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# Remediation of Arsenic from contaminated seawater using manganese spinel ferrite nanoparticles: ecotoxicological evaluation in *Mytilus galloprovincialis*

Francesca Coppola[a], Daniela S. Tavares[b,c], Bruno Henriques[b,d], Rui Monteiro[b,d], Tito Trindade[c], Amadeu M.V.M. Soares[a], Etelvina Figueira[a], Gianluca Polese[e], Eduarda Pereira[b], Rosa Freitas[a]

[a]Departamento de Biologia & CESAM, Universidade de Aveiro, 3810-193 Aveiro, Portugal

[b]Departamento de Química & CESAM, Universidade de Aveiro, 3810-193 Aveiro, Portugal

[c]Departamento de Química & CICECO, Universidade de Aveiro, 3810-193 Aveiro, Portugal

[d]CIIMAR, Universidade do Porto, 4050-123 Porto, Portugal

[e]University of Naples, 80126 Naples, Italy

**Corresponding author:** Rosa Freitas

Address: Departamento de Biologia, Universidade de Aveiro

Campus Universitário de Santiago

3810-193 Aveiro, Portugal

e-mail address: rosafreitas@ua.pt

**ABSTRACT**

In the last decade different approaches have been applied for water remediation purposes, including the use of nanoparticles(NPs) to remove metals and metalloids from water. Although studies have been done on the toxic impacts of such NPs, very scarce information is available on the impacts of water after decontamination when discharged into aquatic environments. In this way, the present study we aimed to evaluate the ecotoxicological safety of seawater previously contaminated with arsenic (As) and remediated by using manganese-ferrite (MnFe2O4) nanoparticles (NPs). For this, mussels *Mytilus galloprovincialis* were exposed for 28 days to different conditions, including clean seawater (control), As (1000 $\mu$g L$^{-1}$) contaminated and remediated (As 70 $\mu$g L$^{-1}$) seawater, water containing manganese-ferrite (MnFe2O4) nanoparticles (NPs) (50 mg L$^{-1}$) with and with the presence of As. At the end of exposure, concentrations of As in mussels tissues were quantified and biomarkers related to mussels' metabolism and oxidative stress status were evaluated. Results revealed that mussels exposed to water contaminated with As and to As+NPs accumulated significantly more As (between 62% to 76% more) than those exposed to remediated seawater. Regarding biomarkers, our findings demonstrated that in comparison to remediated seawater (conditions a, b, c) mussels exposed to contaminated seawater (conditions A, B, C) presented significantly lower metabolic activity, lower expenditure of energy reserves, activation of antioxidant and biotransformation defences, higher lipids and protein damages and greater AChE inhibition. Furthermore, organisms exposed to As, NPs or As+NPs revealed similar biochemical effects, both before and after water decontamination. In conclusion, the present study suggests that seawater previously contaminated with As and remediated by manganese-ferrite (MnFe2O4) NPs presented significantly lower toxicity than As contaminated water, evidencing the potential use of these NPs to remediate seawater contaminated with As and its safety towards marine systems after discharges to these environments.

60

## 1. INTRODUCTION

The increment of pollutants in aquatic environments is closely related with the growth of the world population (Zhang et al., 2015). Studies demonstrated that intense urbanization and industrial activities, with the associated effluents, result in an increase of pollution in the aquatic systems, especially in marine environments (Nardi et al., 2017, Belivermiş et al., 2016; Oliveira, 2015; Ventura-Lima et al., 2011). Often, the final destination of pollutants are lagoons and estuaries (Zhang et al., 2015), with tendency to be accumulated not only in sediments but also by organisms inhabiting these areas (Buffet et al., 2014; Ventura-Lima et al., 2009, 2011). Among the most common pollutants in aquatic environments is arsenic (As), a naturally occurring element (ATSDR, 2015; Saxe et al., 2006) released by natural activities, such as volcanism, dissolution of minerals (particularly into groundwater), but also by human activities, such as mining, metal smelting, combustion of fossil fuels, agricultural pesticide production and use, remobilization of historic sources, including mine drainage water (WHO, 2010; Mandal and Suzuki, 2002; Bhattacharya et al., 2007; Matschullat, 2000; Jang et al., 2016). As a results of its high toxicity, even at trace levels, As presents environmental concerns (IARC, 2012; Quasimeme, 2003; Fattorini et al., 2006). For this reason, currently As is considered the most priority hazardous substance in the environment based on the combination of substance frequency, toxicity and human exposure potential (ATSDR, 2015; Khan et al., 2010). In particular, the presence of As in aquatic systems has already proven to induce toxic impacts in a diversity of species, namely in bivalves, including physiological and biochemical impairments in clams (Freitas et al., 2018) and mussels (Coppola et al., 2018).

Because of aquatic pollution and associated concerns, nowadays an important research topic is the development of new technologies for wastewater decontamination (Gehrke et al., 2015; Davidescu et al., 2015). Different methodologies have been developed to remove pollutants from waters, including oxidation/precipitation (Leupin et. al, 2005; Dutta et. al, 2005; Lee et al., 2002), coagulation/co-precipitation (Hansen et al., 2006; Kumar et al., 2004), sorption, ion-exchange (Baciocchi et al., 2005; Kim and Benjamin, 2004), membrane technologies (Kim et al., 2006; Ballinas et al., 2004), solvent extraction and bioremediation (Kordmostafapour et al., 2006; Iberhan et al., 2003; Katsoyiannis et al., 2002). Some of these

90   techniques have shown a great potential for removing inorganic pollutants from water (Gehrke

91   et al., 2017; Mohan et al., 2006). Among the innovative techniques, one of the most promising

92   approaches to decontaminate water is based on the use of nanoparticles (NPs), with some

93   laboratory studies evidencing their high effectiveness in the removal of metal(loid)s (Tavares et

94   al., 2013; Zhang et al., 2010; Mohan and Pittman, 2007). In particular, manganese-ferrite

95   ($MnFe_2O_4$) nanoparticles (NPs), a common spinel ferrite material has shown to be very effective

96   in decreasing inorganic pollution (including metals and metalloids) in freshwater and seawater

97   (Zhang et al., 2010; Tavares et al., 2013; Jang et al., 2016; Santhosh et al., 2014). However,

98   although the use of $MnFe_2O_4$-NPs for water decontamination is undoubtedly one of the most

99   challenging research areas, important aspects are still missing, such as the potential toxicity of

100  these NPs and the ecotoxicological evaluation of the remediated water (Bhatt and Tripathi,

101  2011; Lovern and Klaper, 2006; Lovern et al., 2007; Smith et al., 2007; Warheit et al., 2007).

102  Together with decontaminated water or resulting from leaching of chemical elements, after

103  application these NPs can end up in aquatic environments, making crucial the assessment of

104  decontaminated water potential impacts towards inhabiting organisms. Until now, different

105  studies have already demonstrated the impacts induced directly by magnetic $MnFe_2O_4$ spinel

106  ferrite NPs in algae, crustaceans and fish, revealing their potential hazard potential to different

107  aquatic species (Bahadar et al., 2016; Beji et al., 2010; Aslibeiki et al., 2016; Federici et al.,

108  2007). Nevertheless, no studies have been carried out to evaluate the toxicity of water

109  decontaminated by these NPs.

110       To evaluate the impacts of the presence of pollutants, including NPs, in the aquatic

111  environment, benthic species are a good biological model as they accumulate and reflect the

112  impacts of different substances (Velez et al., 2015; Attig et al., 2014; Banni et al., 2014a; Hu et

113  al., 2015; Nardi et al., 2017; Coppola et al., 2018; Freitas et al., 2018). Among these species is

114  the mussel *Mytilus galloprovincialis*, identified by several authors as a good bioindicator with the

115  capacity to respond to environmental disturbances, presenting a wide spatial distribution and

116  economic relevance (Coppola et al., 2017; Richir and Gobert, 2014; Freitas et al., 2017; Kristan

117  et al., 2015; Mejdoub et al., 2017). This bivalve is a sedentary filter-feeder and has a large

118  capacity to accumulate pollutants (Coppola et al., 2018; Livingstone et al., 2000; Selvin et al.,

119  2000).

120    Thus, by the above-mentioned, an important topic of research is to understand if the

121    application of NPs to decontaminate seawater still constitutes a threat to aquatic environment,

122    affecting negatively the inhabiting organisms. For this reason, the present study aimed to

123    evaluate the toxicity induced in the mussel *M. galloprovincialis* exposed to seawater previously

124    contaminated with As and decontaminated with $MnFe_2O_4$, NPs. After exposure to

125    decontaminated seawater, biomarkers related to mussels' metabolic, oxidative stress and

126    neurotoxic status were evaluated.

127

128

129 **2. MATERIALS AND METHODS**

130 2.1 Experimental conditions

131 The Mediterranean mussel *Mytilus galloprovincialis* was selected as biological model for

132 this study (e.g. Coppola et al., 2018; Della Torre, 2015; Gomes et al., 2011). Organisms were

133 collected in November 2017, at the Ria de Aveiro lagoon (Portugal), with a mean body weight of

134 $21.3 \pm 6.6$ g, fresh weight (FW).

135 Bivalves were transported from the field to the laboratory in plastic containers, where they

136 were placed in aquaria for depuration and acclimation to laboratory conditions for 2 weeks. To

137 simulate field conditions, in the laboratory organisms were exposed to: temperature $18.0 \pm 1.0$

138 ºC; pH $8.0 \pm 0.1$, photoperiod 12 h light and 12 h dark, and continuous aeration, in artificial

139 seawater (salinity $30 \pm 1$) (Tropic Marin® SEA SALT from Tropic Marine Center). Seawater was

140 renewed daily during the first week and then every three days until the end of the acclimation

141 period.

142 After the acclimation period organisms were distributed in different aquaria according to

143 the conditions described in Table 1. Seven different conditions were evaluated, with 3 aquaria

144 (containing 3 L of seawater each) per condition and 4 individuals per aquarium/replicate (12

145 individuals per condition).

146 Decontaminated seawater was obtained by adding 50 mg $L^{-1}$ of $MnFe_2O_4$ NPs to water

147 previously contaminated with 1000 µg $L^{-1}$ of As. The NPs were removed from seawater after 24

148 hours by applying a magnetic field (although a non-quantifiable residual amount of NPs may

149 hypothetically remain in water) as described by Mohmood et al. (2016).

150 During the experimental period (28 days), water medium was changed weekly and

151 exposure conditions completely re-established, including contaminants concentrations and

152 seawater characteristics (salinity, pH, temperature). Every week, immediately after medium

153 renewal, samples of seawater were collected from each aquarium for As quantification.

154 The concentration of As, 1000 µg $L^{-1}$, was selected according to the emission limit value

155 for this element in wastewater discharges (Decree-Law No. 236/98, in Portuguese), while 70 µg

156 $L^{-1}$ is the residual concentration of As reached in seawater after decontamination with $MnFe_2O_4$,

157 NPs (data from preliminary experiments, not shown).

158   During the entire experimental period (28 days) aquaria were continuously aerated, with a

159 12 light: 12 dark photoperiod. As for the acclimation, temperature (17 ± 1.0 ºC), pH (8.0± 0.1)

160 and salinity (30 ± 1) values were selected considering measurements done at the sampling site

161 (data not provided), and were daily checked and adjusted if necessary.

162   During the experimental period organisms were fed with Algamac protein plus (150,000

163 cells/animal) twice a week. Mortality was also daily checked, with 100% of survival recorded

164 during the experimental period.

165   At the end of the exposure period, organisms were frozen individually with liquid nitrogen

166 and stored at -80ºC, until homogenization of each individual soft tissue using a mortar and a

167 pestle under liquid nitrogen. Each homogenized organism was divided into aliquots (0.5 g each)

168 for biomarkers analyses and As quantification.

169

170  2.2 Synthesis and characterization of $MnFe_2O_4$ nanoparticles

171   $MnFe_2O_4$ nanoparticles were prepared by the chemical oxidative hydrolysis of a mixture

172 of $FeSO_4.7H_2O$ and $MnSO_4.H_2O$ in alkaline conditions. Different techniques were applied to

173 perform the chemical, physical and structural characterization of NPs. The morphology and

174 particle size of the NPs were confirmed by transmission electron microscopy (TEM) using the

175 Hitachi H-9000 TEM microscope operating at 300 kV. For TEM analysis, one drop of sample

176 dispersed in ethanol was placed onto carbon-coated copper grid and then let the solvent

177 evaporate. The surface area of the NPs was determined by $N_2$ adsorption/desorption on a

178 Gemini V2.0 Micromeritics instrument. The crystalline phase of the NPs was identified by x-ray

179 powder diffraction of the powders using a Philips Analytical PW 3050/60 X'Pert PRO (θ/2θ)

180 diffractometer equipped with an X'Celerator detector and with automatic data acquisition (X'Pert

181 Data Collector v2.0b software) by a monochromatized Cu Kα radiation (λ = 1,54056 Å) at 45

182 Kv/40 Ma. The NPs Fourier-Transform Infrared (FT-IR) spectrum was recorded on a Mattson

183 7000 spectrometer, at 4 $cm^{-1}$ resolution, using a horizontal attenuated total reflectance (ATR)

184 cell.

185   The average size distribution of $MnFe_2O_4$ NPs in water at salinity 30 were measured by

186 Dynamic Light Scattering (DLS) at T0 (immediately injected into seawater media), T1 (after 1

187 hour) and T24 (after 24 hours) (Table 2). These time periods were selected based on previous

188    studies (Yao et al., 2014, Yang et al., 2012; Aubery et al., 2011) that showed aggregation and

189    precipitation of different Fe-NPs within 24 h. DLS measurements were performed on a Delsa

190    Nano C from Beckman Coulter, Inc. (Fullerton, CA) equipped with a laser diode operating at

191    658 nm. Scattered light was detected at 165° angle and analysed by using a log correlator over

192    120 accumulations, for 1.0 mL of sample in a UV cuvette semi-micro. Each sample was

193    reproducibly shaken before analysis and exposed to the minimum of DLS measurements

194    needed to obtain at least three valid data. The calculation of particle size distribution and

195    distribution averages was performed by using CONTIN particle size distribution analysis

196    routines through Delsa Nano 3.73 software. The hydrodynamic radius and polydispersity index

197    of the analysed dispersions were calculated on three replicates of each sample by using the

198    cumulant method. Undetected colloidal material at the end of each measurement is indicated as

199    Invalid data (I.d.).

200

201        2.3 Arsenic quantification

202        The quantification of As in water samples collected from each condition (Table 3) was

203    performed by inductively coupled plasma mass spectrometry (ICP-MS), on a Thermo ICP-MS X

204    Series equipped with a Burgener nebulizer. The quantification limit of the method was 1 µg/L (n

205    = 12), with an acceptable relative standard deviation among replicates (n≥2) < 5% (Henriques et

206    al., 2019).

207        Total As concentrations  in $M.$ $galloprovincialis$ whole soft tissues (Table 4) were

208    quantified by ICP-MS, after microwave assisted acid digestion. Samples with 100–200 mg

209    (freeze-dried) were digested in a CEM MARS 5 microwave, firstly with 2 mL of $HNO_3$ (70%) at

210    170 °C for 15 min, followed by a second identical m icrowave cycle with 0.5 mL of $H_2O_2$ (30%).

211    After addition of $H_2O_2$, the mixture was allowed to stand for 15 min so that the microwave

212    reaction was not as violent. The obtained digests were transferred into 25 mL polyethylene

213    vessels and the volume made up with ultrapure water. The quality control was assured by

214    running procedural blanks (reaction vessels with only $HNO_3$ and $H_2O_2$) and certified reference

215    material TORT-2 (Lobster Hepatopancreas; 21.6 ± 1.8 mg $Kg^{-1}$ As) in parallel with samples.

216    Blanks were always below the quantification limit and mean percentage of recovery for As was

217    110 ± 4% (n = 4) (Coppola et al., 2018).

218

## 2.4 Biomarkers evaluation

The whole tissue of mussels was used for biomarkers determination (see section 2.1). For each parameter, 0.5 g of tissue per organism was used, with 2 individuals per replicate (6 per condition). For each condition, metabolic capacity (electron transport system activity, ETS), energy-related (glycogen content, GLY; total protein content, PROT), antioxidant defence (superoxide dismutase activity, SOD; glutathione peroxidase activity, GPx; glutathione S-transferases activity, GSTs), oxidative damage (lipid peroxidation levels, LPO; protein carbonyl levels, PC; glutathione content ratio, GSH/GSSG) and neurotoxicity (Acetylcholinesterase activity, AChE) biomarkers were assessed. Each sample was performed at least in duplicate. All measurements were done using a microplate reader (BioTek, Synergy HT). The extraction for each biomarker was performed with specific buffers: phosphate buffer for SOD, GPx, GSTs, PROT, GLY, CP and AChE; magnesium sulphate buffer for ETS; trichloroacetic acid buffer for LPO and KPE buffer for GSH/GSSG. Each sample was sonicated for 15 s at 4 ºC and centrifuged for 25 min (or 15 min for GSH/GSSG) at 10 000 $g$ (or 3 000 $g$ for ETS) (Coppola et al., 2018; De Marchi et al., 2018; Freitas et al., 2018). Supernatants were stored at -20 ºC and used within a maximum period of 3 weeks.

235

*Metabolic capacity and energy-related biomarkers*

ETS activity was measured based on King and Packard (1975) and the modifications performed by De Coen and Janssen (1997). The absorbance was measured at 490 nm during 10 min with intervals of 25 s. The amount of formazan formed was calculated using $\varepsilon$ = 15,900 $M^{-1}cm^{-1}$ and the results expressed in nmol $min^{-1}$ per g of fresh weight (FW).

241

For GLY quantification the sulphuric acid method was used, as described by (Dubois et al., 1956). A calibration curve was obtained using glucose standards prepared in concentrations between 0 and10 mg $mL^{-1}$. Absorbance was measured at 492 nm and the results were expressed in mg per g FW.

246

247    The PROT content was determined following the spectrophotometric method of Biuret

248    (Robinson and Hogden, 1940), and bovine serum albumin (BSA) was used as standard (0–40

249    mg mL$^{-1}$) to obtain a calibration curve. Absorbance was measured at 540 nm. Concentrations of

250    PROT were expressed in mg per g FW.

251

252    *Antioxidant defences biomarkers*

253    The activity of SOD was quantify following the method of Beauchamp and Fridovich

254    (1971) and was performed with a calibration curve using SOD standards between 0.25 and 60

255    U mL$^{-1}$. The absorbance was measured at 560 nm and the results were expressed in U per g of

256    FW, where U represents the quantity of the enzyme that catalyses the conversion of 50% of

257    nitroblue tetrazolium (NBT).

258

259    The activity of GPx was determined following the method of Paglia and Valentine (1967).

260    Absorbance measurements were performed at 340 nm during 5 min in 10 s intervals and the

261    activity was determined using the extinction coefficient of $\varepsilon = 6.22$ mM$^{-1}$ cm$^{-1}$. Results were

262    expressed in U/g FW, where U corresponds to the quantity of enzyme which catalyzes the

263    conversion of 1 µmol nicotinamide adenine dinucleotide phosphate (NADPH) per min.

264

265    GSTs activity was determined according to Habig et al. (1976). The absorbance was

266    measured at 340 nm. The activity of GSTs was determined using $\varepsilon = 9.6$ mM$^{-1}$ cm$^{-1}$. The

267    enzymatic activity was expressed in U per g of FW where U is defined as the amount of enzyme

268    that catalysis the formation of 1 µmol of dinitrophenyl thioether per min.

269

270

271    *Oxidative damage biomarkers*

272    LPO was determined following the method described by Ohkawa et al. (1979). LPO

273    levels were measured trough the quantification of malondialdehyde (MDA), a by-product of lipid

274    peroxidation. Absorbance was measured at 532 nm ($\varepsilon = 156$ mM$^{-1}$ cm$^{-1}$). LPO levels were

275    expressed in nmol of MDA per g FW.

276

277    PC content was obtained following Levine et al. (1990). Absorbance of samples was

278    measured at 370 nm and the carbonyl content was calculated using an absorption coefficient ε=

279    0.022 mM$^{-1}$ cm$^{-1}$. Results were expressed in nmol of PC groups formed per g FW

280

281    GSH and GSSG glutathione contents were measured at 412 nm (Rahman et al., 2014)

282    and used as standards (0–60 µmol L$^{-1}$) to obtain a calibration curve. Absorbance was measured

283    at 412 nm, for both assays. The results were expressed as nmol per g of FW. The ratio

284    GSH/GSSG was determined taking in account the number of thiol equivalents (GSH/ 2 *

285    GSSG).

286

287

288    *Neurotoxicity biomarker*

289    Acetylthiocholine iodide (ATChI, 470 µmol L$^{-1}$) substrates were used for the determination

290    of Acetylcholinesterase (AChE) following the methods of Ellman et al. (1961) and modification

291    by Mennillo et al. (2017). Enzyme activity was recorded continuously for 5 min at 412 nm and

292    expressed in nmol per g FW.

293

294    2.5 Integrated biomarker response (IBR)

295    The integrated biomarker response (IBR) was calculated according to Beliaeff and

296    Burgeot (2002) aiming to evaluate the general mussel's biochemical response among 6

297    conditions. All biomarkers determined were used in the calculation of the IBR and they were

298    arranged clockwise in the following order: ETS, GLY, PROT, SOD, GPx, LPO, CP, GSH/GSSG,

299    GST, AChE. Values were discussed in terms of a general response given by the final IBR value,

300    where higher values correspond to higher mussels' response.

301

302    2.6 Statistical analyses

303    All the biochemical results (ETS, GLY, PROT, SOD, GPx, GSTs, LPO, PC, GSH/GSSG

304    and AChE) and As concentrations in mussels tissues, obtained from each condition, were

305    submitted to statistical hypothesis testing using permutational analysis of variance, employing

306    the PERMANOVA+add-on in PRIMER v6 (Anderson et al., 2008). The pseudo-F *p*-values in the

307     PERMANOVA main tests were evaluated in terms of significance. When significant differences

308     were observed in the main test, pairwise comparisons were performed. Values lower than 0.05

309     ($p \leq 0.05$) were considered as significantly different. For each biomarker, *p*-value obtained for

310     pair-wise comparisons are represented with *p*-value in Table 5.

311         For As concentrations and each biomarker, the null hypotheses (H0) tested were: i) no

312     significant differences exist among CTL and all the contaminated conditions (CTL, A, B and C).

313     *p*-values are presented in Table 5, with significant differences highlighted in bold; ii) no

314     significant differences exist among decontaminated conditions (CTL, a, b, and c). *p*-values are

315     presented in Table 5, with significant differences highlighted in bold; iii) no significant

316     differences exist between A vs a, B vs b, C vs c conditions. Significant differences between

317     each pair of conditions are represented with an asterisk in figures.

318

319    **3. RESULTS**

320    3.1 Characterization of $MnFe_2O_4$ nanoparticles

321    $MnFe_2O_4$ NPs showed a spheroidal morphology (Figure 1) with a mean diameter and

322    standard deviation of 75 ± 15 nm. The infrared spectrum of the NPs displayed a characteristic

323    band at 537 $cm^{-1}$ related to metal-O stretching vibration of the $MnFe_2O_4$ NPs (Bellusci et al.,

324    2009; Mehran et al., 2016; Tavares et al., 2013). The band at 1107 $cm^{-1}$ was attributed to metal-

325    OH and to metal-$OH_2$ stretching vibrations, which correspond to water sorption on oxide, while

326    1635 $cm^{-1}$ band is due to H-O-H bending and corresponds to molecular water adsorbed or

327    incorporated into the crystalline lattice (Bellusci et al., 2009). The broad band at 3309 $cm^{-1}$

328    corresponds to symmetric and asymmetric stretching of O-H bond (Margabandhu et al., 2016).

329    Powder X-ray diffraction (XRD) pattern show peaks that are characteristics of the presence of

330    $MnFe_2O_4$ with the spinel structure (JCPDS–International centre diffraction data, PDF card 01-

331    071-4919). In seawater, an aggregation of the NPs was observed by DLS, reaching an average

332    size of approximately 60000 nm, after 24 hours. The presence of As in water did not influence

333    NPs aggregation since sizes in conditions A, B, a and b, after 24 hours, the average sizes were

334    statistically indistinguishable.

335

336    3.2 Arsenic quantification in seawater and mussels' tissues

337    Concentrations of As in water samples revealed that real and nominal concentrations

338    were similar, both for A and a conditions. In water samples from conditions without As (B and b)

339    the concentrations of this metalloid were lower than the quantification limit (1.5 µg $L^{-1}$).

340    Concentration of As in water after decontamination was 55 ± 13 µg $L^{-1}$. Because sorption of As

341    by the NPs is extremely rapid, As was not possible to quantify in water from condition C (Table

342    3).

343    The results obtained from As quantification in mussels showed a significant difference

344    between organisms exposed to CTL and those exposed to A and C conditions (Table 4).

345    No significant differences were found between *M. galloprovincialis* submitted to CTL and

346    the organisms exposed to conditions a, b and c (Table 4).

347    Significant differences were observed in terms of As concentrations between mussels

348    exposed to initial (before decontamination) and final (after decontamination) conditions (A vs a

349    and C vs c) (Table 4). Organisms exposed to condition A accumulated more 76% of As than

350    those exposed to condition a, while the contents of As in the mussels exposed to condition C

351    were 62% higher than those in condition c.

352

353    ### 3.3 Biochemical markers

354    *Metabolic capacity and energy-related biomarkers*

355    The ETS activity was significantly higher at control (CTL) in comparison to the values

356    obtained in mussels exposed to As contaminated seawater (conditions A, B, C; resembling

357    initial concentrations, measured before decontamination), with the lowest values at condition A

358    (Figure 2A, Table 5). ETS activity was significantly higher at control (CTL) in comparison to the

359    values obtained in mussels exposed to decontaminated seawater (conditions a, b, c) (Figure

360    2A, Table 5).

361    The ETS activity was significantly higher in organisms exposed decontaminated seawater

362    (conditions a, b, c) in comparison to organisms exposed to As contaminated seawater

363    (conditions A, B, C) (Figure 2A).

364

365    The GLY content was significantly lower in mussels exposed to control (CTL) in

366    comparison to the values observed in mussels exposed to As contaminated seawater

367    (conditions A, B, C) (Figure 2B, Table 5).

368    Significantly lower GLY content was obtained in organisms exposed to decontaminated

369    seawater (conditions a, b, c) in comparison to organisms exposed to As contaminated seawater

370    (conditions A, B, C) (Figure 2B).

371

372    The PROT content was significantly lower in mussels exposed to control (CTL) in

373    comparison to values observed in mussels exposed to As contaminated seawater (conditions A,

374    B), while no significant differences were observed between CTL and C conditions (Figure 2C,

375    Table 5).

376    The PROT content was significantly lower in organisms exposed to decontaminated

377    seawater (conditions a, b, c) in comparison to organisms exposed to contaminated seawater

378    (conditions A, B, C) (Figure 2C).

379

380        *Antioxidant defence biomarkers*

381        The SOD activity was significantly lower at CTL in comparison to values obtained in

382   mussels exposed to As contaminated seawater (A, B, C) (Figure 3A, Table 5). Significantly

383   higher values were obtained in mussels exposed to condition A in comparison to organisms

384   exposed to conditions B and C (Figure 3A, Table 5).

385        The SOD activity was significantly lower in organisms exposed to decontaminated

386   seawater (conditions a, b, c) in comparison to organisms exposed to contaminated seawater

387   (conditions A, B, C) (Figure 3A).

388

389        The activity of GPx was significantly lower at CTL in comparison to values obtained in

390   mussels exposed to contaminated seawater (A, B, C) (Figure 3B, Table 4). Significant

391   differences were observed between organisms exposed to A and C conditions (Figure 3B,

392   Table 5).

393        Regarding to organisms exposed to decontaminated seawater, significantly higher GPx

394   activity was observed at a, b and c conditions in comparison to control (CTL) (Figure 3B and

395   Table 5). No significant differences were observed between organisms exposed to conditions a

396   and b (Figure 3B, Table 5).

397        The GPx activity values were significantly lower in organisms exposed decontaminated

398   seawater (conditions a, b, c) in comparison to organisms exposed to contaminated seawater

399   (conditions A, B, C) (Figure 3B).

400

401        The GSTs activity was significantly lower at CTL in comparison to values obtained in

402   mussels exposed to contaminated seawater (A, B, C) (Figure 3C, Table 5). No significant

403   differences were observed between organisms exposed to A and C conditions (Figure 3C,

404   Table 5).

405        Organisms under control (CTL) conditions showed significantly lower GSTs activity than

406   those exposed to decontaminated seawater (condition c) (Figure 3C and Table 4). No

407   significant differences were observed between organisms exposed to a and c conditions (Figure

408   3C, Table 5).

409     The GSTs activity values were significantly lower in organisms exposed to

410   decontaminated seawater (conditions a, b, c) comparatively to organisms exposed to

411   contaminated seawater (conditions A, B, C) (Figure 3C).

412

413     *Cellular damage biomarkers*

414     The LPO levels were significantly lower at control (CTL) in comparison to values obtained

415   in mussels exposed to contaminated seawater (conditions A, B, C) (Figure 4A, Table 4).

416   Significantly lower LPO values were found in organisms exposed to condition B in comparison

417   to organisms exposed to conditions A and C (Figure 4A, Table 5).

418     Significantly lower LPO levels were observed in organisms exposed to CTL compared to

419   organisms exposed to condition a, b and c (Figure 4A and Table 5). No significant differences

420   were observed among organisms exposed to a, b and c conditions (Figure 4A, Table 5).

421     The LPO levels were significantly lower in organisms exposed to decontaminated

422   seawater (conditions a, b, c) in comparison to organisms exposed to contaminated seawater

423   (conditions A, B, C) (Figure 4A).

424

425     The PC levels were significantly lower in mussels exposed to control (CTL) in comparison

426   to values observed in mussels exposed to contaminated seawater (conditions A, B, C) (Figure

427   4B, Table 5).

428     The PC levels in mussels exposed to control (CTL) were significantly lower than those

429   observed in mussels exposed to conditions a and b (Figure 4B, Table 4). No significant

430   differences were observed among organisms exposed to a, b and c conditions (Figure 4B,

431   Table 5).

432     The PC levels were significantly lower in organisms exposed to decontaminated

433   (conditions a, b and c) seawater comparatively to organisms exposed to contaminated seawater

434   (conditions A, B, C) (Figure 4B).

435

436     The GSH/GSSG values were significantly higher in mussels exposed to control (CTL) in

437   comparison to values observed in mussels exposed to contaminated seawater (conditions A, B,

438   C) (Figure 4C, Table 5).

439    Significantly higher GSH/GSSG values were observed in mussels exposed to control

440    (CTL) in relation to the values observed in mussels exposed to decontaminated seawater

441    (conditions a, b, c) (Figure 4C, Table 5). No significant differences were observed between

442    organisms exposed to a and c conditions (Figure 4C, Table 5).

443    The GSH/GSSG ratio was significantly higher in organisms exposed to decontaminated

444    seawater (conditions a, b, c) than in organisms exposed to contaminated seawater (conditions

445    A, B, C) (Figure 4C).

446

447    *Neurotoxicity biomarker*

448    The AChE activity was significantly higher in mussels exposed to control (CTL) in

449    comparison to the values observed in mussels exposed to contaminated seawater (conditions

450    A, B, C) (Figure 5, Table 5).

451    Significantly higher AChE values were observed in mussels exposed to control (CTL) in

452    comparison to those observed in mussels exposed to decontaminated seawater (conditions a,

453    b, c) (Figure 5, Table 5).

454    Significantly higher AChE values were observed in organisms exposed to

455    decontaminated seawater (conditions a, b, c) than in organisms exposed to contaminated

456    seawater (conditions A, B, C) (Figure 5).

457

458    3.4. Integrated Biomarker Response (IBR)

459    IBR values showed the highest score (16.7) for the mussels exposed to condition B,

460    which indicates higher impacts in organisms under MnFe2O4-NPs (50 mg L$^{-1}$). Moreover,

461    organism exposed to condition c showed the lowest IBR values (1.18), with values for

462    conditions A, a, B, b and C (10.9, 2.31, 16.7, 1.45, 7.34 respectively).

463

## 4.    DISCUSSION

The wide proliferation of nanoparticles for different uses may result in their release and dispersion in the environment with potential harmful effects (Katuli et al., 2014; Keller et al., 2010; Guan et al., 2018; Hanna et al., 2013). Moreover, when in the environment, the capacity of nanoparticles to interact with other pollutants may influence their toxicity (Moore et al., 2006; Fabrega et al., 2011). Studies focused on the toxicity of NPs in aquatic environment showed that NPs toxicity depends on their nature, exposure concentration, shape, size, and surface charge (Sun et al. 2016; Jastrzębska and Olszyna 2015) as well as on the time of exposure, medium composition, route of particle administration and target species (Khosravi-Katuli et al. 2017). Regarding the use of NPs for water decontamination, the available information is very scarce, in particular the one devoted to the possible toxic impacts of remediated water, resulting from incomplete removal of contaminants, prevalence of NPs residues, or other changes in water physicochemical properties induced by the remediation condition. Although the adsorption capacity for both organic and inorganic toxicants of various nanoparticles was evaluated and summarized by several authors (Davidescu et al., 2015; Gehrke et al., 2017; Mohan et al., 2007; Aslibeiki et al., 2015), no knowledge on the possible toxicity of the decontaminated water is available. In the present study we assessed the toxicity of magnetic manganese spinel ferrite nanoparticles, $MnFe_2O_4$, which have high capacity to adsorb As from seawater, as well as the efficiency of the treatment from an ecotoxicological point of view, assessing the toxicity of the remediated seawater towards the mussels *Mytilus galloprovincialis*.

### 4.1 Impact of As single exposure, before and after decontamination (conditions A and a)

Regarding As bioaccumulation in the whole soft tissues, as it was expected, the present study demonstrated that the higher contents of As were found in the mussels exposed to condition A (1000 μg/L). Previous studies also demonstrated a direct relationship between As exposure and element accumulation (Velez et al., 2015; Hsiung and Huang, 2004; Celia et al., 2009).

As a consequence of As exposure and bioaccumulation, higher cellular alterations were observed in mussels exposed to the highest As concentration. In particular, the present findings

494  clearly evidenced that mussels exposed to As at a concentration equal to the maximum

495  permissible value for wastewater discharges (1000 μg/L of As, condition A, previous to 24 h

496  decontamination process) strongly decreased their metabolic capacity (preventing energy

497  expenditure), while increasing their antioxidant defences, cellular damages and neurotoxicity.

498  Furthermore, at a smaller scale, the results also demonstrated that mussels exposed to As in a

499  concentration of 70 μg/L (which corresponds to the value achieved by the NPs-based

500  decontamination methodology – condition a), still presented an oxidative stress and neurotoxic

501  status, with inefficient antioxidant capacity that led to observable cellular damages. In particular,

502  the present study demonstrated that seawater contaminated with As at initial (condition A) and

503  final (condition a) concentration levels (1000 and 70 μg/L, respectively) induced biochemical

504  alterations in mussels that resulted in a general oxidative and neurotoxic status, with higher

505  impacts when organisms were exposed to the highest As concentration (condition A). Mussels

506  exposed to As initial concentration (1000 μg/L) clearly reduced their metabolism, preventing the

507  use of energy reserves. However, when exposed to seawater at As concentration equal to that

508  of decontaminated seawater (70 μg/L), albeit minor, the organisms had a metabolic capacity

509  close to those of control indicating that higher impacts on mussels metabolism result from the

510  exposure to the highest As concentration. The decrease of mussels' metabolism may be related

511  to the capacity of bivalves to close their valves and reduce their filtration and respiration rates

512  when exposed to contaminants (Gosling, 2003; Ortmann and Grieshaber, 2003). Previous

513  studies also demonstrated that metals and metalloids even at lower exposure concentrations

514  induced similar metabolic depression in bivalves (Errahmani et al., 2014; Liu et al., 2012; Velez

515  et al., 2017). The present results also demonstrated that mussels exposed to As 1000 μg/L

516  contaminated seawater strongly increased their antioxidant defences, which may result from the

517  overproduction of reactive oxygen species due to the stress induced by As, which were efficient

518  in limiting the occurrence of LPO. Nevertheless, at this condition, mussels clearly revealed

519  oxidative damages with lower GSH/GSSG ratio compared to control organisms and damages in

520  proteins with higher protein carbonylation values compared to control organisms. Under

521  decontaminated seawater (condition a) no cellular damages were observed, evidenced by lower

522  LPO levels in comparison to control values, which may be explained by higher antioxidant (GPx

523  activity) and biotransformation (GSTs activity) defence capacities at this condition.

524 Nevertheless, still oxidative stress was observed in decontaminated seawater, identified by

525 lower GSH/GSSG values in organisms exposed to condition a in comparison to control. These

526 findings are in accordance with previous studies that demonstrated induced of oxidative stress

527 and metabolic depression in bivalves exposed to pollutants (Freitas et al., 2016; Velez et al.,

528 2016; Moreira et al., 2016; Mejdoub et al., 2017; Coppola et al., 2018; Jaishankar et al., 2014;

529 Mandal and Suzuki 2002). In what regards to the neurotoxic impacts, both conditions A and a

530 inhibited AChE activity, with higher injuries when organisms were exposed to the highest As

531 concentration (contaminated seawater). Rajkumar (2013) also showed that As concentrations

532 (80 µg/L) induced neurotoxicity in mussels. A similar pattern was shown by other authors with

533 clams (e.g. *Ruditapes decussatus* and *R. philippinarum*) and oysters (e.g. *Crassostrea gigas*

534 and *C. angulata*) exposed to As contamination (Velez et al., 2015; Freitas et al., 2012; Moreira

535 et al., 2016a; b).

536

537     4.2 Impact of $MnFe_2O_4$ NPs single exposure, before and after

538 decontamination (conditions B and b)

539     In what regards to $MnFe_2O_4$ NPs exposure conditions, the present study demonstrated

540 that seawater contaminated with NPs at initial (condition B, 50 mg $L^{-1}$, previous to

541 decontamination process) and final (condition b, NPs residuals in non-quantifiable

542 concentration, after decontamination process) concentrations induced biochemical alterations in

543 mussels that resulted in metabolism depression and a general oxidative and neurotoxic status,

544 with higher impacts when organisms were exposed to the highest NPs concentration (condition

545 B). In particular, the present findings demonstrated that mussels decreased their metabolic

546 capacity and reduced energy expenditure when exposed to NPs concentration of 50 mg $L^{-1}$,

547 probably because of valves closure to prevent bioaccumulation of NPs and higher injuries, a

548 behaviour observed in bivalves when exposed to stressful conditions (Anestis et al., 2007;

549 Gosling, 2003). Nevertheless, when mussels were exposed to NPs at final concentration ETS

550 activity and energy reserves concentrations were closer to control condition evidencing the

551 capacity of organisms to maintain their metabolism at lower NPs concentrations. No previous

552 studies evaluated the metabolic impacts derived from exposure to $MnFe_2O_4$ NPs, although

553 some works already demonstrated that other NPs (titanium ($TiO_2$), gold (Au) and copper (CuO))

554  decrease bivalves' metabolism (Xia et al., 2017; Cid et al., 2015; Teles et al., 2016; Gomes et

555  al., 2011). Our results also demonstrated that mussels exposed to NPs increased their

556  antioxidant enzymes activity, a response to higher ROS production due to the presence of NPs.

557  It is known that the presence of NPs (TiO2, Au and CuO NPs) increases the production of ROS,

558  which leads to the activation of antioxidant enzymes in bivalves (Xia et al., 2017; Cid et al.,

559  2015; Gomes et al., 2012; Pan et al., 2012).  As a result of increased antioxidant defences in

560  mussels exposed to NPs at concentration of 50 mg L$^{-1}$ damages of the cellular membrane were

561  prevented. Nevertheless, at this condition, mussels clearly revealed oxidative damages with

562  lower GSH/GSSG ratio compared to control organisms and damages in proteins revealed by

563  higher PC values compared to control organisms. When organisms were exposed to residual

564  levels of NPs (condition b) still oxidative damages were observed, with mussels revealing a

565  limited capacity to eliminate the excess of ROS that originated peroxidation of membrane lipids.

566  Such limited antioxidant capacity may result from lower toxicity induced by condition b in

567  comparison to NPs at initial concentration (condition B). These results agree with studies

568  conducted by Tedesco (2010), which also showed that AuNPs (20 mg/L) induced lipid damage

569  in mussels. Regarding the neurotoxic impacts, both NPs conditions (B and b) led to the

570  inhibition of AChE activity, with higher injuries when organisms were exposed to higher NPs

571  concentration (condition B). These results are in line with different studies conducted with

572  diverse NPs: TiO$_2$, 0.4-10 mg L$^{-1}$, AuNPs 80 µg L$^{-1}$ -100 mg L$^{-1}$ (Guan et al., 2018, Pan et al,

573  2012; Teles et al., 2016; Gomes et al., 2011).

574

575  ## 4.3 Impact of As and MnFe$_2$O$_4$ NPs combined exposure before

576  decontamination

577  Concerning the impacts derived from the combined exposure to As and NPs (condition

578  C), the present study demonstrated that initial concentrations of As and NPs (1000 µg L$^{-1}$ and

579  50 mg L$^{-1}$, respectively) reduced mussels' metabolism, increased oxidative stress and

580  neurotoxicity compared to control organisms. In particular, organisms exposed to condition C

581  decreased their metabolic capacity while increasing their energy reserves and increased their

582  antioxidant defences, which were not enough to prevent cellular damages, with lower

583  GSH/GSSG ratio and higher PC values in comparison to control values. This response pattern

584    was similar to those observed in organisms at single exposures (conditions A and B), revealing

585    that the combination of contaminant and nanoparticles did not induce an additive or synergetic

586    response. Although no previous studies showed biochemical stress induced by the combination

587    of As and MnFe$_2$O$_4$ NPs in bivalves, former works demonstrated that NPs and metal(loid)s

588    (such as As) had similar impacts, including metabolism alteration and increased antioxidant

589    defences when bivalves were exposed to combination of both pollutants (De Marchi et al., 2017;

590    Velez et al., 2016a; Monteiro et al., 2018; Della Torre et al., 2015). Nevertheless, the present

591    results are in agreement with studies conducted by Freitas et al. (2018), which showed that

592    functionalized NPs (MWCNTs, 0.1 mg L$^{-1}$) in combination with As (1000 µg L$^{-1}$) induced

593    reduction of metabolic capacity, increase of oxidative stress and lipid damage in mussels, with a

594    similar effect when organisms were exposed to As and NPs separately. Conversely, results

595    obtained from a study conducted with AuNP and cadmium chloride (CdCl$_2$) on *M. edulis* by

596    Tedesco et al. (2010) showed the highest oxidative stress and cellular damage in organism

597    when exposed to these NPs and CdCl$_2$ contamination. Regarding the neurotoxicity activity,

598    As+NPs (condition C) induced reduction of AChE activity, which is in accordance with former

599    studies that analysed this biomarker in different invertebrates' species after exposure to

600    different pollutants such as metals and NPs (Monteiro et al. 2018; Fan et al., 2018; Freitas et

601    al., 2018; Xia et al., 2017; Xiong et al., 2011).

602

603    4.4 Impact of As and MnFe$_2$O$_4$ NPs acting in combination after seawater

604    decontamination

605    The present study demonstrated that organisms exposed to the decontaminated water

606    (condition c, As 70 µg/L and non-quantifiable concentration of NPs) changed their biochemical

607    performance in comparison to control organisms, namely reducing their metabolism, increasing

608    their oxidative stress and neurotoxic status. In comparison to organisms exposed to conditions

609    a and b, where each contaminant was acting individually, the impacts induced were similar, with

610    no significant differences for most of the biomarkers analysed among conditions (a, b, c).

611    Nevertheless, the impacts induced in organisms exposed to decontaminated seawater

612    (condition c) were significantly lower than the impacts observed in organisms exposed to both

613    contaminants at initial concentrations (condition C). In fact, organisms exposed to the

614 decontaminated seawater presented higher metabolism than organisms exposed to the water

615 enriched with As+NPs (condition C). Higher metabolic capacity did not result into higher

616 antioxidant capacity, which probably was not activated due to low stress induced at this

617 condition, originating in turn higher LPO levels and lower GSH/GSSG values at this condition.

618 Furthermore, greater inhibition of AChE was observed when organisms were exposed to

619 condition C compared to condition c, indicating the highest neurotoxic potential of As+NPs initial

620 conditions.

621

622 ## 5. CONCLUSION

623 The present study demonstrated that As decontaminated seawater (condition c) still

624 generates oxidative stress in mussels, with increased cellular damage and oxidative stress in

625 comparison with the control conditions (CTL), but contaminated conditions A, B and C clearly

626 caused higher oxidative stress than the decontaminated seawater (conditions a, b and c) with

627 higher increase in antioxidant defences, neurotoxicity and reduction in metabolism followed by

628 increase of energy reserves. Overall, these results are innovative since, up to our knowledge,

629 no published information is available on the ecotoxic effects induced in mussels when exposed

630 to As contaminated seawater remedied by $MnFe_2O_4$, NPs.

631

651

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**Figure 1.** Transmission Electronic Microscopy image of $MnFe_2O_4$ nanoparticles.

**Figure 2.** A: Electron transport system activity (ETS); B: Glycogen content (GLY); C: Total protein content (PROT) in *Mytilus galloprovincialis* exposed to different conditions (CTL, a, A, b, B, c and C) at the end of the experiment. Results are mean + standard deviation. Significant differences between conditions A vs a, B vs b, C vs c are presented with asterisks.

**Figure 3**. A: Superoxide dismutase activity (SOD); B: glutathione peroxidase activity (GPx); C: Glutathione S-transferases activity (GSTs), in *Mytilus galloprovincialis* exposed to different conditions (CTL, a, A, b, B, c and C) at the end of the experiment. Results are mean + standard deviation. Significant differences between conditions A vs a, B vs b, C vs c are presented with asterisks.

**Figure 4.** A: Lipid peroxidation levels (LPO); B: protein carbonyl levels (PC); C: ratio between reduced and oxidized glutathione (GSH/GSSG), in *Mytilus galloprovincialis* exposed to different conditions (CTL, a, A, b, B, c and C) at the end of the experiment. Results are mean + standard deviation. Significant differences between conditions A vs a, B vs b, C vs c are presented with asterisks.

**Figure 5.** Acetylcholinesterase activity (AChE), in *Mytilus galloprovincialis* exposed to different conditions (CTL, a, A, b, B, c and C) at the end of the experiment. Results are mean + standard deviation. Significant differences between conditions A vs a, B vs b, C vs c are presented with asterisks.

**Table 1.** Experimental conditions.

| CONDITIONS | | DESCRIPTION |
|---|---|---|
| CTL | | Seawater with As 0 µg $L^{-1}$ + NPs 0 mg $L^{-1}$ |
| Water before As decontamination | A | Seawater with As 1000 µg $L^{-1}$ |
| | B | Seawater with NPs 50 mg $L^{-1}$ |
| | C | Seawater with As 1000 µg $L^{-1}$ and NPs 50 mg $L^{-1}$ |
| Water after As decontamiantion | a | Seawater with As 70 µg $L^{-1}$ |
| | b | Seawater after 24h in contact with NPs (50 mg $L^{-1}$), which were afterwards separated from seawater |
| | c | Seawater previously contaminated with As (1000 µg $L^{-1}$), then remediation using NPs (50 mg $L^{-1}$) during 24 h (which were afterwards separated from seawater). |

**Table 2.** Aggregation of NPs MnFe2O4 in seawater (nm), at different time (T0, T1, T24) after the beginning of the experiment.

| Time (hours) | MEAN (nm) | SD | CV% |
|---|---|---|---|
| **T0** | 3987 | 614 | 15 |
| **T1** | 14045 | 498 | 35 |
| **T24** | 67013 | 152 | 23 |

**Table 3.** Arsenic concentration ($\mu g\ L^{-1}$) measured in water samples collected immediately after the weekly water renewal. Results correspond to the mean value and standard deviation of the four weeks.

| [As] water µg L$^{-1}$ | | |
|---|---|---|
| CTL | | <1.5 |
| As | **A** | 947 ± 17 |
| | **a** | 82 ± 15 |
| NP | **B** | <1.5 |
| | **b** | <1.5 |
| As + NP | **C** | * |
| | **c** | 55 ± 13 |

*Because sorption of As by the NPs is extremely rapid, its quantification in this condition was not performed.

**Table 4.** Arsenic concentration in mussels (mg Kg$^{-1}$), 28 days after the beginning of the experiment. Concentrations were measured in organisms from different conditions: (CTL, a, A, b, B, c and C). Asterisks represent differences between A vs a, B vs b and C vs c conditions, while different lowercase letters represent differences between CTL vs a, CTL vs b, CTL vs c and uppercase CTL vs A, CTL vs B, CTL vs C conditions.

As concentration (mg Kg$^{-1}$)

| | | | |
|---|---|---|---|
| **CTL** | | $7.4\pm1.5^{A,a}$ | |
| **As** | A | $12\pm2.6^{B}$ | * |
| | a | $6.8\pm2.2^{a}$ | |
| **NP** | B | $5.2\pm0.9^{A}$ | |
| | b | $4.4\pm0.2^{a}$ | |
| **As+NP** | C | $11\pm2.7^{B}$ | * |
| | c | $6.8\pm2.2^{a}$ | |

**Table 5.** *p*-values obtained by pair-wise comparisons between conditions (CTL vs A, CTL vs B, CTL vs C CTL vs a, CTL vs b, CTL vs c, A vs B, A vs C, B vs C a vs b, a vs c and b vs c) for each biomarker: ETS, electron transport system activity; GLY, glycogen content; PROT, total protein content; SOD, superoxide dismutase activity, GPx, glutathione peroxidase activit; GSTs, glutathione S-transferases activity; LPO, lipid peroxidation levels; PC, protein carbonyl levels; glutathione ratio, GSH/GSSG; acetylcholinesterase activity, AChE. Significant differences (*p* ≤ 0.005) are highlighted in bold.

| | ETS | GLY | PROT | SOD | GPx | GSTs | LPO | CP | GSH/GSSG | AChE |
|---|---|---|---|---|---|---|---|---|---|---|
| **CTL vs A** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** |

| | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|---|
| **CTL vs B** | **0.0001** | **0.0001** | 0.2061 | **0.0001** | **0.0001** | **0.0006** | **0.0001** | **0.0001** | **0.0001** | **0.0001** |
| **CTL vs C** | **0.0001** | **0.0001** | **0.0003** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** | **0.0001** |
| **CTL vs a** | **0.0001** | 0.0550 | 0.2100 | 0.1983 | **0.0003** | 0.0707 | **0.0003** | **0.0293** | **0.0045** | **0.0093** |
| **CTL vs b** | **0.0001** | 0.9800 | 0.0770 | 0.2334 | **0.0168** | 0.8774 | **0.0020** | **0.0448** | **0.0003** | **0.0018** |
| **CTL vs c** | **0.0001** | 0.1936 | 0.0609 | 0.1310 | **0.0001** | **0.0054** | **0.0021** | 0.2694 | **0.0026** | **0.0043** |
| **A vs B** | **0.0076** | **0.0001** | 0.9018 | **0.0002** | 0.1127 | **0.0001** | **0.0001** | **0.0001** | 0.2629 | 0.8423 |
| **A vs C** | **0.0438** | **0.0017** | **0.0032** | **0.0009** | **0.0055** | 0.3656 | 0.7131 | 0.8632 | 0.0868 | 0.9424 |
| **B vs C** | 0.5399 | 0.9497 | **0.0331** | 0.4324 | 0.1733 | **0.0001** | **0.0001** | **0.0001** | 0.9478 | 0.7912 |
| **a vs b** | 0.1146 | 0.8939 | 0.3760 | 0.759 | 0.1757 | **0.0013** | 0.7522 | 0.8243 | **0.0298** | 0.3500 |
| **a vs c** | 0.1934 | 0.6053 | 0.1222 | 0.4063 | **0.0384** | 0.0694 | 0.066 | 0.1524 | 0.9786 | 0.4114 |
| **b vs c** | 0.3265 | 0.4136 | 0.4084 | 0.2369 | **0.0006** | **0.0001** | 0.1131 | 0.2086 | **0.0188** | 0.9893 |

100 nm

**A**

## ETS



**B**

## GLY



**C**

## PROT

**A** SOD

**B** CAT

**C** GSTs

**A**

**LPO**



**B**

**PC**



**C**

**GSH/GSSG**

## AChE
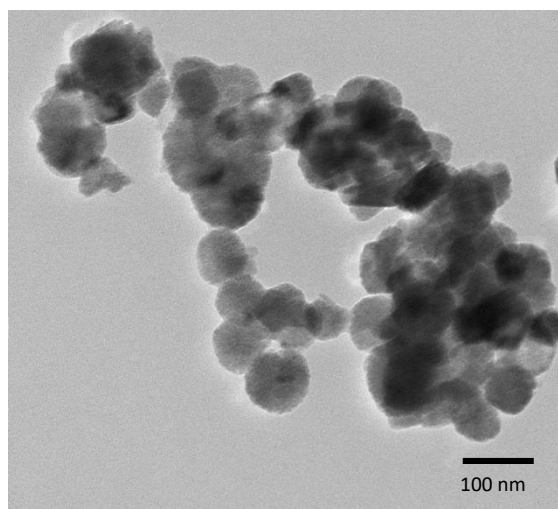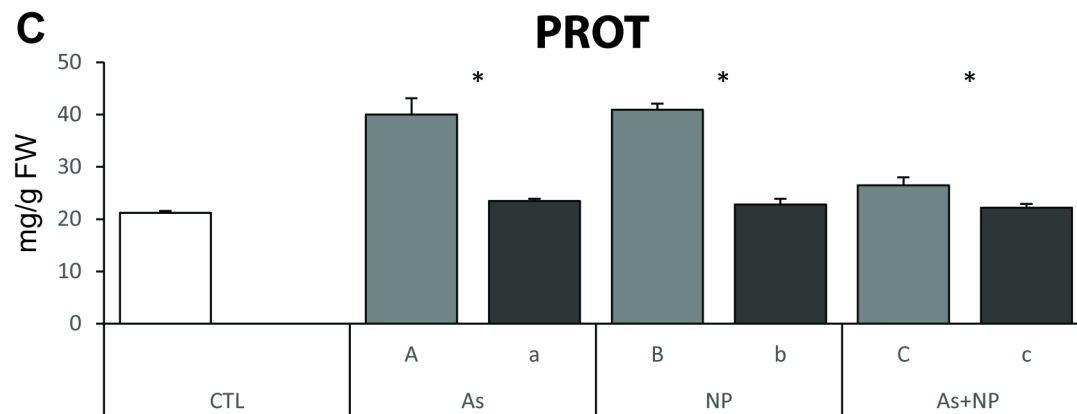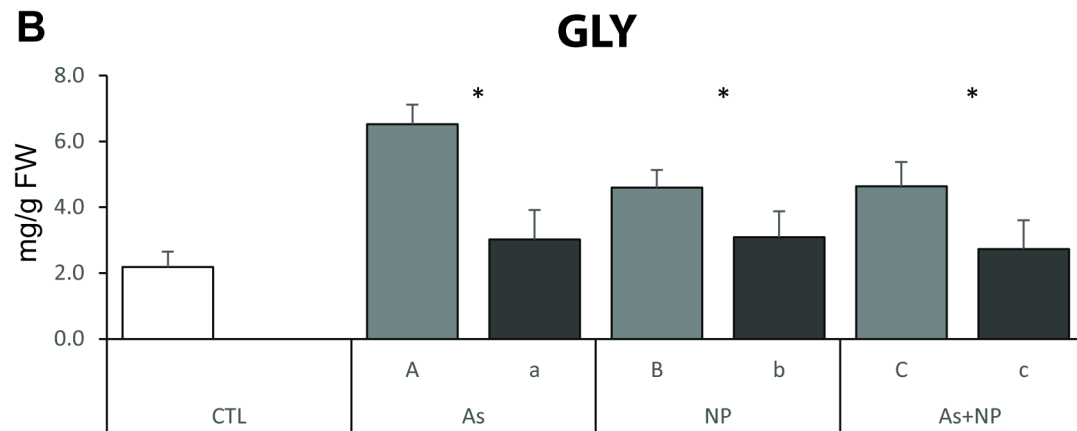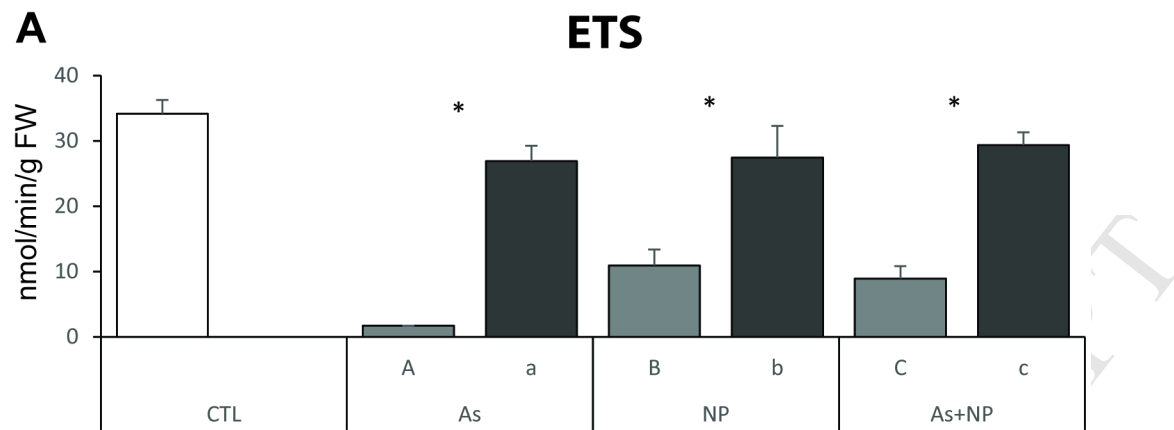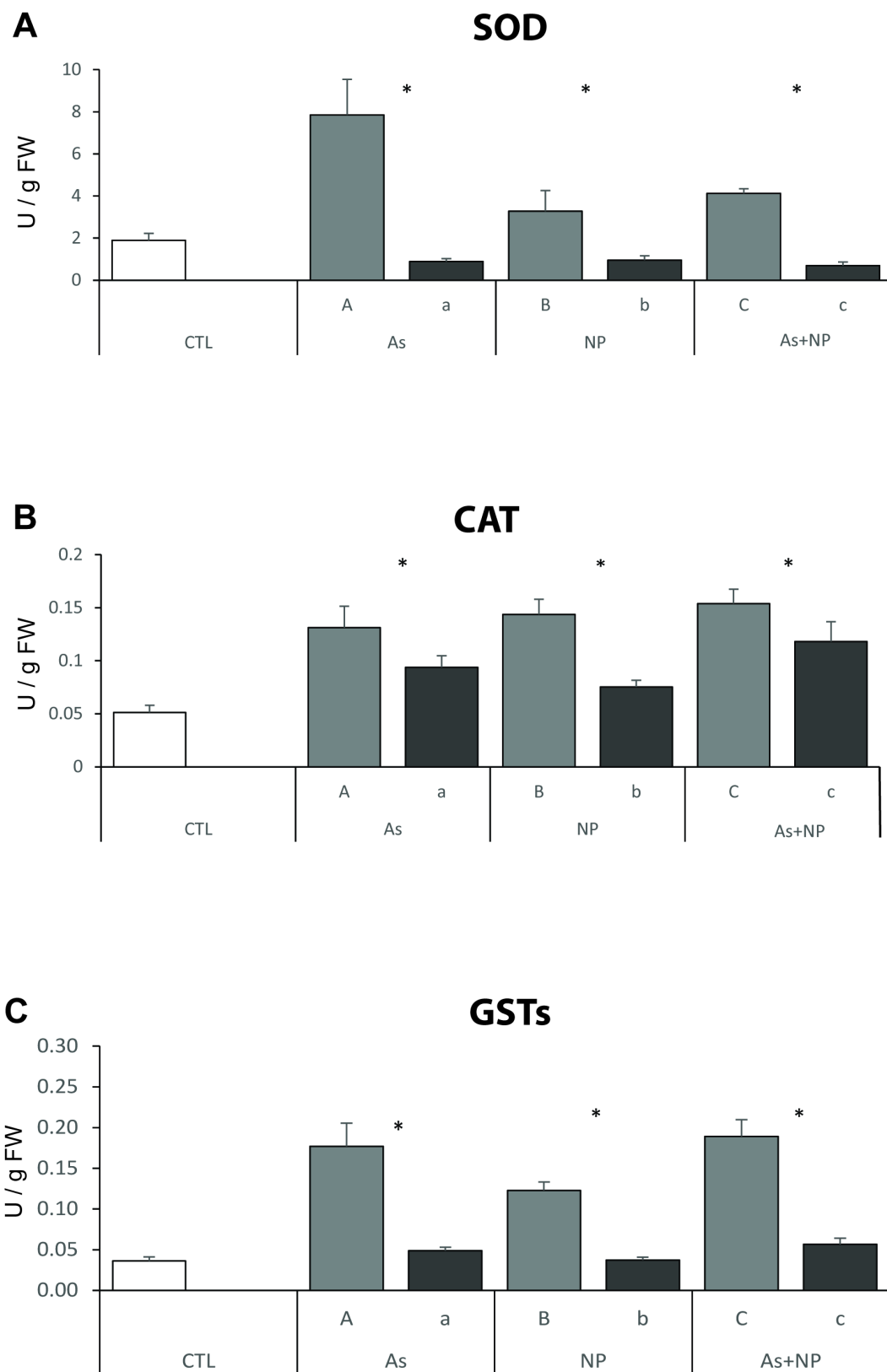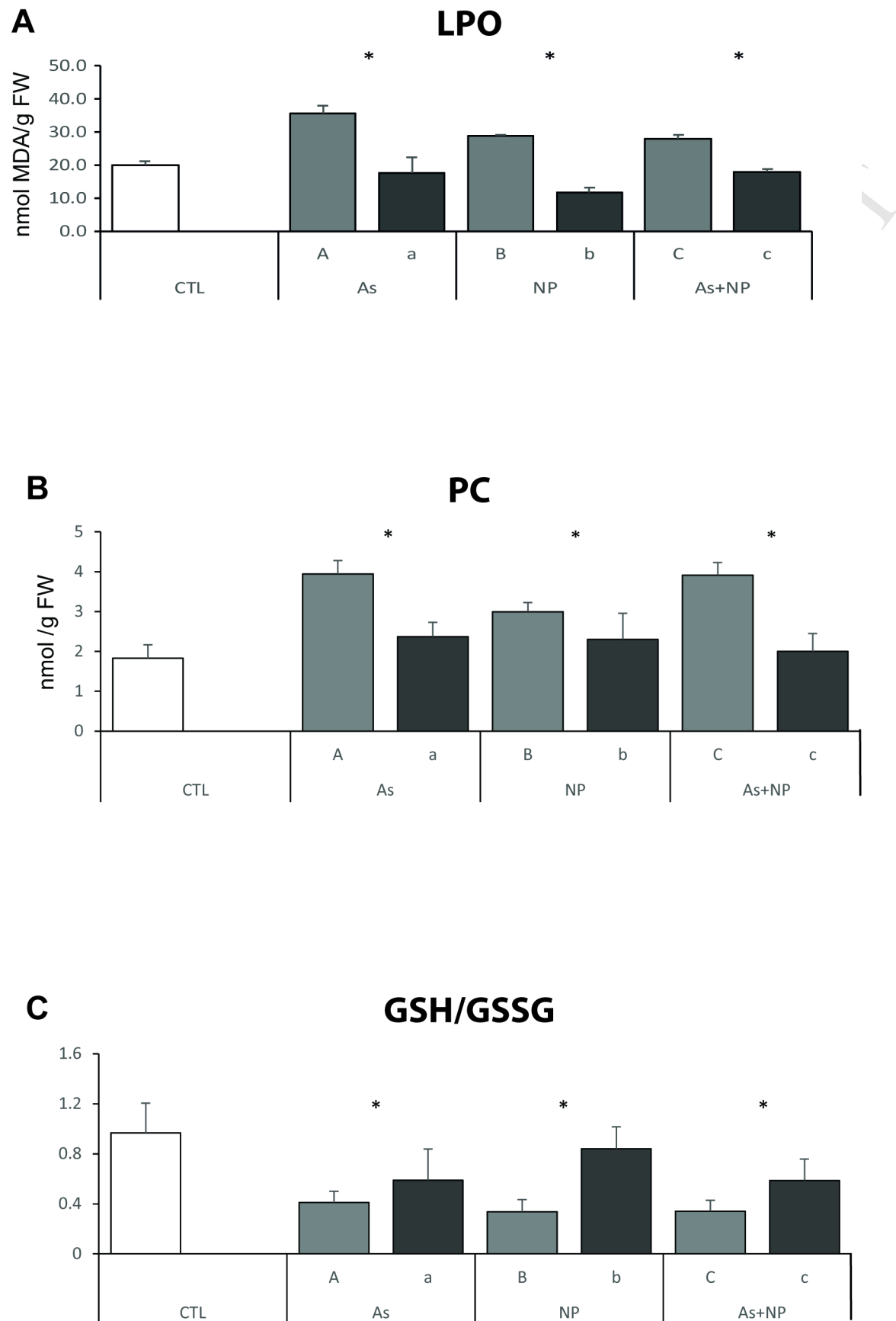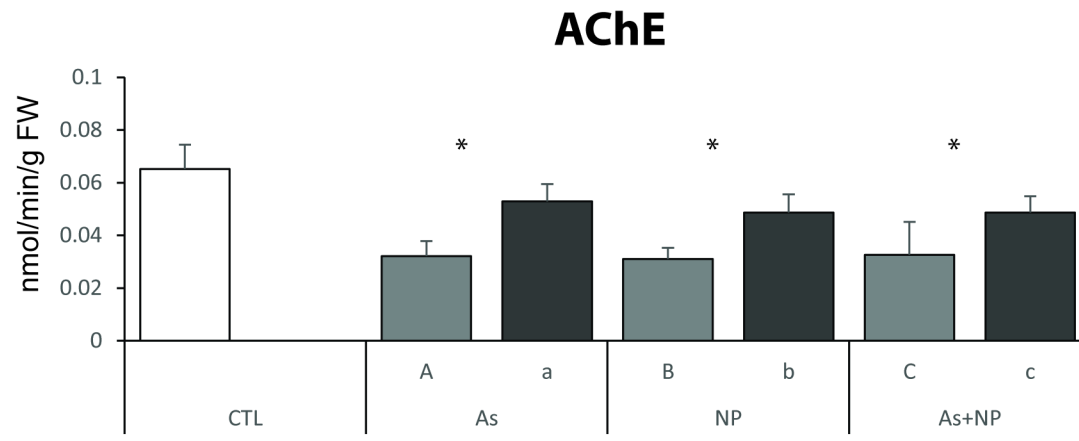
- Decontaminated seawater did not affect mussels metabolic capacity

- Contaminated mussels enhanced their antioxidant and biotransformation enzymes activities

- No cellular damages were observed in mussels exposed to decontaminated seawater

- Neurotoxicity was induced in contaminated mussels