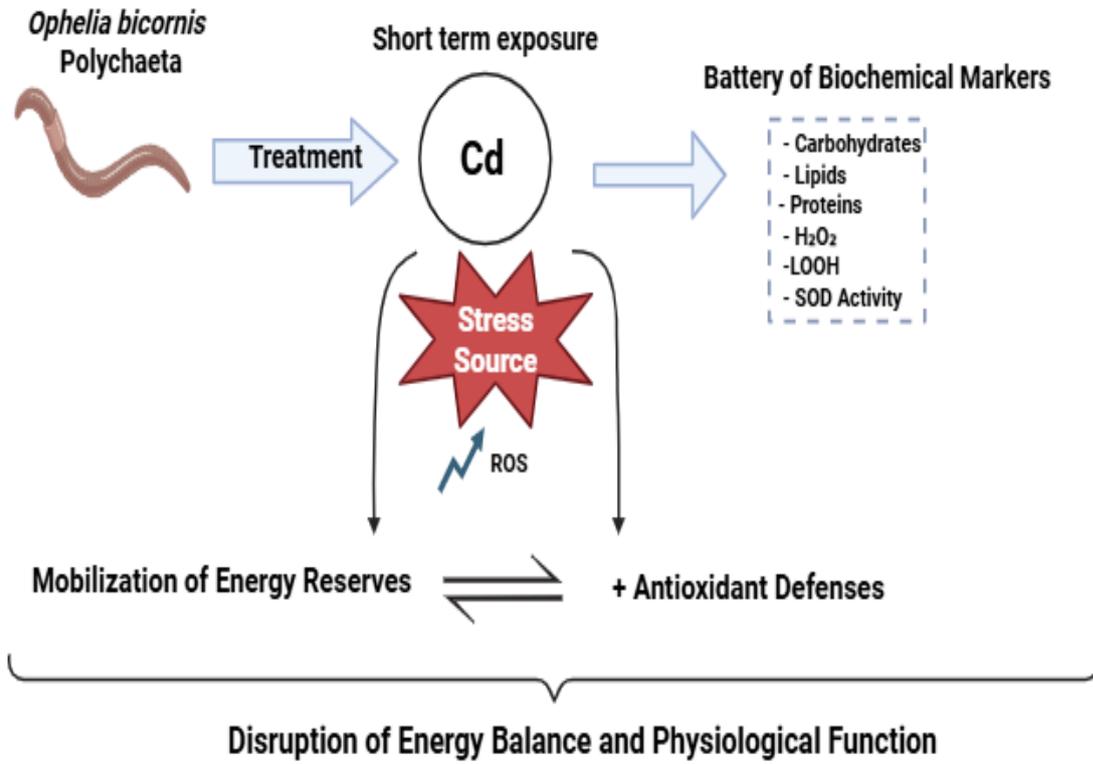


1 **Graphical abstract**

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16 **Short-term effects of cadmium exposure on energy reserves and stress markers**  
17 **on *Ophelia bicornis* Savigny, 1822 (Polychaeta: Opheliidae)**

18 **Belabed Soumeya<sup>1\*</sup>, Labiod Ryma<sup>2</sup>, Boukrouma Nadhra<sup>3</sup>, hadhbi Lazhar<sup>4</sup> and Soltani**  
19 **Noureddine<sup>1</sup>**

20 <sup>1</sup> Laboratory of Applied Animal Biology (LBAA), Department of Biology, Faculty of Sciences,  
21 Badji Mokhtar University, 12, P.O. Box, 23000 Annaba, Algeria.

22 <sup>2</sup> Laboratory of Organic Chemistry and Interdisciplinary (LOCI), University of Souk-Ahras, P.O.  
23 Box 1553, 41000 Souk Ahras, Algeria.

24 <sup>3</sup> Laboratory of Aquatic and Terrestrial Ecosystems (LEAT), Department of Biology, Faculty of  
25 Natural and Life Sciences, Mohamed Chérif Messaâdia University, BP1553, Annaba Road, 41000  
26 Souk Ahras, Algeria.

27 <sup>4</sup> Laboratory of Ecology, Biology and Physiology of Aquatic Organisms, Faculty of Sciences of  
28 Tunis, University Tunis El Manar, 2092, Tunis, Tunisia.

29 \* Correspondence: [s.belaabed@univ-soukahras.dz](mailto:s.belaabed@univ-soukahras.dz) ; phone: +213552835465

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43 **Abstract**

44 *Ophelia bicornis* (Polychaeta: Opheliidae) has been identified as a potential bioindicator for  
45 ecotoxicological studies. This study evaluates the short-term physiological and biochemical effects  
46 of cadmium (Cd) exposure on *O. bicornis* by examining energy reserves (lipids, carbohydrates,  
47 proteins) and oxidative stress biomarkers (hydrogen peroxide, lipid peroxide (LOOH), and  
48 Superoxide dismutase (SOD). Cd treatment was conducted at a sublethal concentration LC<sub>10</sub>  
49 obtained after 48h of exposure (31.11 mg/L) as previously determined. Statistical analyses revealed  
50 significant interactions between Cd treatment and exposure time affecting several biochemical  
51 parameters. Amounts of Carbohydrates increased notably after 48h in Cd-exposed individuals,  
52 indicating a time-dependent energy mobilization. Lipid contents were also affected by treatment and  
53 exposure duration, reflecting energy reserve use under metal stress. In contrast, protein remained  
54 stable, with no significant variation across treatment or time. Oxidative stress biomarkers were  
55 markedly influenced by Cd exposure. LOOH levels increased significantly with treatment and time,  
56 peaking at 48h. Similarly, H<sub>2</sub>O<sub>2</sub> increased with Cd and exhibited a significant time-dependent  
57 interaction, highlighting a progressive oxidative challenge. SOD activity was primarily modulated  
58 by exposure duration, reaching its maximum at 48h. Overall, the present experiment shows that  
59 sublethal Cd disrupts energy metabolism and triggers oxidative stress response in *O. bicornis*.

60 **Keywords:** *Ophelia bicornis*, Bioindicator, Toxicity, Energy metabolism, Oxidative stress,  
61 Biomarkers.

62 **1. Introduction**

63 In recent years, it has become evident that, to fully assess the impact of anthropogenic trace metal  
64 inputs on aquatic ecosystems, it is essential to understand the chemical and physiological processes  
65 that govern metal accumulation in organisms, as well as how these metals influence energy reserves  
66 and antioxidant activities. Many toxicants produce a sublethal effect by binding to or interfering  
67 with essential cellular components such as enzymes and metabolites, thereby disrupting

68 fundamental processes within the organism (Holmstrup *et al.* 2010). They can disrupt energy  
69 metabolism, deplete energy stores, and induce oxidative stress, ultimately affecting overall health  
70 and survival of aquatic organisms (Song *et al.* 2023). These substances are particularly known for  
71 their toxic effects, persistence, and tendency to transfer and accumulate through all the trophic  
72 levels (Di Salvatore *et al.* 2013). Trace metals are among the most extensively studied groups of  
73 xenobiotics (Pan and Wang 2012). Besides, various chemical substances, including cadmium, can  
74 exert multiple mechanisms of toxicity, each contributing to the overall harmful effects (Valavanidis  
75 *et al.* 2006; Regoli and Giuliani 2014). Cadmium (Cd) is a heavy metal hazardous to all living  
76 organisms, and its effects have been studied on various organisms (Sharma *et al.* 2015). It is a non-  
77 essential element that can be toxic even at trace levels for aquatic species. Cadmium is a highly  
78 toxic environmental pollutant that poses a significant threat due to its tendency to accumulate in  
79 marine ecosystems, its ability to easily cross cell membranes, and its persistence in the  
80 environment, which can be found at high concentrations in seawater (Amira *et al.* 2018).  
81 Polychaetes play an important role in the functioning of benthic communities (Hutchings 1998).  
82 They represent 35 to 50% of benthic macrofauna and are often the numerically dominant  
83 macrobenthic taxon. Found in coastal and littoral ecosystems, where the distribution of species in  
84 soft-bottom habitats is primarily influenced by sediment particle size (Gambi and Giangrande 1986)  
85 and organic matter content, they are frequently exposed to various contaminants. Due to their  
86 sensitivity and ecological significance, polychaetes serve as excellent indicators for studying the  
87 effects of pollutants and the transfer of xenobiotics through food chains (Dean 2008). Consequently,  
88 they have been extensively used in coastal studies for environmental monitoring and are considered  
89 a key group in ecotoxicological research (Scaps 2000). Furthermore, Polychaete Annelids are a  
90 major link in food webs and are widely consumed by many species, such as crustaceans, fishes, and  
91 birds. Among them, the endobenthic species *Ophelia bicornis* (Savigny 1818) plays a crucial role in  
92 evaluating the structure and functioning of estuarine ecosystems. This polychaete is particularly  
93 notable for its physiological tolerance to various environmental stressors, especially its resistance to

94 sediment-bound contaminants. Accordingly, reduced energy reserves and elevated metabolic rates  
95 in animals have associated to metal toxicity, especially in the case of cadmium, Zinc and copper  
96 (Pook *et al.* 2009). Recent studies have explored advanced analytical and computational approaches  
97 for environmental pollution assessment. Deep learning and optimization-based models have been  
98 applied to classify and predict air pollution levels and analyze environmental data (Mohandas *et al.*  
99 2025b, 2025a). Additionally, hybrid recurrent neural network models combined with Internet of  
100 Things (IoT) data have been proposed for urban air pollution prediction (Mohandas *et al.* 2025c).  
101 Beyond atmospheric pollution, graph-based and remote sensing-driven approaches have been  
102 developed for environmental monitoring (Sivasubramanian *et al.* 2025), and early detection of  
103 ecological stress, such as pest disease identification in agricultural systems (Maruthai *et al.* 2025;  
104 Mohandas *et al.* 2025a). While these studies provide valuable large-scale insights, experimental  
105 ecotoxicological investigations remain essential for understanding species-specific biochemical  
106 responses to metal exposure. *Ophelia bicornis*, which commonly inhabits organic-rich  
107 environments, is considered a key species for ecological assessments and a reliable bioindicator for  
108 monitoring heavy metal pollution (Bat *et al.* 2016). In this context, *Ophelia bicornis* could be  
109 effectively used as a bioindicator species in ecotoxicological studies of marine and coastal  
110 ecosystems along the eastern Algerian coast to assess metal contamination (Labiod and Belabed  
111 2024). This species is distributed across the North, Mediterranean, Black Sea, and Atlantic coasts of  
112 North Africa and Europe, typically inhabiting high-energy intertidal zones and fine to medium  
113 sediments (Maltagliati *et al.* 2005).

114 The presence of a xenobiotic in the environment always poses a risk to living organisms. However,  
115 the relationship between toxic concentrations and the toxic response is fairly complex and difficult  
116 to predict, as it depends on several factors, including toxicokinetics and genetic variability. One  
117 method of quantifying xenobiotic interaction and its potential impact on living organisms is through  
118 biomarker monitoring. Therefore, biomarkers measure exposure, toxicity, and individual  
119 susceptibility to environmental chemicals, making them valuable tools for assessing and monitoring

120 the risk of long-term effects associated with exposure to xenobiotics(Depledge 2020). Generally,  
121 biochemical biomarkers are frequently used to identify early signs of stress or toxicity in benthic  
122 fauna (Mayer *et al.* 2018; Depledge 2020). One of the main consequences of metal toxicity is  
123 additional energy costs, and the resulting metabolic load can lead to disruption in oxidative  
124 metabolism and increased anaerobic activity (Gashkina 2024). Furthermore, metal toxicity mainly  
125 involves changes at biological levels of cellular activities, including reactive oxygen species (ROS)  
126 generation, which can cause damage associated with the different pathological processes  
127 (Makhdoumi *et al.* 2020). Likewise, Cd accumulation has also been reported to alter key  
128 biochemical and physiological functions (Wright and Welbourn 1994)and to induce the production  
129 of reactive oxygen species (ROS) such as hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and lipid peroxide (LOOH)  
130 (Benedetti *et al.* 2015). A significant increase in the tissue concentration appeared 48h after  
131 exposure in *Donax trunculus*, following treatment with LC<sub>50</sub> (2.59 mg/L) and LC<sub>25</sub> (1.32 mg/L) of  
132 Cd (Belabed and Soltani 2018). Moreover, the presence of cadmium chloride (CdCl<sub>2</sub>) in seawater  
133 led to mortality in the marine benthic species *Donax trunculus* at concentrations corresponding to  
134 LC<sub>10</sub> (0.72 mg/L) and LC<sub>50</sub> (2.59 mg/L) after 96h of exposure, and also affected the biomarker  
135 responses such as acetylcholinesterase (AChE) and Catalase (CAT) activities (Belabed and Soltani  
136 2022). Thus, some biomarkers such as Superoxide dismutase, catalase, glutathione peroxidase,  
137 glutathione, and glutathione reductase are indicative of oxidative stress caused by exposure to pro-  
138 oxidants (Demirci-Çekiç *et al.* 2022), whereas metallothionein contents are widely used as  
139 biomarkers of metal contamination by binding and removing toxic metals (Cosson and Amiard  
140 2021). These antioxidant defenses aim to protect cells and tissues from oxidative damage by  
141 neutralizing the toxicity of ROS (Singh *et al.* 2024).

142 In previous studies (Belabed and Soltani 2018, 2022; Merad *et al.* 2024), Cd was tested on a  
143 Molluscan species, *Donax trunculus*, widely used as bioindicator of marine pollution. The present  
144 study aimed to investigate the short-term effects (24, 48h) of cadmium exposure at a sublethal  
145 concentration LC<sub>10</sub> (31.11 mg/L) (Labioud and Belabed 2024) in the marine organism *O. bicornis*, to

146 assess the extent of metal contamination. Acute toxicity data on *O. bicornis* and its responses to  
147 toxic substances remain scarce in the scientific literature. In light of the limited data available, a  
148 battery of stress indices was employed, including SOD activity, the generation of free radicals, and  
149 the biochemical composition. Characterization of the response of stress indices could determine the  
150 possible adverse effects of cadmium on worms and evaluate the potential harmful impacts on  
151 aquatic environments.

## 152 **2. Materials and methods**

### 153 *2.1 . Sampling and contamination protocol*

154 The annelid *O. bicornis* was collected at low tide by hand from littoral sediments in the intertidal  
155 zone (depth greater than 20 cm) at El Hennaya Beach (36°54'24.00" N, 08°7'43.62" E), that is  
156 located in the district of Berrihane, which belongs to the province of El Tarf (Northeast  
157 Algeria). This site is not exposed to any source of pollution because of its location, which is quite  
158 remote from the various discharges. Once in the laboratory, the samples were transported in plastic  
159 buckets with sediment from the collection site, and they were acclimatized for 48 hours.  
160 Afterwards, the collected worms were separated from the residue, cleaned from debris with  
161 seawater from the site of origin of worms, and they were transferred to the glass Petri dishes  
162 containing filtered natural seawater for another period of acclimatization (24h) in the dark at a  
163 temperature of 16°C. This acclimatization phase establishes stable experimental conditions and  
164 enhances the accuracy of toxicity test results, reflecting the actual effect of the toxic substance on  
165 organisms. Sublethal concentrations of Cd (LC<sub>10</sub>) were administered to lots (n = 30 for each  
166 treatment). Thus, every 24 hours throughout the exposure period, worms (0.3 g in weight) were  
167 sampled and immediately homogenized (10% w/v) in Tris-HCl buffer (100 mM; pH = 7.4). The  
168 homogenates were centrifuged at 9000×g for 20 min at 4 °C. The supernatant was then preserved at  
169 -80°C until biochemical analysis. Analyses were performed to assess the total lipid, carbohydrate,  
170 and protein contents as well as oxidative stress responses by measuring Superoxide dismutase  
171 (SOD) activity and quantifying Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and Lipid hydroperoxides (LOOHs)

172 levels. Assays were conducted on six individuals in both the treated and control groups, spanning a  
173 treatment period of 48 hours.

#### 174 2.2 . *Energy reserve analysis*

175 In this experimental procedure, biochemical analyses concerned only the animals that survived the  
176 two days of Cd exposure. The extraction of protein, carbohydrate and lipid contents is determined  
177 according to Shibko *et al.*(1966). The method involves treating a homogenate by adding 1 ml of  
178 20% trichloroacetic acid (TCA) to the sample. After grinding with ultrasound (Sonifier B-30),  
179 centrifugation at  $5000 \times g$  for 10 minutes at  $4^{\circ}C$  yields a supernatant 1 that is collected use in the  
180 total carbohydrate analysis, while the pellet 1 undergoes washing with 1 ml chloroform ether  
181 (1V/1V), centrifuged again ( $5000 g$  for 10 min). To determine the carbohydrate content of the  
182 supernatant fraction, 4 ml of anthrone was added to 100  $\mu$ l of supernatant, and quantification was  
183 performed by measuring absorbance at 620 nm against a standard curve of glucose standard (Sigma,  
184 USA) (Duchateau and Florkin 1959). The supernatant 2 recovered will be used for the  
185 determination of the total lipids based on the vanillin method of (Goldsworthy *et al.* 1972) and  
186 absorbance of the samples was measured at 530 nm. The remaining pellet 2 was re-suspended in  
187 1ml NaOH (0,1 N) used for assessing the total protein content (Bradford, 1976a) (Sigma, USA).  
188 The absorbance was measured at 595 nm using bovine serum albumin (Sigma, USA) as a standard.  
189 The contents of carbohydrates, proteins and lipids were expressed in  $\mu$ g/mg of body weight.

#### 190 2.3 . *Superoxide dismutase (SOD) activity*

191 The superoxide dismutase (SOD) activity was determined by monitoring the photochemical  
192 reduction of NBT according to the method of Beauchamp and Fridovich (1971). One unit of SOD  
193 activity corresponded to the amount of enzyme required to cause 50% inhibition of NBT reduction  
194 at 560 nm measured by a spectrophotometer. The SOD activity was expressed as U/mg of protein.

#### 195 2.4 . *Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>)*

196 Analysis of hydrogen peroxyde ( $H_2O_2$ ) levels were carried out by the method described by Ou and  
197 Wolff (1996), using the ferrous ion oxidation xylenol orange assays (FOX1). The amount of  $H_2O_2$   
198 in the supernatant was determined at 560 nm using a spectrophotometer. Values were expressed as  
199 mmol/mg of protein.

#### 200 2.5 . Lipid hydroperoxides (LOOHs) content

201 Lipid hydroperoxides (LOOH) formation was estimated according to the method described by Jiang  
202 *et al.* (1992) using the ferrous oxidation in xylenol orange assay. Hydroperoxides content was  
203 colorimetrically determined at 560 nm. Results were expressed as mmol/ mg of protein.

#### 204 2.6 . Total Protein quantification

205 Protein content was done by the method of Bradford (1976), which is based on the binding of  
206 Coomassie brilliant blue G -250, present in the Bradford reagent, to the total protein present in the  
207 sample, yielding a stable staining compound. The presence of this compound can be  
208 spectrophotometrically quantified at 595 nm.

#### 209 2.7 . Statistical analysis

210 All statistical analyses were conducted using R software (version 4.5.0; R Core Team 2025). Prior  
211 to hypothesis testing, data normality and homogeneity of variances were assessed using the  
212 Shapiro–Wilk test and Levene’s test, respectively. For carbohydrate content, which met normality  
213 and homogeneity assumptions ( $p > 0.05$ ), two-way ANOVA was applied to evaluate the effects of  
214 Exposure Time, Treatment, and their interaction. Significant effects were followed by Tukey’s HSD  
215 post hoc test for pairwise comparisons while controlling for multiple testing. For all other measured  
216 biomarkers (lipids, proteins, LOOH,  $H_2O_2$ , and SOD), which did not meet normality assumptions,  
217 the non-parametric Scheirer–Ray–Hare test was used as a factorial alternative to ANOVA, with  
218 significant effects further analyzed using Dunn’s test for multiple comparisons. Exploratory data  
219 analysis was performed to visually assess variable distributions across experimental groups defined  
220 by Exposure Time and Treatment. Boxplots summarizing median, interquartile range, overall

221 dispersion, and potential outliers were generated using the *ggplot2* package (version 3.4.2;  
222 Wickham 2016), supporting the evaluation of model assumptions. Bar plots with error bars (mean  $\pm$   
223 SD) were used to facilitate visual comparison of central tendencies and variability. To explore  
224 overall variation patterns and relationships among biomarkers, Principal Component Analysis  
225 (PCA) was performed using the *FactoMineR* package (version 2.7; Lê *et al.* 2008), and results were  
226 visualized with the *factoextra* package (version 1.0.7; Kassambara and Mundt 2020). Redundancy  
227 Analysis (RDA) via the *vegan* package (version 2.6-4; Oksanen *et al.* 2022) quantified the  
228 proportion of variance in biomarker responses explained by Exposure Time and Treatment. All  
229 results are presented as mean  $\pm$  standard deviation (SD), and the number of biological replicates  
230 used in each assay is reported in the Results section. Statistical significance was set at  $p < 0.05$ .

### 231 3. Results

#### 232 3.1. Changes in the energy reserves

233 All energy reserve components exhibited large variations. Generally, changes in energy reserve  
234 responses are detected after prolonged exposure, over several hours or days. Proteins were  
235 quantitatively the most important energy fractions in the biochemical composition of *O. bicornis*  
236 (28,88  $\mu\text{g}/\text{mg}$  of tissue), whereas carbohydrates and lipids represented respectively only 3,93 and  
237 1,93  $\mu\text{g}/\text{mg}$  of tissue.

##### 238 3.1.1. Carbohydrates contents

239 The values ( $\mu\text{g}/\text{mg}$  of Tissue) obtained in *Ophelia bicornis* are presented in figure 1. The two-way  
240 ANOVA revealed a marginally significant interaction between the factors Time and Treatment ( $p =$   
241 0.0569), suggesting that the effect of the treatment on Carbohydrates contents depends on the  
242 duration of exposure. No significant main effects were detected for Time ( $p = 0.0672$ ) or for  
243 Treatment ( $p = 0.1534$ ), indicating that these factors alone do not account for the observed  
244 variations (Figure 1, A). However, the interaction between these two factors highlights a combined  
245 influence on the measured response.

246 Post hoc analysis using Tukey's test showed a significant increase in Carbohydrates after 48 hours  
247 in the Cd treated series compared to the initial time point (0 hours) ( $p = 0.0287$ ), indicating a Time-  
248 dependent effect of the treatment (Figure 1, B). In contrast, no significant differences were observed  
249 within the controls group over time, confirming that the variations in concentration over time are  
250 primarily attributable to the treatment.

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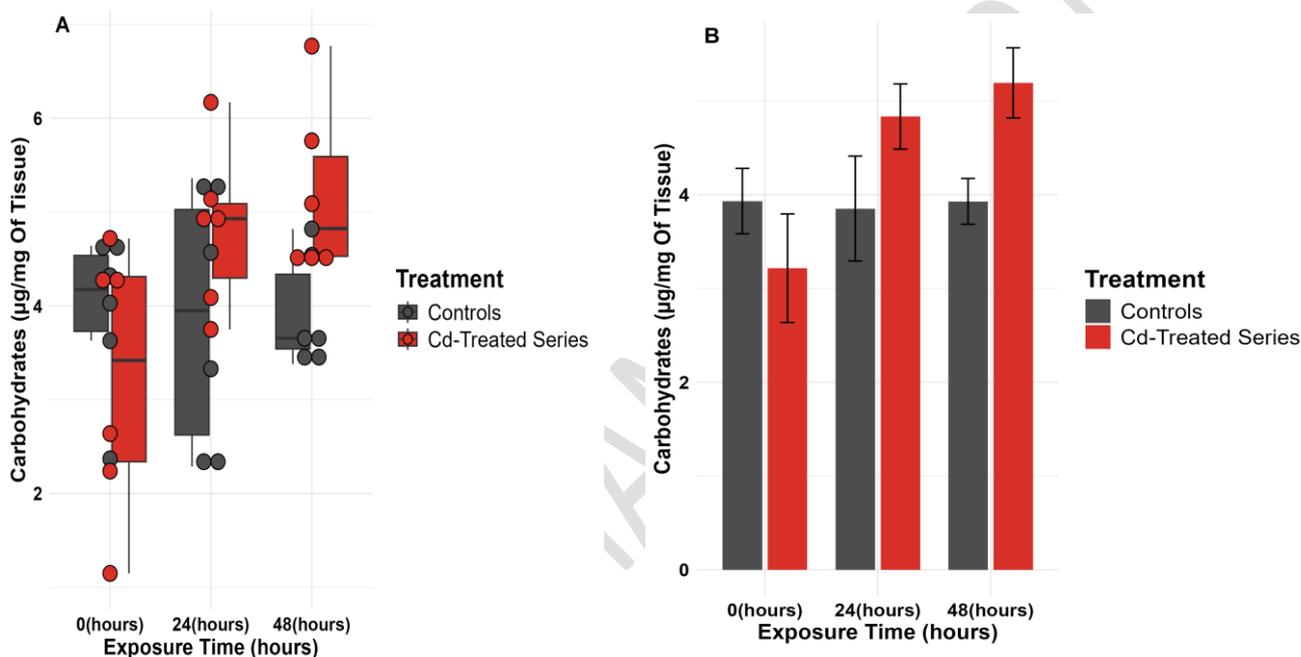
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259 **Figure 1.** Effect of cadmium on carbohydrate content ( $\mu\text{g}/\text{mg}$  Of Tissue) in *O. bicornis* ( $n = 6$ ).

260 (A) Boxplot showing mean carbohydrate content after exposure to cadmium (mean  $\pm$  SD).

261 (B) Error bar plot representing the standard deviation of carbohydrate content (mean  $\pm$  SD).

### 262 3.1.2. Lipids contents

263 The Scheirer-Ray-Hare test, was used to assess the effects of Exposure Time and Treatment on  
264 Lipid contents ( $\mu\text{g}/\text{mg}$  of Tissue) in 36 observations. The test revealed a statistically significant  
265 effect of Treatment ( $H = 4.91$ ,  $p = 0.0267$ ) and a significant Time  $\times$  Treatment interaction ( $H = 7.26$ ,  
266  $p = 0.0265$ ), suggesting that the impact of treatment depends on exposure duration (Figure 3).  
267 However, the main effect of Time alone was not statistically significant ( $H = 4.51$ ,  $p = 0.1048$ )  
268 (Figure 2, C).

269 Post-hoc Dunn tests with Bonferroni correction showed a significant difference in lipids between  
270 Cd-treated series and Control groups ( $Z = -2.22$ ,  $p = 0.0267$ ), while no significant pairwise  
271 differences were found between time points, including between 24 and 48 hours (*adjusted p* =  
272 0.1197) (Figure 2, D).

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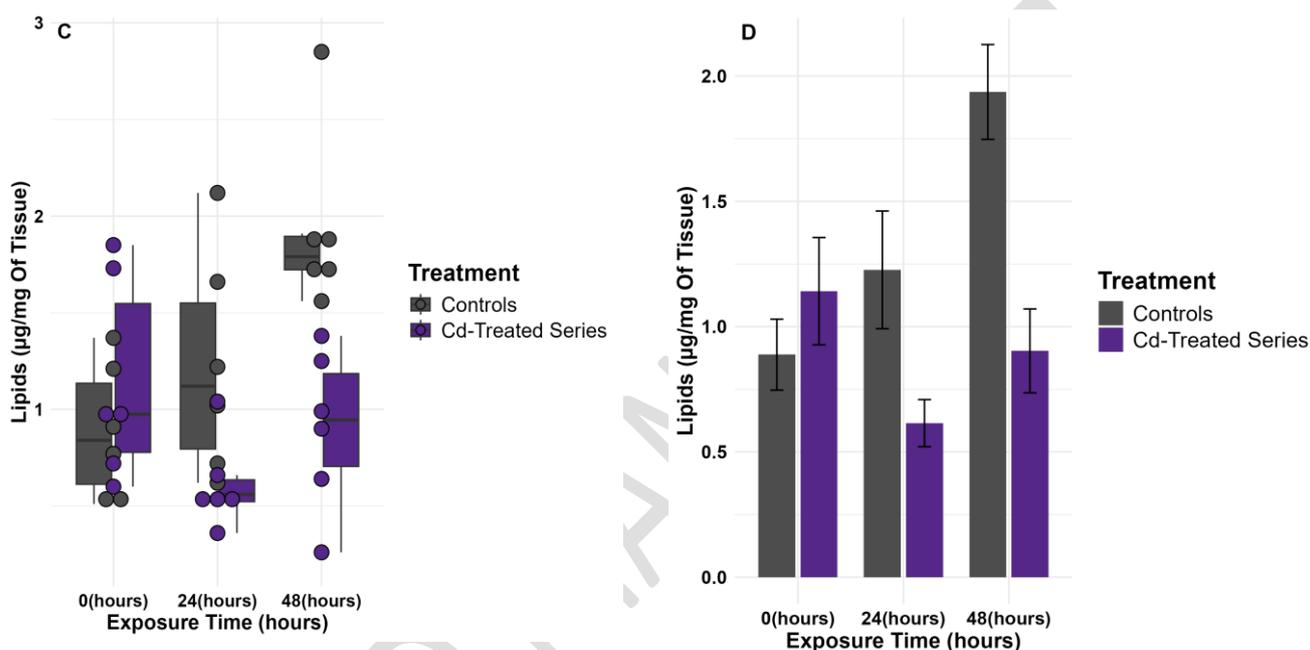
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281 **Figure 2.** Effect of cadmium on lipid content (µg/mg Of Tissue) in *O. bicornis* (n = 6).

282 (C) Boxplot showing mean lipid content after exposure to cadmium (mean ± SD).

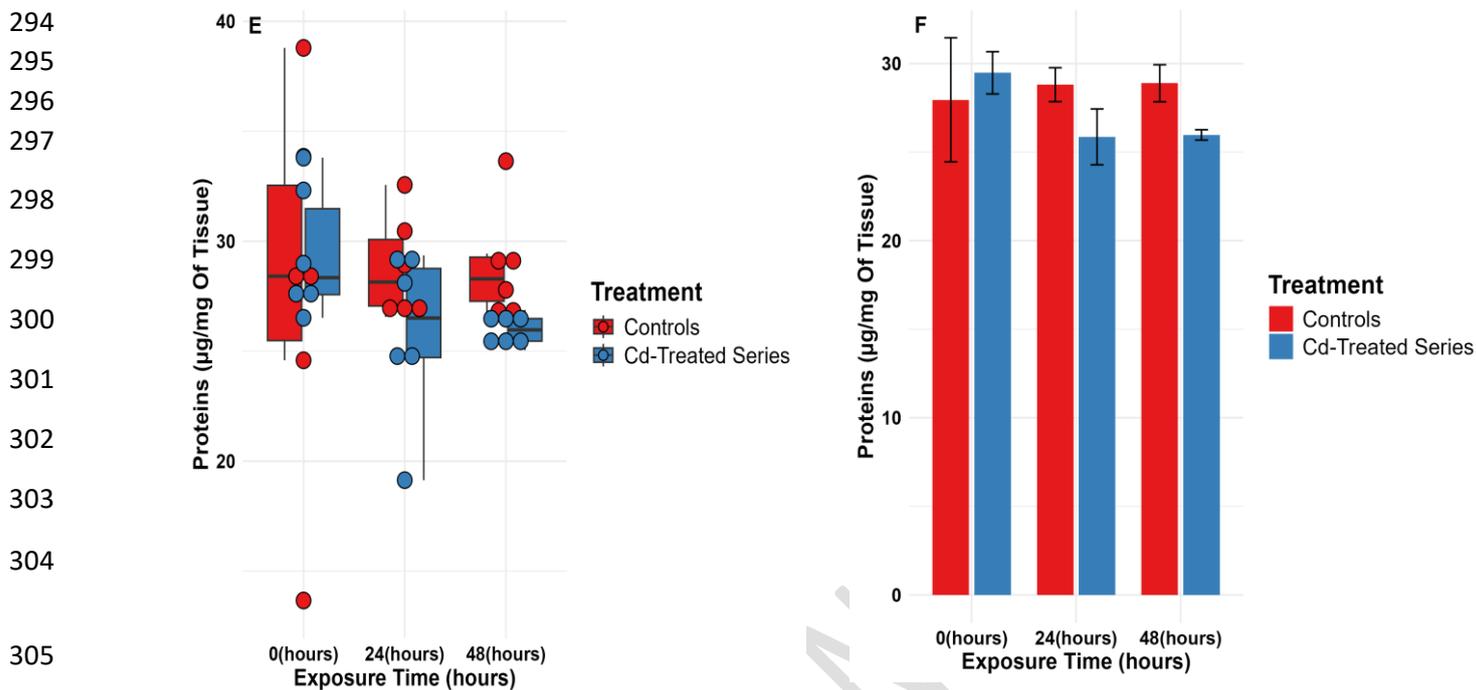
283 (D) Error bar plot representing the standard deviation of lipid content (mean ± SD).

### 284 3.1.3. Proteins contents

285 The Scheirer-Ray-Hare test, was used to examine the effects of Time, Treatment, and their  
286 interaction on Proteins contents (µg/mg of Tissue) across 36 observations. The results showed that  
287 none of the factors reached statistical significance at the 0.05 level (Figure 3, E). The effect of Time  
288 on Proteins (µg/mg of Tissue) was not significant ( $H = 1.7643$ ,  $p = 0.41388$ ), indicating that  
289 Protein's expression did not vary significantly across the different Time points.

290 Similarly, the Treatment factor was also not significant ( $H = 2.6043$ ,  $p = 0.10658$ ), suggesting no  
291 strong evidence that the treatment alone influenced protein levels. The interaction between Time

292 and Treatment was likewise non-significant ( $H = 3.5013$ ,  $p = 0.17366$ ), meaning there was no  
293 combined effect of Time and Treatment on Proteins ( $\mu\text{g}/\text{mg}$  of Tissue) (Figure 3, F).



306 **Figure 3.** Effect of cadmium on protein content ( $\mu\text{g}/\text{mg}$  Of Tissue) in *O. bicornis* ( $n = 6$ ).

307 **(E)** Boxplot showing mean protein content after exposure to cadmium (mean  $\pm$  SD).

308 **(F)** Error bar plot representing the standard deviation of protein content (mean  $\pm$  SD).

#### 309 3.1.4. $\text{H}_2\text{O}_2$ levels (mmole/mg of Proteins)

310 The Scheirer-Ray-Hare test revealed a marginally non-significant effect of Time on  $\text{H}_2\text{O}_2$  Levels  
311 (mmole/mg of Proteins) ( $H = 5.39$ ,  $p = 0.0676$ ), a significant effect of Treatment ( $H = 8.02$ ,  $p =$   
312  $0.0046$ ), and a significant Time  $\times$  Treatment interaction ( $H = 9.33$ ,  $p = 0.0094$ ) (Figure 4, G).

313 Post-hoc Dunn tests with Bonferroni correction showed that  $\text{H}_2\text{O}_2$  Levels (mmole/mg of Proteins)  
314 were significantly higher in the Cd-treated group compared to controls ( $p = 0.0046$ ). Regarding  
315 Time, no pairwise comparisons reached significance after correction, although the difference  
316 between 0 and 48 hours approached significance ( $p = 0.0945$ ). These results indicate that treatment  
317 and the interaction between Time and Treatment significantly influence  $\text{H}_2\text{O}_2$  levels (mmole/mg of

318 Proteins), with Cd treatment increasing levels overall and a time-dependent variation depending on  
319 treatment (Figure 4, H).

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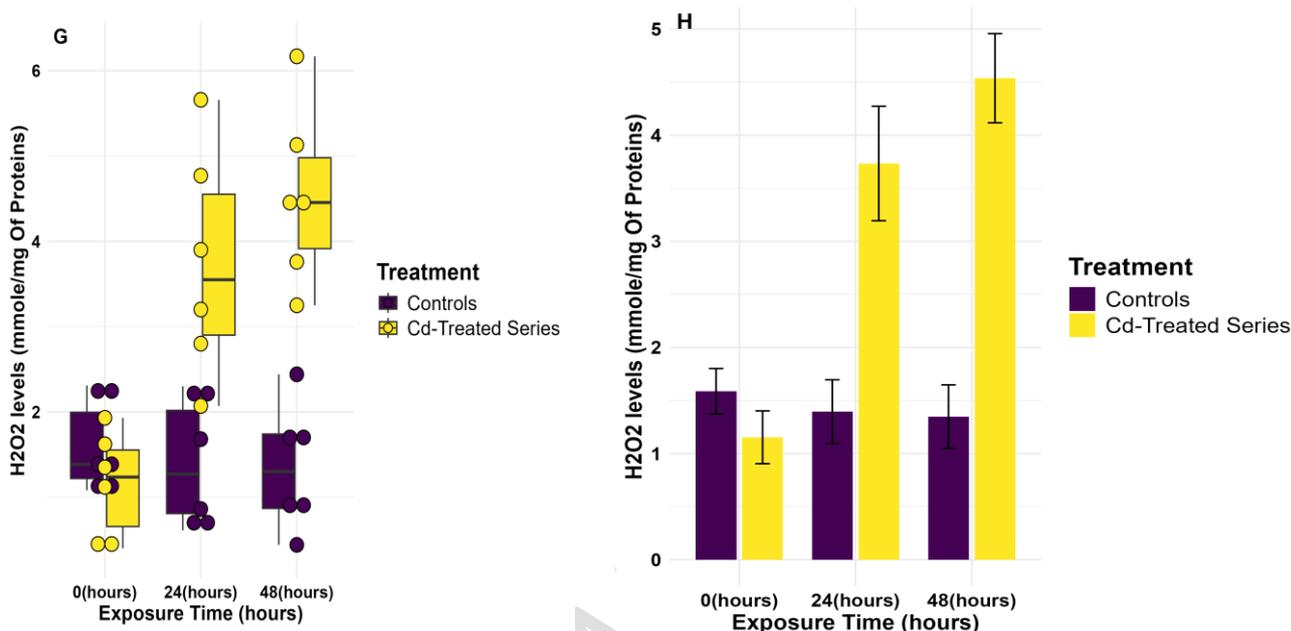
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339 **Figure 4.** Effect of cadmium on H<sub>2</sub>O<sub>2</sub> levels (mmole/mg Of Proteins) in *O. bicornis* (n = 6).

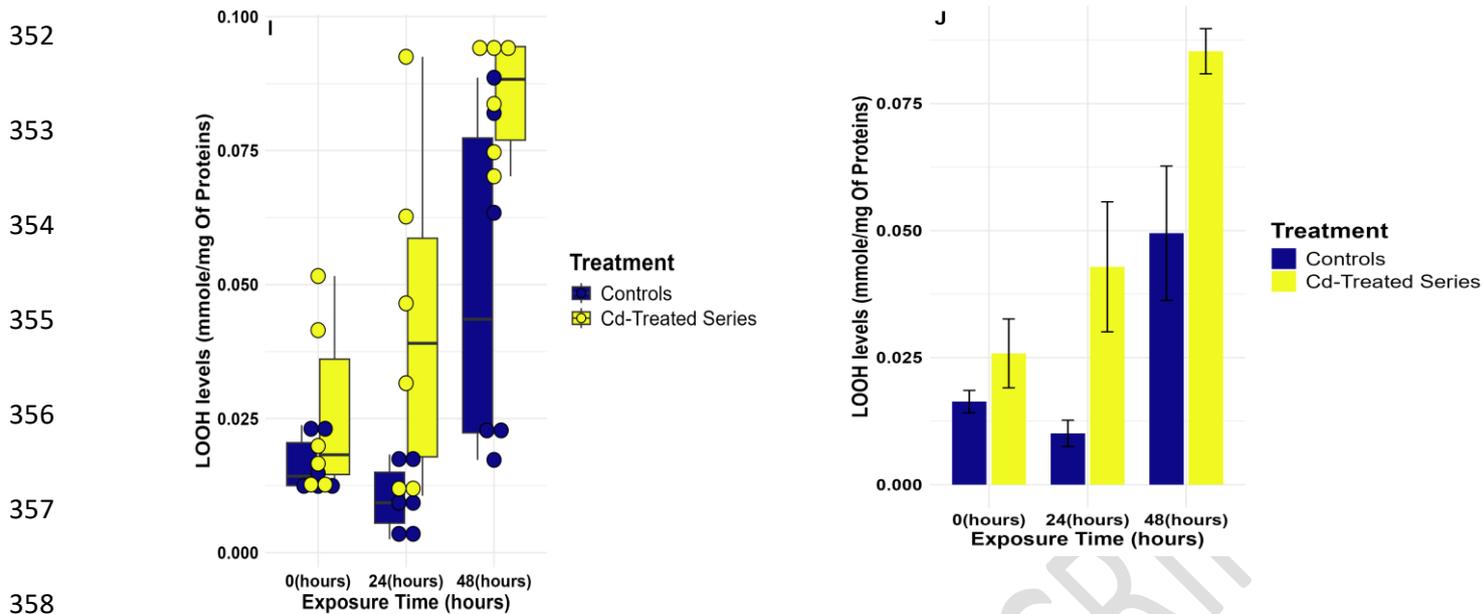
340 **(G)** Boxplot showing mean H<sub>2</sub>O<sub>2</sub> levels after exposure to cadmium (mean ± SD).

341 **(H)** Error bar plot representing the standard deviation of H<sub>2</sub>O<sub>2</sub> levels (mean ± SD).

### 342 3.1.5. LOOH levels (mmole/mg of Proteins)

343 The Scheirer-Ray-Hare test showed significant effects of Time ( $H = 14.45$ ,  $p = 0.00073$ ) and  
344 treatment ( $H = 5.71$ ,  $p = 0.01690$ ) on LOOH levels (mmole/mg of Proteins), while the interaction  
345 between Time and Treatment was not significant ( $H = 1.31$ ,  $p = 0.52$ ) (Figure 5, I).

346 Post-hoc Dunn tests with Bonferroni correction revealed that LOOH levels (mmole/mg of Proteins)  
347 at 48 hours were significantly higher than at 0 hours ( $p = 0.0049$ ) and 24 hours ( $p = 0.0019$ ), with  
348 no difference between 0 and 24 hours. Additionally, Cd-treated samples showed significantly higher  
349 LOOH levels (mmole/mg of Proteins) compared to controls ( $p = 0.0169$ ). These results indicate that  
350 both Time and Treatment independently affect LOOH levels (mmole/mg of Proteins), with  
351 increases observed at 48 hours and following Cd exposure (Figure 5, J).



359 **Figure 5.** Effect of cadmium on LOOH levels (mmole/mg Of Proteins) in *O. bicornis* (n = 6).

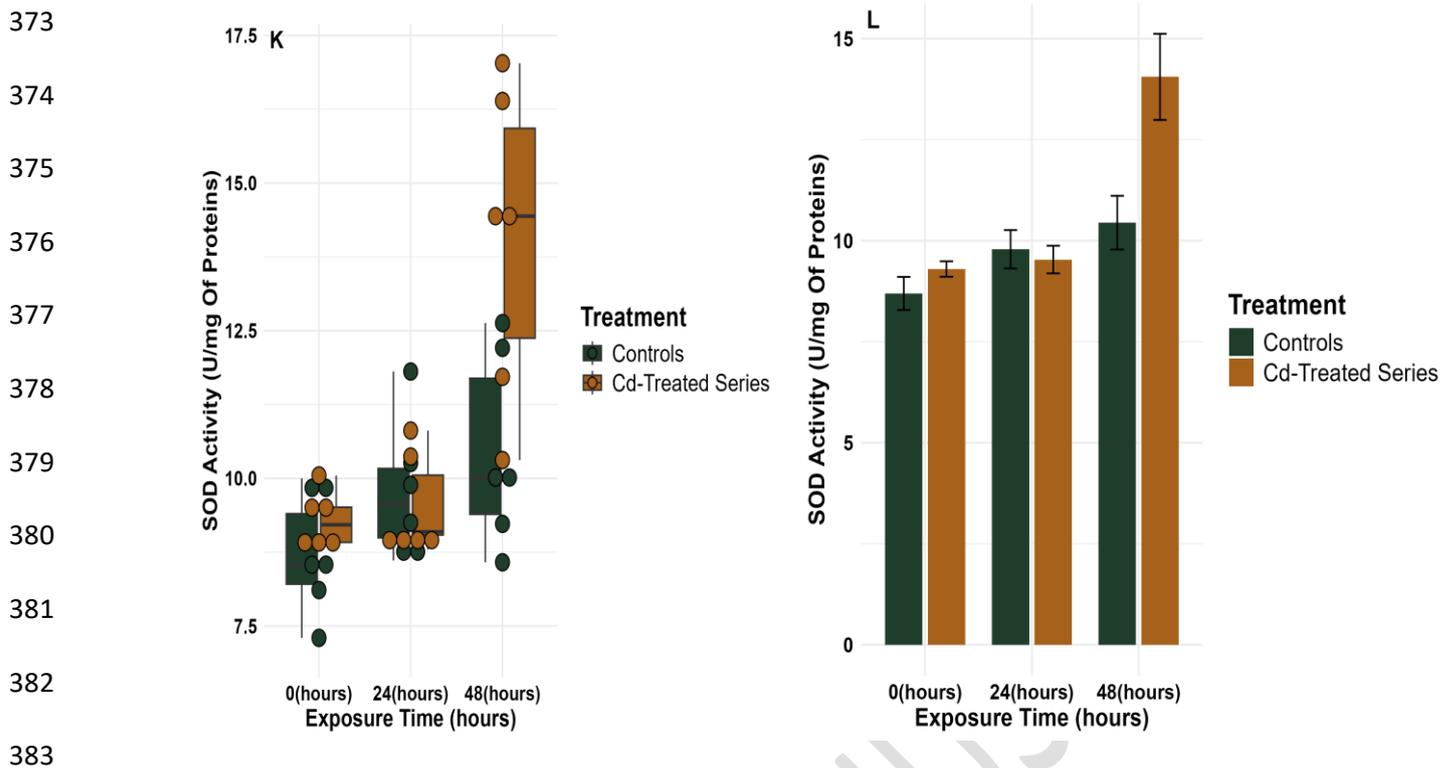
360 **(I)** Boxplot showing mean LOOH levels after exposure to cadmium (mean  $\pm$  SD).

361 **(J)** Error bar plot representing the standard deviation of LOOH levels (mean  $\pm$  SD).

### 362 3.1.6. SOD Activity (U/mg of Proteins)

363 The Scheirer-Ray-Hare test showed a significant effect of Time on SOD Activity (U/mg of Proteins)  
 364 (H = 12.71, p = 0.00174), while neither Treatment (H = 2.26, p = 0.13286) or the Time  $\times$  Treatment  
 365 interaction (H = 1.99, p = 0.36975) were significant (Figure 6, K).

366 Post-hoc Dunn tests with Bonferroni correction indicated that SOD activity at 48 hours was  
 367 significantly different from 0 hours (p = 0.0012), while differences between 0 and 24 hours and  
 368 between 24 and 48 hours were not statistically significant after adjustment (p = 0.51 and p = 0.09,  
 369 respectively). No significant difference was found between Cd-treated series and control groups (p  
 370 = 0.13). These results suggest that SOD Activity (U/mg of Proteins) changes significantly over  
 371 Time, particularly between 0 and 48 hours, but is not significantly affected by treatment or the  
 372 interaction of treatment and Time (Figure 6, L).

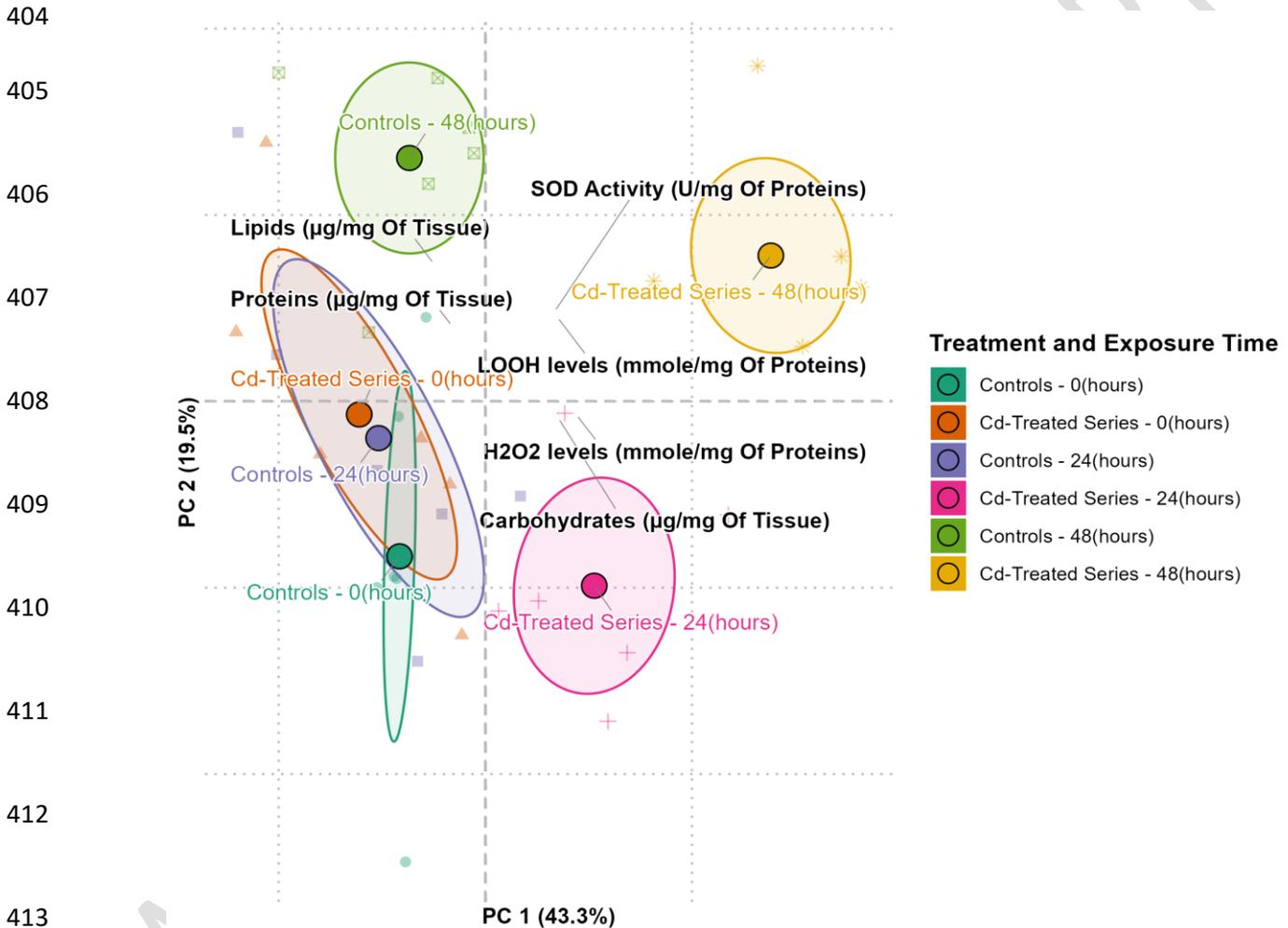


384 **Figure 6.** Effect of cadmium on SOD activity (U/mg Of Proteins) in *O. bicornis* (n = 6).  
 385 **(K)** Boxplot showing mean SOD activity after exposure to cadmium (mean  $\pm$  SD).  
 386 **(L)** Error bar plot representing the standard deviation of SOD activity (mean  $\pm$  SD).

388 *3.2. Principal Component Analysis (PCA) and Redundancy Analysis (RDA) of Biochemical*  
 389 *Biomarkers in Ophelia bicornis: Effects of Exposure Time and Treatment*

390 The combined application of PCA and RDA provided complementary insights into the biochemical  
 391 responses to treatment and exposure time. PCA, an unconstrained ordination method, revealed that  
 392 the first two principal components explained 62.82% of the total variance (PC1: 43.3%; PC2:  
 393 19.5%). PC1 was showing strong positive correlations with H<sub>2</sub>O<sub>2</sub> levels (r = 0.89, p < 0.001),  
 394 Carbohydrate (r = 0.71, p < 0.001), and LOOH levels (r = 0.70, p < 0.001). This axis was positively  
 395 associated with the Cd-treated group, particularly at 48 hours. PC2 was mainly correlated with  
 396 Lipid (r = 0.74, p < 0.001) and SOD activity (r = 0.49, p = 0.003) and was significantly associated  
 397 with the 48-hour Exposure Time (Figure 7).

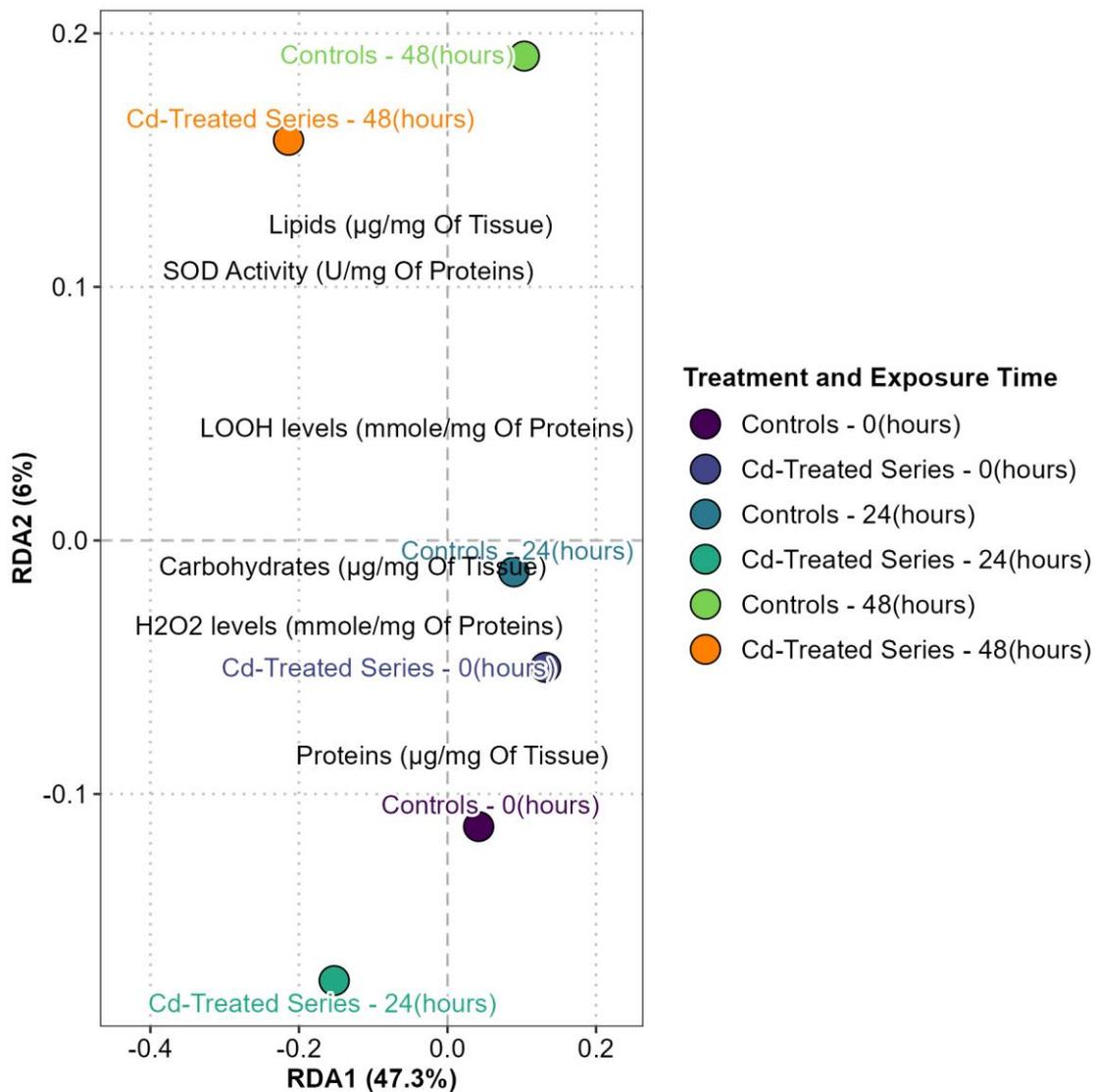
398 In parallel, RDA (constraining the ordination by Treatment and Exposure Time explained 47.3%  
 399 and 6% of the variance on Axis 1 and Axis 2, respectively. Axis 1 was positively associated with  
 400 H<sub>2</sub>O<sub>2</sub> levels ( $r = 0.33$ ,  $p < 0.001$ ) and SOD activity ( $r = 0.12$ ,  $p < 0.01$ ), and negatively with protein ( $r$   
 401  $= -0.19$ ) and lipid ( $r = -0.08$ ), indicating increased oxidative stress in exposed samples. Axis 2 was  
 402 primarily correlated with Lipid ( $r = 0.13$ ,  $p < 0.01$ ) and SOD activity ( $r = 0.09$ ,  $p < 0.05$ ), reflecting  
 403 Time-dependent metabolic responses (Figure 8).



414 **Figure 7.** Principal Component Analysis (PCA) Biplot of Biochemical Biomarkers in Relation to  
 415 Treatment and Exposure Time in *Ophelia bicornis*. **Legend:** The PCA Biplot illustrates the distribution of  
 416 biochemical parameters ( $n = 6$ ) measured in Tissue samples exposed to Cd-Treated series and control conditions over  
 417 varying Exposure Times. Variables include Carbohydrates contents ( $\mu\text{g}/\text{mg}$  Of Tissue), Lipids ( $\mu\text{g}/\text{mg}$  Of Tissue),  
 418 Proteins ( $\mu\text{g}/\text{mg}$  Of Tissue), H<sub>2</sub>O<sub>2</sub> levels (mmole/mg Of Protein), LOOH levels (mmole/mg Of Protein), and SOD  
 419 activity (U/mg Of Protein). These variables are represented as vectors, where their direction and length indicate the

420 magnitude and contribution to the principal components. Sample groups, defined by Treatment (Controls vs. Cd-  
 421 Treated series) and Exposure Time (0, 24, 48 hours), are displayed as points with colored 95% confidence ellipses  
 422 delineating group clustering. Points oriented in the direction of a vector suggest higher values of the corresponding  
 423 biomarker. (PC1: 43.3%; PC2: 19.5%).

424



425 **Figure 8.** Redundancy Analysis (RDA) Biplot of Biochemical Biomarkers Constrained by  
 426 Cadmium Treatment and Exposure Time. **Legend:** The RDA Biplot shows the relationship between  
 427 biochemical parameters (Carbohydrates, Lipids, Proteins, H<sub>2</sub>O<sub>2</sub> levels, LOOH levels, and SOD activity) and the  
 428 explanatory variables Treatment (Controls vs. Cd-Treated series) and Exposure Time (0, 24, 48 hours). Biochemical  
 429 variables are represented as vectors indicating their strength and direction of influence on the ordination axes. Sample  
 430 points are grouped according to treatment and exposure duration, with 95% confidence ellipses highlighting clustering  
 431 patterns. The first two RDA axes explain 27% and 5.1% of the variance, respectively. Vectors pointing toward sample  
 432 clusters indicate higher biomarker expression under those conditions.

#### 433 4. Discussion

434 Short-term tests were designed to measure not only mortality rates, but also to determine tolerance  
435 ranges of metabolic systems most sensitive to metal stress. The toxicity tests are necessary in water  
436 pollution evaluation, as chemical and physical parameters are insufficient to determine potential  
437 effects on aquatic organisms. Studying the responses of organisms to environmental stressors plays  
438 a key role in ecotoxicological studies. The established lethal concentration values could be used as  
439 key indicators in toxicity tests to demonstrate the effects of toxic pollutants, particularly for  
440 ecotoxicological evaluation of marine ecosystems in risk assessments. This work aims to evaluate  
441 the toxic effects of cadmium and its capacity to alter several variations in organism function.  
442 Namely, the estimation of biochemical contents, and the evaluation of enzymatic activities in *O.*  
443 *bicornis* species. The most toxicants exert their effects at basic level of the organism by reacting  
444 with enzymes or metabolites and other functional components of the cell (Ho Yu 2000). Thus,  
445 organisms' energy reserves provide a valuable indicator of their ability to cope with stress long  
446 before negative effects appear at more complex biological levels (Smolders *et al.* 2003). To examine  
447 energy metabolism perturbations, many bioenergetic parameters are available such as levels of  
448 energy reserves as carbohydrates, lipids and proteins (Meyer *et al.* 2002), cellular energy allocation  
449 (De Coen and Janssen 2003), scope for growth (Widdows *et al.* 2002). This research represents the  
450 first attempt to evaluate cadmium toxicity using *Ophelia bicornis* as a test organism. In this context,  
451 the energy status of *O. bicornis* was studied to gain insight into Cd-induced lethal effects and the  
452 ability of cadmium to induce disturbances on different levels of organism function. Cadmium is a  
453 very toxic metal, and even though it is unable to undergo Fenton-type reactions and therefore is  
454 considered a non-redox metal, and it has nevertheless been associated to increased production of  
455 reactive oxygen species (ROS) to marine organisms (Unsal *et al.* 2020; Liu *et al.* 2022).

456 Our results revealed that cadmium exposure and duration both influenced the biochemical responses  
457 in *O. bicornis*. Energy-related compounds such as carbohydrates and lipids were mobilized over  
458 time in the cadmium-treated series compared to the control group, reflecting a physiological

459 adjustment to metal-induced stress. However, protein levels remained relatively constant,  
460 suggesting that they were not significantly affected by treatment or exposure time. These findings  
461 are consistent with the notion that the earliest toxic effects of chemicals manifest primarily at the  
462 cellular and biochemical levels. As highlighted by Zhou *et al.* (2004), recent studies have  
463 increasingly focused on measurement at these levels to assess whether exposure to pollutants causes  
464 harmful effects. The observed changes in carbohydrate and lipid contents, along with the stability of  
465 protein levels, reflect such early biochemical responses to cadmium-induced stress. The  
466 biochemical composition (carbohydrates, lipids and proteins) indeed varies between taxonomic  
467 groups, but also among species, developmental stages, analyzed tissues, and environmental  
468 conditions. Moreover, this study suggests that the main storage materials in *O. bicornis* are proteins,  
469 whereas carbohydrates and lipids were present at a relatively low level.

470 The increase in carbohydrates levels after 48h oh exposure likely reflects a stress-induced  
471 mobilization of glycogen or other carbohydrate stores to provide readily available energy for  
472 detoxification possesses. The observed increase in carbohydrate levels in cd-exposed *Ophelia*  
473 *bicornis* may reflect a typical stress response in marine invertebrates. However, the utilization of  
474 carbohydrates by aquatic animals is relatively low compared to that of terrestrial animals. In marine  
475 invertebrates, carbohydrates are often the first source of energy utilized during acute stress, as they  
476 can be quickly metabolized through glycolysis to produce ATP (Wang *et al.* 2016).Increases in  
477 circulating glucose in the hemolymph have been reported in response to a variety of stressors such  
478 as salinity, temperature, hypoxia, or exposure to heavy metals (Wang *et al.* 2012). Notably,  
479 exposure to cadmium induced in the crayfish *Procambarus clarkii* a significant increase in  
480 hemolymph glucose concentration (Reddy *et al.* 1994).Carbohydrates and lipids play an important  
481 role in building up reserves for embryo development in Annelids. At the same time, the  
482 reproductive effort is high with up to 80% of the total energy allocated to gamete biomass (Hoeger  
483 and Schenk 2024). With increasing levels of stress, the maintenance of physiological integrity  
484 becomes more challenging for organisms. As an immediate response to the toxicity, organisms tend

485 to reduce their metabolic rate (Hand and Hardewig 1996). Some studies have suggested that the  
486 lipid contents of the exposed animals were significantly decreased compared to the control (Lucia *et*  
487 *al.* 2010; Yang *et al.* 2013), and this was confirmed by our study. In addition to carbohydrates, total  
488 lipid content significantly decreased in *Ophelia bicornis* under cadmium exposure, particularly after  
489 48h, suggesting a depletion of lipid reserves in response to metal-induced stress. This finding is  
490 consistent with previous studies in crustaceans such as crabs, where lipid levels in the  
491 hepatopancreas and ovary decreased after prolonged cadmium exposure compared to controls (Liu  
492 *et al.* 2016). Lipid reserves are mobilized or degraded under metal-induced stress, likely to meet  
493 elevated energy demands or due to oxidative damage. Luis and Passos (1995) found that lipids are  
494 crucial during the life cycle of *N. diversicolor*, particularly for reproduction. Lipids accumulate as  
495 an energetic reserve in oocytes and support metabolism during gametogenesis when the animals  
496 stop feeding. In aquatic invertebrates, reductions in the sequestration of reserves have been shown  
497 to affect the number or quality of gametes produced (Mathieu and Lubet 1993).

498 In contrast, protein contents remained relatively stable. The stability in protein levels suggests that  
499 *O. bicornis* reserves proteins, avoiding their breakdown unless under severe or prolonged stress.  
500 Proteins were not preferentially mobilized as energy sources under these exposure conditions.  
501 Similarly, according to the hypothesis proposed by Ketata *et al.*(2007), proteins are not primarily  
502 involved in energy metabolism in the clam *Ruditapes decussatus* exposed to cadmium. This  
503 observation aligns with our results, which also revealed no significant change in protein levels  
504 across treatment conditions or exposure durations. This contrasts with the findings of Barber and  
505 Blake (2006) reported that proteins can serve as a significant energy source, particularly when other  
506 reserves are depleted. Bivalves store energy primarily as carbohydrates (mainly glycogen) and  
507 lipids to maintain physiological integrity, during reproductive periods, whereas annelids tend to rely  
508 more on proteins to support their fundamental metabolic activities, it has been shown that proteins  
509 are the main components of muscle tissue. According to the research of (Tripp-Valdez *et al.* 2019)  
510 proteins, complex macromolecules described as biopolymers, are the most abundant of cellular

511 organic molecules. Under severe stress conditions, proteins may serve as an alternative energy  
512 source. However, this process compromises their structural and functional roles, since proteins are  
513 not primarily synthesized or stored for energy supply (Le Gal *et al.* 1997). On the other hand, Cd  
514 exposure has been shown to alter protein metabolism in bivalves, affecting both synthesis and  
515 degradation pathways in various tissues. The effects of cadmium on protein levels were tissue-  
516 specific and varied with the duration or exposure (Ivanina *et al.* 2008). Cadmium is known to affect  
517 reproduction and energy metabolism in marine organisms (Baudou *et al.* 2017; Louis *et al.* 2021).  
518 Concurrently, oxidative stress-induced protein oxidation can result in structural alterations and loss  
519 of function, disrupting cellular processes dependent on properly functioning proteins (Boguszevska-  
520 Mańkowska *et al.* 2015). A significant decrease in lipids, proteins and carbohydrates contents in  
521 muscles and gills of *Carcinus Aestuarii* as compared to *Portunus segnis* has been observed (Chetoui  
522 *et al.* 2021). Thus, seasonal changes can influence the physiology of crabs by affecting their  
523 metabolic activity and response, leading to disturbances in biochemical components such as lipids,  
524 proteins and carbohydrates. These alterations appear to be linked to the accumulation of trace  
525 elements, environmental factors and the reproductive cycles of aquatic organisms (Chetoui *et al.*  
526 2021).

527 At the cellular level, oxidative stress plays an important role in the development of damage.  
528 Reactive oxygen species (ROS) can be free radicals derived from oxygen, such as superoxide anion  
529 ( $O_2^-$ ) and hydroxyl radicals (OH), or non-radical derivatives such as hydrogen peroxide ( $H_2O_2$ ).  
530 These molecules are produced through several mechanisms (Fouzai *et al.* 2020). While ROS play  
531 essential roles in cellular functions, their excessive accumulation beyond the antioxidant capacity of  
532 the organism causes oxidative damage to biomolecules, leading to disruptions in cellular functions  
533 and structures (Sinenko *et al.* 2021). The degradation of these radicals is controlled by antioxidants  
534 systems, which adapt to the level of radicals present. The antioxidant system of marine organisms  
535 consists of low molecular weight scavengers and antioxidant enzymes which interact in a  
536 sophisticated network. Several studies on aquatic organisms have demonstrated the importance of

537 antioxidant enzymes in protecting cellular systems from oxidative damages induced by xenobiotic.  
538 The impact of oxidative stress has been shown to extend beyond individual cellular processes,  
539 influencing various downstream effects of metal toxicity (Chen *et al.* 2018). The deleterious effects  
540 of metals like cadmium are responsible for oxidative damage in living organism. Previous  
541 investigations indicated that cadmium induces the formation of oxygen free radicals in tissues and  
542 inhibits the activity of some enzymes of the antioxidant defense system (Unsal *et al.* 2020). When  
543 organisms are exposed to elevated levels of trace metals, oxidative stress can damage to essential  
544 biomolecules such as proteins, lipids will be happened, impairing physiological function (Singh *et*  
545 *al.* 2019; Haidar *et al.* 2023). In response, organisms have developed a variety of defense systems to  
546 mitigate the toxic effects of metals and maintain internal homeostasis. One of these mechanisms is  
547 the production of antioxidant enzymes such as superoxide dismutase (SOD), catalase, and  
548 glutathione, which work synergistically to neutralize reactive oxygen species (ROS) generated  
549 during metal exposure (Jomova *et al.* 2023). Cope with oxidative stress produced by ROS  
550 generation, SOD as the most powerful and the primary detoxification enzyme, removes the  
551 superoxide radical through the process of dismutation to oxygen and hydrogen peroxide. In the  
552 present study, changes in the activity of SOD were observed in *O. bicornis* under cadmium  
553 treatment. Similar to findings in other marine invertebrates exposed to Cu (Brown *et al.* 2004), or  
554 reported in field studies (Pérez *et al.* 2004), Cd may induce oxidative damage in *O. bicornis*,  
555 potentially through the generation of ROS. In response to oxidative stress, organisms exposed to  
556 various organic and metal contaminants often exhibit increased activity of antioxidant enzymes.  
557 However, these responses vary depending on the species, the enzymes involved, and whether the  
558 exposure is to a single or a mixture of contaminants (Regoli and Giuliani 2014). SOD activity in *O.*  
559 *bicornis* exposed to cadmium remained constant at 24h but showed an increase after 48h of  
560 exposure, with values still higher than those of the control at day 0. Our results are in full agreement  
561 with those reported in literature. Similar results were obtained in previous investigations of several  
562 polychaete species. According to (Sun and Zhou 2008), the activity of SOD in *N. diversicolor*

563 exposed to Cd and petroleum hydrocarbons (PHCs) increased after a 6 day exposure, and (Moreira  
564 *et al.* 2006) also reported a significant induction of this enzyme when the species was exposed to  
565 contaminated sediments. The increase in SOD activity suggested that SOD was induced as  
566 consequence of generation of  $O_2^-$ . This response likely reflects the role of the antioxidant enzyme  
567 SOD in removing excess free radicals, mainly  $O_2^-$  and  $H_2O_2$  generated to counteract Cd-induced  
568 stress and mitigate their toxicity. However, (Chaâbane *et al.* 2020) showed that SOD activity  
569 enhanced in the gills and digestive gland of chromium (VI)-exposed *Venus verrucosa*. ROS  
570 accumulation is a widely documented effect of cadmium exposure in marine organisms leading to  
571 oxidative damage and cellular dysfunction (Liu *et al.* 2022). Cd does not induce ROS production  
572 directly; however, it can cause indirect damage and generate free radicals (Patra *et al.* 2011). This  
573 indirect damage was elucidated in the present study by a significant increase of  $H_2O_2$  levels  
574 confirming the metal potential to boost pro-oxidants' production. The enhanced  $H_2O_2$  generation in  
575 *O. bicornis* was associated with an increase in LOOH levels, as compared to the control group.  
576 Hydrogen peroxide is a central redox signaling molecule, capable of serving as messenger to carry a  
577 redox signal from the site of its generation to a target site (Sies *et al.* 2017). The levels of  $H_2O_2$  were  
578 found to be increased in *D. magna* adults exposed to the metal mixture (Cu/Cd), oxidative stress  
579 was found as a common mechanism underlying the toxicity of these metals (Majid 2024). These  
580 findings are in line with those of (Chetoui *et al.* 2022), and (Chaâbane *et al.* 2020). Our study found  
581 that Cd exposure at a sublethal concentration ( $LC_{10}$ ) induced significant oxidative stress responses  
582 in *Ophelia bicornis*. Lipid hydroperoxides (LOOH) and hydrogen peroxide ( $H_2O_2$ ) increased  
583 significantly over time, indicating heightened ROS production and lipid peroxidation. Concurrently,  
584 superoxide dismutase (SOD) activity increased after 48h, suggesting an enzymatic attempt to  
585 counteract ROS accumulation. However, the sustained elevation of  $H_2O_2$  and LOOH, despite SOD  
586 activation, highlights an oxidative imbalance. This underscores the vulnerability of *O. bicornis*  
587 tissues to damage caused by Cd exposure. These findings regarding energy metabolism and stress  
588 responses may help predict the effects of Cd on marine organisms, particularly annelids such as *O.*

589 *bicornis*, which are highly exposed to metal contaminants in their habitats. *O. bicornis* is especially  
590 important in ecotoxicological studies, living in direct contact with contaminated sediments.

## 591 **5. Conclusion**

592 The physiological and biochemical responses of the annelid *Ophelia bicornis* were investigated  
593 under laboratory conditions after short exposure to cadmium (Cd), focusing on antioxidant enzyme  
594 activity and the content of major biochemical components. These findings provide insight into the  
595 molecular responses triggered by Cd exposure and the underlying mechanisms in *O. bicornis*,  
596 emphasizing the significant role of oxidative stress in mediating the cellular response to cadmium-  
597 induced damage. responses of the antioxidant system and biochemical parameters in this species  
598 are not well-documented; thus, our study provides the first evidence addressing these aspects  
599 regarding cadmium toxicity. This research highlights the ecotoxicological relevance of *O. bicornis*  
600 as a sensitive bioindicator of cadmium exposure. Even at sublethal concentrations, Cd disrupted the  
601 energy balance in exposed organisms and induced oxidative stress. this is evidenced by increased  
602 carbohydrates mobilization, alterations of lipid contents, elevated levels lipid hydroperoxides  
603 (LOOH) and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and a time dependent modulation of superoxide dismutase  
604 (SOD) activity. These physiological and biochemical changes indicate that *O. bicornis* relies on its  
605 energy reserves and antioxidant defenses to cope with metal-induced stress. The consistent  
606 correlation between of oxidative stress biomarkers and cadmium exposure further underscores their  
607 value as early warning indicators. Overall, these findings enhance our understanding of how benthic  
608 invertebrates respond to metal-contaminated environments and confirm the sensitivity of *O.*  
609 *bicornis*. This establishes its use as a bioindicator in environmental monitoring and provides a  
610 baseline for evaluating its physiological responses to metal exposure, and opens a new perspective  
611 for its application in future ecotoxicological and biomonitoring studies.

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