's gonadal antioxidant system. The effect is enzyme-dependent and when significant effects are observed, acidification and/or hypoxia decreased the activity of CAT and GST, but increased GPX, GSH activity and MDA level. pH had a stronger effect on the antioxidant response than DO. These changes in the antioxidant system are associated with a decrease of GSA. A more specific link between the antioxidant system and gonad development is worth exploring in the future. A partial recovery of the antioxidant system of mussels exposed to low pH and DO was observed after a 10-day recovery but without positive consequences for GSA. A full recovery may take longer time and the short and long-term consequences of these changes need to be further investigated.

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Author contribution JY, MH, HK, and YD conceived the study. GX, HK, XC, and HC carried out the experiment. HK, GX, and SD analyzed

the data and wrote the manuscript. All authors read and approved the final manuscript.

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Data availability The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate Animals were used following international, national, and/or institutional guidelines.

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Competing interests The authors declare no competing interests.

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