

Metal body burden and tissue oxidative status in the bivalve *Venerupis decussata* from Tunisian coastal lagoons

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HIGHLIGHTS

- The health status of bivalves reflects natural and anthropogenic stressors
- Boughrara and South Tunis lagoons are the most impacted transitional waters
- Higher temperature and metal content enhance oxidative and toxicopathological traits
- Multivariate PCA discriminates environmental status of Tunisian transitional waters

1 **ABSTRACT**

2 Coastal transitional waters are exposed to many anthropogenic threats. This study aims
3 to assess the trace metals' pollution status of transitional waters by evaluating its
4 biological effects in the clam *Venerupis decussata*. Among the studied sites along the
5 Tunisian littoral, South Tunis and Boughrara were the most impacted, since clams from
6 these two lagoons presented significant differences in: (i) trace metal contents, (ii) in-
7 cell hydrogen peroxide, (iii) enzymatic and non-enzymatic defenses, (iv) damage to
8 lipids and proteins, and (v) protein post-translational modifications. These changes
9 related to evident histopathological traits. PCA showed a clear separation between the
10 digestive gland and gills tissues and illustrated an impact gradient in Tunisian coastal
11 lagoons. Water temperature was revealed as an added natural stressor that, when
12 concurring with high pollution, may jeopardize an ecosystem's health and contribute to
13 the accumulation of hazardous metals in organisms.

14

15 *Keywords:*

16 Clams; Histopathology; Molecular biomarkers; Oxidative stress; Temperature; Trace
17 metals; Transitional waters

18

19 *Abbreviations:* AOPP, advanced oxidation protein product; BG, Boughrara Lagoon; BZ,
20 Bizerte Lagoon; CAT, catalase; DNPH, 2,4-dinitrophenylhydrazine; EL, Ellouza; GPx,
21 glutathione peroxidase; GSH, reduced glutathione; mBBr, monobromobimane; MDA,
22 malondialdehyde; NPSH, non-protein thiols; NT, North Tunis Lagoon; PAHs,
23 polycyclic aromatic hydrocarbons; PCA, principal component analysis; PCO, protein
24 carbonyl; PRS, protein redox state; ROS, reactive oxygen species; SOD, superoxide

- 25 dismutase; SPM, Suspended particulate matter; ST, South Tunis Lagoon; TBP,
26 tributylphosphine; TCA, trichloroacetic acid.

27 **1. Introduction**

28 Transitional waters are those situated between the land and the sea, including
29 fjords, estuaries, lagoons, deltas and rias. Due to their strategic location, coastal lagoons
30 usually exert vast ecological and economic impacts on their surrounding areas.
31 However, lagoons are characterized by being relatively isolated from the open sea,
32 which makes them highly vulnerable to anthropogenic pollution, as they are exposed to
33 urbanization, industrialization and intensive agricultural activities (Alves Martins et al.,
34 2015; Bilgin and Uluturhan-Suzer, 2017). The Tunisian coastline contains several of
35 these ecosystems that are exploited for fishery and shellfish farming, particularly
36 Bizerte, North and South Tunis, and Boughrara lagoons (Alves Martins et al., 2015;
37 Lahbib et al., 2018).

38 As a consequence of an intense human activity, transitional water-bodies
39 accumulate increasing amount of toxic chemicals, most of them highly persistent (Bald
40 et al., 2005; Carreira et al., 2013). As marine pollution increases worldwide, there is a
41 concomitant need to develop strategies to monitor the biological effects of
42 contaminants.

43 Ecosystem pollution can be assessed by using sentinel organisms as
44 bioindicators. Bivalve mollusks are used worldwide as marine pollution sentinels due to
45 their sessile nature, filter-feeding habits, and ability to concentrate pollutants (Bebiano
46 et al., 2004; Costa et al., 2013; Chalghmi et al., 2016a; Funes et al., 2006). The
47 *Venerupis decussata* clam (Linnaeus, 1758) (also known as *Ruditapes decussatus* and
48 *Tapes decussatus*) lives in the muddy sand sediments of coastal environments (Parache,
49 1982). *V. decussata* is widely used in fisheries and aquaculture in southwestern Europe
50 and Mediterranean areas, where this mollusk has a high economic impact (Carreira et
51 al., 2013). Due to its wide distribution, and sensitivity to contaminants, *V. decussata* is

52 also a common bioindicator of environmental pollution, particularly in confined coastal
53 environments (Bebianno et al., 2004; Costa et al., 2013).

54 Trace metals constitute one of the most dangerous sources of pollution in marine
55 ecosystems due to their toxicity, persistence, and non-degradability (Park et al., 2017).
56 As many other toxicants, a well-described effect of exposure to transition metals is
57 oxidative stress, by catalyzing the generation of highly hazardous and mutagenic
58 reactive oxygen species (ROS), such as: hydroxyl radical (HO·), superoxide anion
59 radical ($O_2^{\cdot-}$) or hydrogen peroxide (H_2O_2) (Park et al., 2017; Roberts et al., 2009; Sies,
60 1986). Organisms respond to oxidative threats by activating protective mechanisms that
61 include enzymatic and non-enzymatic antioxidants. As first line defense antioxidants,
62 superoxide dismutase (SOD) catalyzes the dismutation of $O_2^{\cdot-}$ to H_2O_2 , while catalase
63 (CAT) and glutathione peroxidase (GPx) convert H_2O_2 into water and diatomic oxygen.
64 GPx also reduces reactive lipid-hydroperoxides using reduced glutathione (GSH), thus
65 avoiding the formation of malondialdehyde (MDA) (Funes et al., 2006; López-Barea,
66 1995; Rodriguez-Ortega et al., 2002; Sies, 1986). Thiol-containing compounds are a
67 very important class of non-enzymatic antioxidants. Among low-Mr non-proteins thiols,
68 GSH plays a pivotal protective role against electrophilic xenobiotics and oxidative
69 stress (Pompella et al., 2003). The antioxidant properties of ascorbic acid (vitamin C)
70 are attributed to its capacity both to directly interact with oxidizing radicals, and to
71 regenerate other small antioxidant molecules, including GSH (Ahmad et al., 2012;
72 Gebicki, 2016). Among the thiol-containing proteins, metallothioneins (MTs) stand out
73 as very sensitive metal-pollution biomarkers. MTs are small proteins with a high Cys
74 content involved in the homeostasis of essential metals, sequestration and detoxification
75 of trace metals, and scavenging of ROS (Amiard et al., 2006; Romero-Ruiz et al., 2008;
76 Viarengo et al., 1999). When the ROS production rate exceeds that of the quenching

77 antioxidant detoxification mechanisms, damage can reach different biomolecules
78 impairing many physiological processes (Park et al., 2017; Sies, 1986). MDA is a
79 highly reactive by-product of lipid peroxidation (Draper and Hadley, 1990), while
80 formation of irreversible carbonyl groups (PCO) and advanced oxidation protein
81 products (AOPP) are major forms of protein oxidation (Kalousova et al., 2005; Kayali
82 et al., 2006; Yan and Forster, 2011). Because of their abundance and high reactivity,
83 proteins are the main targets in oxidative stress scenarios since they absorb around 70%
84 of ROS (Davies, 2005). As a consequence of ROS exposure, proteins are regulated by
85 post-translational modifications (PTMs) that influence their structural conformation,
86 biological activity, protein-protein interactions, turnover rates, and targeting to
87 subcellular localizations (Cabiscol and Ros, 2006; Eaton, 2006; Klomsiri et al., 2011;
88 Sheehan et al., 2010).

89 Changes at the molecular level constitute rapid, early-warning and reversible
90 responses to environmental alterations, so they have high toxicological relevance
91 (López-Barea, 1995). At a higher level, histopathological analyses in biomonitoring
92 programs provide additional relevant information on the health status of the organisms
93 (Costa et al., 2013).

94 Besides water and sediment pollution, mainly of anthropogenic origin, marine
95 organisms are also exposed to natural stressors, such as: fluctuations in temperature,
96 salinity or pH, desiccation, pathogens, and changes in O₂ or CO₂ levels. Synergistic
97 interactions between the effects of various natural stressors and pollutants are the most
98 common phenomena (Holmstrup et al., 2010; Sokolova and Lannig, 2008).

99 The aim of the present study is to develop a multivariate approach for the
100 evaluation of the trace metal contamination status of transitional waters, the generated
101 oxidative stress, and its biological effects. The clam *V. decussata* has been selected as

102 bioindicator, and several lagoons along of the Tunisian coastline were chosen as case
103 studies. To implement this approach, we have carried out: (i) physico-chemical
104 measurements in waters, and (ii) trace metal content, (iii) reactive oxygen species
105 generation, (iv) enzymatic and non-enzymatic antioxidative defenses, (v) oxidative
106 damage to biomolecules, (vi) global post-translational modifications, and (vii)
107 histopathological analysis in clam tissues.

108 **2. Materials and methods**

109 *2.1. Areas of study*

110 Widely distributed areas of bivalve mollusks production along the Tunisia
111 coastline have been studied in this work (Fig. 1). Ellouza (EL), located in the western
112 coast of the Gabes gulf (Sfax city), was used as the reference site because previous
113 studies have shown that it is a low impacted open-sea coastal area (e.g.: Annabi et al.,
114 2013; Chalghmi et al., 2019; Kessabi et al., 2012). The other sites studied are lagoons,
115 closed or semi-closed transitional water ecosystems with a great ecological value
116 (Bilgin and Uluturhan-Suzer, 2017; Carreira et al., 2013). At the same time, lagoons
117 might be subjected to a high anthropogenic pressure. Thus, the Bizerte Lagoon (BZ),
118 with an extension of over 128 Km² and an average depth of 7 m, is surrounded by many
119 industries (oil and its derivatives, plastic, clothing, agri-food, etc.), agricultural activities
120 and urban areas (Alves Martins et al., 2015; Bejaoui et al., 2017). The Tunis Lagoon
121 (40 Km²) is divided into two parts, North (NT) and South (ST), separated by the Tunis-
122 Goulette road and a navigation channel. NT receives urban discharges daily, while ST is
123 mostly affected by fishing and navigation, port and industrial activities (Chalghmi et al.,
124 2019). The Boughrara Lagoon (BG) is an exceptional site for both terrestrial and marine
125 biodiversity, although its ecological future is in jeopardy. This lagoon is characterized
126 by high salinity and temperature (Khedhri et al., 2017). BG is strongly impacted by
127 many activities such as aquaculture farms and fishing boat traffic, thus receiving an
128 important amount of organic matter. Moreover, high levels of phosphogypsum have
129 been discharged into this lagoon, disturbing its quality, biodiversity and all ecosystem
130 functions (Rekik et al., 2012).

131 *2.2. Physicochemical analysis*

132 Surface water samples were analyzed *in situ* at the different sites where clams were
133 collected. Temperature, salinity, pH and suspended particulate matter were measured in
134 February 2017, using a WTW multi-parameter probe (model WTW LF.325).

135 2.3. *Specimens and sample preparation*

136 *Venerupis decussata* clams were collected in February 2017 at the different
137 sampling sites shown in Fig. 1. Animals were immediately transferred alive to the
138 laboratory on ice and selected those with similar shell dimensions and weight
139 (Supplementary Table 1). The specimens were briefly rinsed with ultrapure water to
140 eliminate any exogenous materials, and then dissected on ice for excision of the
141 digestive gland and gills.

142 For metal analysis, the tissues of twenty clams per sampling site were pooled
143 into six independent replicates and freeze-dried. Fresh tissue samples were excised for
144 histopathology and fixed immediately, as described below (Section 2.9). For
145 biochemical analyses, individual tissues were snap-frozen, ground with a Freezer/Mill®
146 Grinder (SPEX Sample PreP) and stored at -80°C until pooled in eight independent
147 samples (twenty clams per pool). After that, the pooled samples were homogenized in
148 50 mM Tris-HCl buffer (pH 7.4, 5 mL/g) containing 1 mM EDTA and 1 mM PMSF,
149 using an electrically driven Potter-Elvehjem glass homogenizer and a Teflon pestle. The
150 homogenates were then centrifuged at 16,200 g for 15 min. The remaining supernatants
151 were aliquoted and stored at -80°C until used for the biochemical analyses. The protein
152 content was determined in the supernatants by dye-binding (Bradford, 1976), using
153 bovine serum albumin (BSA) as standard.

154 2.4. *Trace element analysis*

155 Freeze-dried organ samples (0.5 g) were digested in a mixture of 6 mL of HNO₃
156 and 1 mL of H₂O₂ in Teflon reactors using a programmable microwave Touch Control
157 Terminal 320 system (Milestone, type Ethos). The first 5 min consisted of a heating
158 ramp up to 175 ± 5°C at 1200 W. After heating, the residue obtained was re-dissolved
159 in 5 mL of HNO₃, and the solution heated for a further 30 min at 150°C. After cooling
160 to room temperature, the solutions were transferred to new flasks and their final volume
161 was adjusted to 50 mL with ultrapure water. Concentrations of cadmium (Cd), copper
162 (Cu), lead (Pb), zinc (Zn), iron (Fe), nickel (Ni) and aluminum (Al) were determined by
163 Inductively Coupled Plasma-Mass Spectrometry (Triple Quadrupole ICP-MS, Thermo
164 Fisher), including blanks and internal standards. The metal contents are expressed as µg
165 g⁻¹ sample dry weight. The reference material NIST SRM 2976 (trace elements and
166 methylmercury in mussel tissue) was analyzed using the same procedure, with recovery
167 rates being found to be within 95% of the certified range. The quality control of
168 standard of each tested element was as follows: Cd, 0.81 ± 0.11; Cu, 4.01 ± 0.31; Pb,
169 1.18 ± 0.15; Zn, 138 ± 12; Fe, 171.0 ± 4.9; Ni, 0.94 ± 0.10; Al, 134 ± 34 µg g⁻¹.

170 2.5. *Measurement of hydrogen peroxide*

171 The concentration of H₂O₂ in the digestive gland and gills of *V. decussata* was
172 determined spectrophotometrically at 560 nm following the Ferrous Oxidation in
173 Xylenol orange (FOX) reaction (Ou and Wolff, 1996). Standard solutions of H₂O₂ were
174 used to calibrate the assay.

175 2.6. *Enzymatic and non-enzymatic antioxidant defenses*

176 Enzymatic and non-enzymatic antioxidants were determined to evaluate the
177 defensive responses to oxidative stress. Cuvette-based colorimetric assays were used for
178 the different determinations. Catalase (CAT) was determined by recording the
179 breakdown of H₂O₂ at 240 nm (Aebi, 1984). Glutathione peroxidase (GPx) was

180 measured using GSH as a conjugation substrate coupling the assay to NADPH
181 consumption, determined at 340 nm (Flohe and Gunzler, 1984). Superoxide dismutase
182 (SOD) was determined spectrophotometrically at 580 nm (Beauchamp and Fridovich,
183 1971), based on the ability of superoxide dismutase to inhibit the reduction of nitro-blue
184 tetrazolium (NBT) by superoxide anion.

185 Due to the relevant role of thiolic compounds as antioxidants, they were
186 quantified as metallothionein-like proteins (MTLPs) after precipitating proteins from
187 the extracts used for biochemical analysis (Section 2.3), and as non-protein thiols
188 (NPSH) on the remaining soluble cytosol. MTLPs and NPSH were determined
189 spectrophotometrically at 412 nm (Ellman, 1959; Petrovic et al., 2001; Viarengo et al.,
190 1999). For both, the concentration was estimated using GSH as a reference standard.
191 Vitamin C (Vit C) was determined at 529 nm by the DNPH and copper sulfate assay
192 (Jacques-Silva et al., 2001). The ascorbic acid concentration was estimated from a
193 previously prepared standard curve.

194 2.7. *Biochemical oxidative damage*

195 The level of thiobarbituric acid-reactive substances was assayed to assess
196 malondialdehyde (MDA) content, a known marker of lipid peroxidation (Draper and
197 Hadley, 1990). The levels of MDA were determined spectrophotometrically at 532 nm,
198 according to a standard curve generated from serial dilutions of 1,1,3,3-
199 tetraethoxypropane (TEP), an MDA precursor.

200 The oxidative damage to proteins was evaluated by determining advanced
201 oxidation protein products (AOPP) and protein carbonyl (PCO) levels. The AOPP
202 levels were determined spectrophotometrically at 340 nm (Kayali et al., 2006). PCO
203 was determined by a method based on the reaction of carbonyls groups with 2,4-

204 dinitrophenylhydrazine (DNPH), and determined at 370 nm (Reznick and Packer,
205 1994).

206 2.8. *Electrophoresis-based evaluation of post-translational modifications*

207 All protein separations were performed by SDS-polyacrylamide gel
208 electrophoresis on 4% (w/v) stacking gel and 12% resolving gel using the Laemmli
209 buffer system. The samples were separated at 200 V constant in a Mini-PROTEAN 3
210 Cell (Bio-Rad, Hercules, CA, USA). After electrophoresis, the gels were scanned using
211 a ChemiDoc™ MP Imaging System (Bio-Rad). Image Lab software (Bio-Rad) was used
212 for the acquisition of gel images and all subsequent image analyses.

213 The Pro-Q® Diamond gel stain (Molecular Probes®, Invitrogen) was used to
214 selectively stain the phosphoproteins in polyacrylamide gels. 50 µg of protein was
215 analyzed by SDS-PAGE and, after electrophoresis, the manufacturer's
216 recommendations were followed for the specific staining with Pro-Q Diamond of the
217 phosphoproteins in the gels, that were scanned at 555/580 nm excitation/emission
218 detection wavelengths to show the phosphorylated proteins. The gels were then stained
219 with SYPRO Ruby (Bio-Rad) and rescanned at 532/555 nm excitation/emission
220 detection to show total proteins. Determining the ratio of Pro-Q Diamond dye to
221 SYPRO Ruby dye signal intensities provided a measure of the phosphorylation level
222 normalized to the total amount of protein.

223 A fluorescence-based electrophoretic assay was used to quantify both the global
224 level of reversibly oxidized and reduced thiols in proteins whose ratio indicates their
225 redox status. Cysteiny protein thiols were directly labeled with monobromobimane
226 (mBBBr) for native reduced groups. For oxidized thiol detection, all free thiol groups
227 were first blocked with iodoacetamide (IAM), reversibly oxidized thiols were then
228 reduced with tributylphosphine (TBP), and finally the reduced thiols generated were

229 labeled with mBBr. For the reduced thiol groups, 100 µg protein was denatured by
230 incubating for 10 min at 37°C in 50 µl Tris-HCl 0.1 M, pH 7.4, and 1% SDS. mBBr
231 was then added to 200 µM final concentration and the samples were incubated for 30
232 min at 37°C in the dark. After adding Laemmli sample buffer, the proteins were loaded
233 and separated on 12% SDS-PAGE gels, avoiding light. To minimize the background,
234 the gels were washed with ethanol:acetic acid (50:3) before scanning at 395/490 nm
235 excitation/emission detection wavelength to show the fluorescent labeled proteins. For
236 normalization, the gels were re-stained with Coomassie blue, and then the image
237 analysis was carried out as indicated above. For determination of the oxidized thiol
238 groups, the initial reduced groups in 200 µg protein extracts were blocked by incubating
239 for 30 min at 37°C in Tris-HCl 0.1 M, pH 7.4, 1% SDS and 200 mM IAA. The proteins
240 were precipitated with cold 20% TCA (w:v) followed by three washes with
241 ethanol:ethyl acetate (50:50, v:v) to remove excess IAA. The final pellets were
242 resuspended in Tris-HCl 0.1 M, pH 7.4, 1% SDS and 1 mM TBP to reduce the oxidized
243 thiols, and then the procedure continued as for the initially reduced thiol groups.
244 Approximately 20 µg of protein was loaded on the gels for both the reduced and
245 oxidized thiol determinations.

246 2.9. *Histopathological analysis*

247 The preparation of the samples (5 gills and 6 digestive glands per site) for the
248 histopathological appraisal follows the standard methods previously described (Martoja
249 and Martoja-Pierson, 1967). In brief, tissue samples were fixed in 10% neutral-buffered
250 formalin and embedded in paraffin. Sections (5 µm) were stained with hematoxylin and
251 eosin (H&E). Histopathological lesions and alterations were identified according to
252 specific literature on *V. decussata* histology and histopathology (Costa et al., 2013).

253 2.10. *Statistical analysis*

254 Data are expressed as mean \pm SD of six replicates for the trace elements, eight
255 for the other biochemical analyses, and three for the post-translational modifications and
256 physico-chemical parameters. The results were analyzed using the Statistica software
257 (Version 8). The normality of the data was evaluated using the Kolmogorov–Smirnov
258 and the Shapiro–Wilk *W* tests, while homogeneity of variance was verified using
259 Levene’s test. The comparison between the groups was made through one-way
260 ANOVA followed by Tukey’s test (for normal data), or through Kruskal-Wallis test (for
261 non-normal data). Principal component analysis (PCA) was performed using R (Ihaka
262 and Gentleman, 1996). The Spearman correlation matrix was also calculated to study
263 the relationship between the different parameters analyzed using the GraphPad Prism
264 8.0.2 software.

265 **3. Results**

266 *3.1. Physico-chemical characteristics of the transitional waters*

267 Physico-chemical characteristics were determined in the surface water at the
268 different locations (Table 1). Although no significant variations were observed in
269 salinity and pH, compared to the EL reference site, marked increases were observed in
270 temperature and suspended matter. Thus, the temperature was significantly higher in ST
271 ($p < 0.05$) and BG ($p < 0.01$), while the concentration of suspended particulate matter
272 greatly increased at all the lagoons studied ($p < 0.001$).

273 *3.2. Accumulation of trace metals in clam tissues*

274 Table 2 shows the results of the metal concentrations in the gills and in the
275 digestive gland of clams collected from five locations along the Tunisian coastline. The
276 lowest metal concentrations were found in the tissues from specimens collected at
277 Ellouze (EL), considered here as the reference site. Otherwise, the highest metal levels
278 were found at ST and BG, where the differences were statistically significant for Al
279 (gills), Zn and Fe (digestive gland), and Cd, Pb, Cu and Ni (both tissues). Compared to
280 the reference site, we wish to highlight the ≥ 2 -fold increase in Pb, Cu and Ni (gills),
281 Cd, Pb and Ni (digestive gland) contents at BG, and in Pb (digestive gland) and Ni
282 (both tissues) levels at ST. Even though the ST and NT water bodies are connected, the
283 ST specimens accumulated more metals than those from NT, where only Ni (gills) and
284 Pb (digestive gland) increased significantly. Finally, clam tissues from BZ showed
285 levels slightly superior to the reference site, although the differences were of little
286 significance ($p < 0.05$) for Ni in gills, and Pb in both tissues.

287 *3.3. Oxidative stress and antioxidant defensive responses in clam tissues*

288 Table 3 shows the levels of H₂O₂ determined in the gills and in the digestive
289 gland of clams. In both tissues, significantly higher levels of H₂O₂ were detected in the
290 clams from BZ, ST and BG compared to EL, the reference site.

291 Table 3 shows the levels of enzymatic and non-enzymatic antioxidative
292 responses in the gills and in the digestive gland of *V. decussata* clams. In agreement
293 with the accumulation of trace metals in their tissues, the clams from BG and ST
294 presented significant increments in all the antioxidative responses in both tissues,
295 compared to the EL reference site. The most relevant changes were found for CAT
296 activity in ST and BG (around 70% increase in both tissues), for GPx in the digestive
297 gland, which increased 46% (ST) and 84% (BG), and for MTLPs in the gills, with a
298 91% and 74% rise at BG and ST, respectively. This last parameter also increases
299 approximately 55% in the digestive gland of the specimens from ST and BG.
300 Additionally, around 50% (NPSH) and 30% (SOD and Vit C) increments were obtained
301 in both tissues of the clams from these two sites. Finally, significantly higher levels of
302 CAT and MTLPs ($p < 0.01$) in the digestive gland, and GPx ($p < 0.05$) in the gills, were
303 detected in the clams from BZ.

304 *3.4. Oxidative damage and post-translational modifications in the clam tissues*

305 Fig. 2A shows the evaluation of the oxidative damage to the lipids and proteins
306 in the gills and in the digestive gland of clams captured at the different sites studied.
307 Malondialdehyde (MDA) content is a consequence of the oxidative damage to lipids.
308 According to their metal contents, MDA levels show top values in both tissues of the
309 ST and BG clams, with significant increases of up to 64% above the levels found at the
310 EL reference site. Meanwhile, lipid peroxidation levels in the BZ and NT specimens
311 were similar to those determined in the EL animals.

312 Global protein damage was determined by measuring both the protein carbonyl
313 (PCO) derivatives, and the advanced oxidation protein products (AOPP) (Fig. 2A). PCO
314 levels showed no differences between the two tissues analyzed, although again, in both
315 tissues, samples from the ST and BG lagoons presented significant increments in
316 protein oxidation ranging from 21 to 35 %, respectively, in comparison to those from
317 EL. Otherwise, AOPP levels increased in both tissues of clams from all the sampling
318 sites in comparison to EL, even though, again ST and BG were the two locations
319 showing the highest levels of protein oxidative damage.

320 Global protein phosphorylation status is shown in Fig. 2B. Our results show that:
321 (i) the digestive gland presented lower protein phosphorylation levels than the gills, and
322 (ii) in both tissues, phosphorylation was significantly reduced in the clams from BZ, ST
323 and BG compared to those from the EL reference site. As also shown in Fig. 2B, the
324 clams from ST and BG presented significant changes in their protein redox status.
325 Protein Cys were significantly more oxidized in the two tissues analyzed from both sites
326 compared to the reference site, an opposite trend to their phosphorylation levels.

327 *3.5. Histological alterations in the clam tissues*

328 The general histopathological state of the digestive gland of the clams collected
329 at different locations along the Tunisian coast is shown in Fig. 3A. The clams from EL
330 (Fig. 3A-EL), the reference site, essentially presented a normal digestive gland
331 architecture, comprising many digestive tubules lined by a single layer of epithelial cells
332 and a narrow lumen. The tubules are separated by intertubular connective tissue. In the
333 clams from the different lagoons, pathological traits were evident. Specifically, the
334 lesions in the BZ (Fig. 3A-BZ) and NT (Fig. 3A-NT) clams were scarce and of
335 relatively low biological significance, consisting mostly of minor foci of haemocyte
336 infiltration and lumen occlusion. However, the clams with the higher levels of metals

337 and molecularly more impacted, *i.e.* ST (Fig. 3A-ST) and BG (Fig. 3A-BG), presented
338 additional and more significant alterations. These included necrotic tubules and foci of
339 intertubular tissue necrosis, with moderate infiltration of haemocytes, as well as
340 epithelial cell vacuolization (most obvious in the BG clams) and atrophic tubules (in the
341 digestive gland of the clams from ST).

342 The gills of the clams from the reference site (EL) mostly presented the normal
343 structure of the frontal, intermediate and abfrontal zones (Fig. 3B). Ctenidia presented a
344 well-defined structure with a regular arrangement of filaments and lamellae. The
345 lamellar epithelial cells in the frontal zone presented intact cilia. The haemolymphatic
346 sinuses presented few haemocytes. Cells containing lipofuscin-like substances, like
347 rhogocytes, were sparse. Similar observations were made from the gills of the
348 specimens collected at the NT, BZ and ST lagoons (Fig. 3B-BZ, 3B-NT, 3B-ST).
349 However, the histological sections of the gills of the clams containing the highest metal
350 levels (BG) revealed moderate abrasion of ctenidia, foci of lamellar deformation and
351 diffuse epithelial desquamation (Fig. 3B-BG).

352 *3.6. Relationships between the abiotic and biotic parameters*

353 Principal Component Analysis (PCA) yielded Principal Component 1 (PC1)
354 accounting for 46.0% of the variability amongst the data while PC2 accounted for
355 12.9% (Fig. 4). Samples from the different tissues, the digestive gland and gills,
356 grouped differently mainly related to PC2. The locations presented the pattern that
357 has been perceived from all the results described. Thus, NT and BZ sites present
358 similar trends and the analysis of their data places them close to the reference site
359 EL. On the other hand, the ST and BG lagoons analyses overlap in both tissues and
360 are grouped differently from the other sites. Thus, the sites studied follow a gradient
361 along the PC1 axis: EL<NT<BZ<ST<BG (Fig. 4).

362 The Spearman correlation coefficients were also determined for cross-
363 validation of the relationships between the different variables (Supplementary Table
364 2). Most physico-chemical parameters, such as SPM, salinity and pH, were either
365 only weakly, or not related with the other variables analyzed. However, a high
366 correlation was found between temperature and the content of metals, including Cd
367 ($p < 0.0001$), Al ($p < 0.0002$), Pb, Cu and Ni ($p < 0.0021$). Significantly, a very
368 strong correlation ($p < 0.0002$) was found between temperature and in-cell H₂O₂, a
369 very important ROS consequence of oxidative stress. Accordingly, a high
370 correlation was found between H₂O₂ and metal concentrations, highlighting Cd ($p <$
371 0.0001), Pb, Cu, Al and Ni ($p < 0.0021$). Strong correlations were found among the
372 oxidative threats (temperature and trace metals), hydrogen peroxide, and the
373 molecular consequences of the oxidative stress generated (antioxidative defenses,
374 damage to biomolecules and PTMs). It should be emphasized that perfect
375 correlations ($p < 0.0001$) were established between H₂O₂ and SOD/MTLP, between
376 MDA and Zn/Fe, and between PCO and Vit C. Additionally, very strong
377 correlations ($p < 0.0002$) were found between temperature and CAT, Cd and
378 MTLP/AOPP/PRS, Pb and MTLP, Cu and PSR, Al and CAT/PCO. Finally, it is
379 noteworthy that significant negative correlations were obtained between protein
380 phosphorylation and several variables.

381 **4. Discussion**

382 Transitional waterbodies are important ecological systems and highly diverse
383 environments in terms of socio-economic drivers, physico-chemical status, geophysics
384 and nature of anthropogenic impacts (Bald et al., 2005; Bilgin and Uluturhan-Suzer,
385 2017; Carreira et al., 2013). The approach used here was designed to assess the status of
386 metal pollution in transitional waters and its biological effects on the organisms at the
387 biochemical and tissue levels. For that purpose, several lagoons along the Tunisian
388 coastline were evaluated, using the bivalve mollusk *V. decussata* as the bioindicator
389 organism.

390 In environmental studies, choosing an appropriate reference site is quite a
391 difficult task, coupled with the difficulty of establishing the physico-chemical elements
392 which correspond to low or undisturbed/non-impacted conditions (Bald et al., 2005).
393 The clams from Ellouze (EL) presented the lowest levels of trace elements (Table 2),
394 which was considered here as an appropriate reference area, in agreement with previous
395 reports (Annabi et al., 2013; Chalghmi et al., 2019; Kessabi et al., 2012). Since EL is in
396 an open-sea coastal area, the suspended particulate matter in the water was significantly
397 lower than in the four lagoons studied. Lagoons are relatively isolated from the open
398 sea, and that makes them more vulnerable to the impacts of pollution (Bilgin and
399 Uluturhan-Suzer, 2017). Many studies have provided information about the presence of
400 high metal contents in the sediments of different Tunisian lagoons. However, those
401 studies focus on local restricted areas, particularly BZ, and the degrees of pollution
402 described vary among reports (Barhoumi et al., 2016; Ben Said et al., 2010; Ghedira et
403 al., 2016; Ghedira et al., 2011; Hellal Mel et al., 2011; Zaaboub et al., 2016; Zakhama-
404 Sraieb et al., 2019). Furthermore, several other pollutants have been detected in the BZ
405 lagoon, including polycyclic aromatic hydrocarbons (PAH), pesticides and

406 polychlorinated biphenyls (Bancon-Montigny et al., 2019; Ben Said et al., 2010; Hellal
407 Mel et al., 2011; Mhadhbi et al., 2019; Triki et al., 2017). Nevertheless, when studied,
408 elevated levels of metals have also been identified in ST (Chalghmi et al., 2016a;
409 Chalghmi et al., 2016b), NT (Oueslati et al., 2018) and BG (Kharroubi et al., 2012),
410 and, additionally, a recent study describes ST as being moderately to highly PAH-
411 contaminated (Chalghmi et al., 2019).

412 Trace metal levels determined here were compared with the soft tissue
413 accumulation levels in the same species found at other contaminated areas in previous
414 studies. Although the concentration of most metals, Pb, Cu, Zn, Fe, Al and Ni,
415 determined here were similar or even lower to those found in other areas, the levels of
416 Cd determined were much higher than those found in all other polluted areas (Cravo et
417 al., 2012; Chalghmi et al., 2016a, 2016b; Bilgin and Uluturhan-Suzer, 2017). Cd is an
418 increasingly common contaminant of great ecological and human concern, due to its
419 widespread industrial use and because it is one of the most hazardous known substances
420 (Ghedira et al., 2011; Serafim and Bebianno, 2007). Significant ecological effects at the
421 population level have been shown in freshwater bivalves (*Pyganodon grandis*)
422 chronically exposed or transferred to a Cd gradient. Increased Cd concentration leads to
423 elevated mortality rates and to decreased population densities, growth, biomass,
424 secondary production, turnover ratio, and reproductive success, which results in the
425 impairment of population health status (Perceval et al., 2004, 2006). In a recent study,
426 lower condition and gonad indices were found in *V. decussata* clams from BG than
427 those from the Ghar El Melh Lagoon (northeast Tunisia), in agreement to the highest
428 levels of Cd and Pb determined in BG clams' tissues (Bejaoui et al., 2020).

429 It must be emphasized that the levels of metals act, in this study, as a mere
430 surrogate of global contamination, in the absence of comprehensive data on organic

431 pollutants in the studied areas. However, it cannot be assumed that aquatic organisms
432 employ the same strategies to accumulate different classes of pollutants, even among
433 trace metals. In aquatic environments, trace metal species are distributed among
434 different water-soluble agents, colloids, suspended matter and sedimentary phases. In
435 general, these pollutants are retained in the sediments, and their mobility and
436 bioavailability can be influenced by many factors including pH, temperature, salinity,
437 redox potential, organic matter, other contaminants or even microbial activity
438 (Filgueiras et al., 2004). Some of the water physico-chemical parameters determined in
439 this study, such as salinity or pH, did not show differences when lagoons were
440 compared with EL. However, the water temperature was higher in the different lagoons,
441 especially at ST and BG (Table 1). BG is a shallow lagoon (4 m average depth) and is
442 therefore highly influenced by the local dry and sunny climate (Khedhri et al., 2017).
443 Among other environmental factors, temperature plays a key role, significantly in
444 polluted coastal and transitional waters (Cabral et al., 2019; Lannig et al., 2006; Nardi et
445 al., 2018; Sokolova and Lannig, 2008). It has been shown that an increase in
446 temperature facilitates the solubility and mobilization of trace metals, which finally
447 results in a higher bioavailability, thus, increasing uptake rates and bioaccumulation in
448 aquatic ectotherms (Mubiana and Blust, 2007; Nardi et al., 2017, 2018; Richards and
449 Chaloupka, 2009; Sokolova and Lannig, 2008). Due to the favored rate of metal uptake
450 and bioaccumulation, an elevated temperature tends to enhance the toxic effects of
451 metals on organisms, resulting in elevated mortality rates (Lannig et al., 2006; Sokolova
452 and Lannig, 2008). Here, we found a high correlation between temperature and the
453 metals contents such as: Cd, Al, Pb, Cu and Ni. Significantly, a perfect correlation ($R =$
454 0.960 , $p < 0.0001$) was found between temperature and Cd (Supplementary Table 2).
455 Thus, specimens from the ST and BG lagoons, where the water temperature is the

456 highest, accumulated the highest levels of metal in their tissues, suggesting that these
457 are the most impacted of all the ecosystems analyzed. Otherwise, the clams from BZ
458 and NT presented moderate levels of trace metals, with significant increases only for Pb
459 and Ni (Table 2).

460 In ectotherms, such as bivalves, the body temperature readily changes with the
461 ambient temperature, resulting in alterations of the rates of most physiological, cellular
462 and biochemical processes (Sokolova and Lannig, 2008). Additionally, when combined
463 with an elevated environmental temperature, pollution has a direct, synergistic impact
464 on these processes (Cabral et al., 2019; Lannig et al., 2006; Nardi et al., 2017, 2018;
465 Sokolova and Lannig, 2008). The gills and the digestive gland play crucial roles in
466 balancing gas and ions and digestion plus nutrient absorption, respectively. Whereas the
467 gills are a major organ of apical entry for waterborne toxicants, such as dissolved
468 metals, the digestive gland absorbs ingested toxicants and it is most likely involved in
469 their detoxification and storage, as an analogue of the vertebrate liver (Louiz et al.,
470 2018). In any case, both organs may have to cope with toxicological challenges and
471 their consequences, such as oxidative stress and cell-level alterations due to metabolic
472 impairment, when the levels of toxicants overcome the organism's defense mechanisms.
473 However, establishing cause-effect relationships between adverse effects and exposure
474 to specific toxicants is challenging. Still, the current histological studies in *V. decussata*
475 show a global pattern that is consistent with increasing contamination at ST and BG,
476 where specimens accumulated the highest concentrations of trace metals in their tissues.
477 Overall, the changes observed range from foci of mild reversible lesions (like foci of
478 haemocytic infiltration) to severe alterations (such as necrosis) that may heavily impair
479 organ functioning if becoming diffuse. Our results are comparable with those observed
480 in clams collected from low-moderately contaminated sites in the Mediterranean basin

481 or adjacent areas, such as Tunisia and Southern Portugal (Costa et al., 2013; Chalghmi
482 et al., 2016b). Altogether, the histopathology of both the gills and the digestive gland
483 suggest that moderate but persistent exposure to mixed environmental toxicants results
484 in chronic adverse effects. These lesions may, in time, reduce the fitness of the
485 population in the area, thus rendering the clams more susceptible to environmental
486 changes and parasites. With this respect, the importance of the digestive gland and gills
487 in detoxification and gas/ion balance must be highlighted, as these processes alter all
488 downstream metabolism.

489 Damage to the respiratory surfaces in the gills results in a less efficient oxygen
490 uptake and an associated greater need for active ventilation, which in turns leads to an
491 increased uptake of metals. Further, oxygen transport can also be impaired by exposure
492 to metals and elevated temperatures, which interferes with the normal O₂ delivery to
493 metabolizing tissues (Sokolova and Lannig, 2008). Besides the limited oxygen supply,
494 it must be added that trace metals strongly affect mitochondrial function by reducing the
495 activity of the electron transport chain and ATP production, which results in an increase
496 in mitochondrial ROS production (Sokolova and Lannig, 2008). A higher H₂O₂ content
497 was found here at BG and ST, followed by BZ (Table 3), in agreement with the
498 temperature levels and metal concentrations, such as Cd, Pb, Cu, Al, Ni, and to a lesser
499 extent Zn and Fe (Supplementary Table 2). In fact, very strong ($R = 0.935$; $p < 0.0002$)
500 and perfect ($R = 0.964$; $p < 0.0001$) correlations were found between H₂O₂ and
501 temperature, and H₂O₂ and Cd, respectively. Cd is a potent disruptor of mitochondrial
502 function, whose sensitivity is strongly enhanced at elevated temperatures, suggesting
503 synergism between both environmental stressors (Cherkasov et al., 2006a; Cherkasov et
504 al., 2006b; Cherkasov et al., 2010).

505 Metal-induced formation of free radicals and the consequent oxidative stress is
506 the basis of the toxicity of many trace metals. Thus, while some metals, such as Fe and
507 Cu, undergo redox-cycling reactions, in a second group, including Cd, Pb and Ni, the
508 mechanism is through the depletion of glutathione and bonding to sulfhydryl groups of
509 proteins (Stohs and Bagchi, 1995; Valko et al., 2005). Antioxidants provide protection
510 against deleterious free radical attacks (López-Barea, 1995; Sies, 1986). This includes
511 primary enzymatic (CAT, GPX and SOD) and non-enzymatic (MTLPs, NPSH and Vit
512 C) antioxidants, that significantly increased in the clams from ST and BG with the
513 higher metal loads in their tissues (Table 3). Thus, high correlations were found
514 between trace metals, the nonradical species H₂O₂, and most antioxidants (Fig. 4 and
515 Supplementary Table 2). Significantly, metallothionein-like proteins (MTLP) showed a
516 very strong correlation ($R = 0.924$ and 0.900 ; $p < 0.0002$) with metals such as Cd and
517 Pb, and a high correlation ($R = 0.888$, 0.796 and 0.827 ; $p < 0.0021$) with Cu, Zn and Fe.
518 MTs play crucial biological roles in the detoxification of trace metals (Ag, Cd, Co, Fe,
519 Hg, Ni, Pb) and in the homeostasis of essential metals (Cu, Zn) (Kägi, 1991).
520 Furthermore, a perfect correlation ($R = 0.954$; $p < 0.0001$) was found between MTLP
521 and H₂O₂, according to the relevant protective role of these proteins against oxidative
522 threats (Kägi, 1991; Viarengo et al., 1999; Viarengo et al., 2000). Protective
523 mechanisms, including antioxidant, detoxifying and damage-repairing proteins,
524 contribute to elevated metabolic rates and maintenance costs in metal-exposed
525 organisms, especially at elevated temperatures (Cherkasov et al., 2006a; Sokolova and
526 Lannig, 2008). The extra-energy expenditure for emergency maintenance, might
527 seriously threaten essential energetically demanding processes (*i.e.*: growth,
528 reproduction, immunity) and, as a consequence, the whole-organism's physiology and
529 survival (Cherkasov et al., 2006a; Lannig et al., 2006; Sokolova and Lannig, 2008).

530 If toxic chemicals exceed pollutant-elicited defense mechanisms, especially in
531 limited energy supply conditions, key biomolecules are damaged (Alhama et al., 2017).
532 Oxidative damage to lipids (MDA) and proteins (PCO and AOPP) was evident here in
533 the specimens with the highest metal concentrations in their tissues, those from BG and
534 ST (Fig. 2A). The perfect correlation between MDA, and Zn and Fe should be
535 highlighted, as well as the very strong correlation between PCO and Al, and between
536 AOPP and Cd (Supplementary Table 2). Proteins are major targets of oxidative stress,
537 which leads to post-translational modifications (PTMs) that can alter their structure and
538 modify their function, and are important in cell signaling (Cabicol and Ros, 2006;
539 Davies, 2005). Reversible changes in the redox state thiols and in the phosphorylation
540 level of proteins showed an opposite trend, both in the digestive gland and gills. Thus,
541 the most oxidatively-threatened clams, those from BG and ST, presented the highest
542 oxidized redox state in their protein cysteines. On the contrary, specimens from those
543 areas, and those from BZ, presented low phosphorylation levels. We had previously
544 described similar patterns in *Mus spretus* testis exposed to the persistent pollutant *p,p'*-
545 DDE (Alhama et al., 2018). Damage and modifications of key biomolecules may reach
546 higher organizational levels when faced with large and long-lasting deleterious effects
547 from shifts in the internal oxidative status, with an emphasis on proteins, which form
548 the primary structure and metabolic machinery of cells. Thus, several histopathological
549 traits were evident in the clams with the highest levels of trace metals which were also
550 more molecularly impacted, corresponding to samples from ST and BG digestive gland,
551 and BG gills.

552 PCA analysis was performed to obtain a global vision of the results. Multivariate
553 analysis grouped differentially the digestive gland and gills. In both tissues, sampling
554 sites along the Tunisian littoral were separated following a pollution gradient:

555 EL<NT<BZ<ST<BG. This gradient is associated with the increasing concentration of
556 most trace metals (Cd, Pb, Cu and Ni) and with temperature, which results in higher
557 levels of oxidative stress biomarkers (H₂O₂, MTLP, AOPP and SOD).

558

559 **5. Concluding remarks**

560 The clam *Venerupis decussatam* has shown its usefulness as biomonitor
561 organism for detecting the impact of pollution in lagoon aquatic ecosystems. Our work
562 demonstrates that multivariate analysis is efficient to separate among sampling sites
563 along the Tunisian littoral, either using the digestive gland or gills tissues. Clams from
564 the different sites are differentially impacted following a pollution gradient:
565 EL<NT<BZ<ST<BG. The battery of biochemical parameters measured correlates with
566 trace metal pollution. The synergistic interaction between temperature and trace metals'
567 bioaccumulation may elicit oxidative stress that, if severe and long-lasting, may
568 overcome the antioxidant capacity of the clams and trigger histopathological traits in
569 their tissues, the digestive gland and gills. Altogether, this study highlights the
570 importance of conducting multifaceted in-field analyses to understand the effects of
571 chemical stressors in complex marine ecosystems, and puts the spotlight on
572 temperature, as an added natural stressor to be considered when organisms face seasonal
573 changes and in a global climate change scenario.

574

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585 **Figure Legends**

586 **Fig. 1.** Map showing the areas studied along the Tunisian coast. The location of the
587 sampling sites and the main polluting activities are indicated.

588 **Fig. 2.** Oxidative damage and post-translational modifications in the clam tissues.
589 **A.** Changes in malondialdehyde (MDA), and advanced oxidation protein products
590 (AOPP) and protein carbonyl (PCO) levels, as biomarkers of oxidative damage to
591 lipids and proteins, respectively. **B.** Electrophoresis-based evaluation of post-
592 translational modification levels, phosphorylation (arbitrary fluorescence units,
593 AUF) and the redox state of cysteine residues (oxidized/reduced thiols ratio), in
594 proteins. Determinations made in the gills (blue, left) and in the digestive gland
595 (red, right) of *V. decussata* are shown. Bars represent the mean \pm SD of at least
596 eight independent determinations (A) or of three independent determinations (B)
597 from twenty pooled clams. Site abbreviations are as follows: Ellouze (EL) and
598 Bizerte (BZ), North (NT) and South Tunis (ST), and Boughrara (BG) lagoons.
599 Statistically significant compared to the EL reference site are expressed as: *, $p <$
600 0.05, **, $p < 0.01$, ***, $p < 0.001$.

601 **Fig. 3.** Histological alterations in the clam tissues. Micrographs of the digestive gland
602 and gill sections of *V. decussata* sampled from the reference site Ellouza (EL), and
603 four Tunisian lagoons suffering different degrees of anthropogenic pollution,
604 Bizerte (BZ), North Tunis (NT), South Tunis (ST) and Boughara (BG), are shown.
605 Six digestive glands and five gills were analyzed per site. **A.** The digestive gland of
606 the clams from Ellouza were characterized by a normal morphoanatomy of the
607 digestive glands, which consisted in digestives tubules (DT) with well-
608 distinguishable lumen (L), formed by a single layer of epithelial cells (EC). The
609 tubules are surrounded by interstitial (connective-like) tissue (IT). The digestive

610 gland of the clams from the four potentially impacted lagoons were mostly
611 characterized by lumen occlusion (LO), haemocyte infiltration within intertubular
612 tissue (HI), atrophic tubules (AT), focal necrosis (NF) and epithelial cell
613 vacuolization (TV). **B.** The clams from the reference site were characterized by a
614 normal morphoanatomy of gills, which consisted in: frontal (F), intermediate (I) and
615 abfrontal (AF) zones; regular filaments arrangements (FA), normal structure of cilia
616 (C), lipofuscin granules (LG) and haemolymphatic sinuses (HS). The most
617 significant alterations detected in the specimens from Boughrara, the potentially
618 most impacted area, were lamellar deformation (LD) with epithelial desquamation
619 (ED) and abrasion of ctenidia (AC).

620 **Fig. 4.** Principal component analysis (PCA). Plot of physico-chemical parameters in
621 the water, and metal contents and biochemical determinations (green) in the gills
622 (blue) and in the digestive gland (red) of *V. decussata* clams, PC1 vs PC2, are
623 represented. Ellipses show 95% confidence intervals. Sites are represented as
624 follows: the Ellouza (●), and Bizerte (◆), North (■) and South (✱) Tunis, and
625 Boughrara (✕) lagoons. Biomarker abbreviations are as follows: AOPP, advanced
626 oxidation protein products; GPx, glutathione peroxidase; CAT, catalase; MDA,
627 Malondialdehyde; MTLP, metallothionein-like proteins; NPSH, non-protein
628 sulfhydryls; PCO, protein carbonyl levels; PP, protein phosphorylation; PRS,
629 protein redox state; SOD, superoxide dismutase; SPM, suspended particulate
630 matter; Vit C, vitamin C.

631

632

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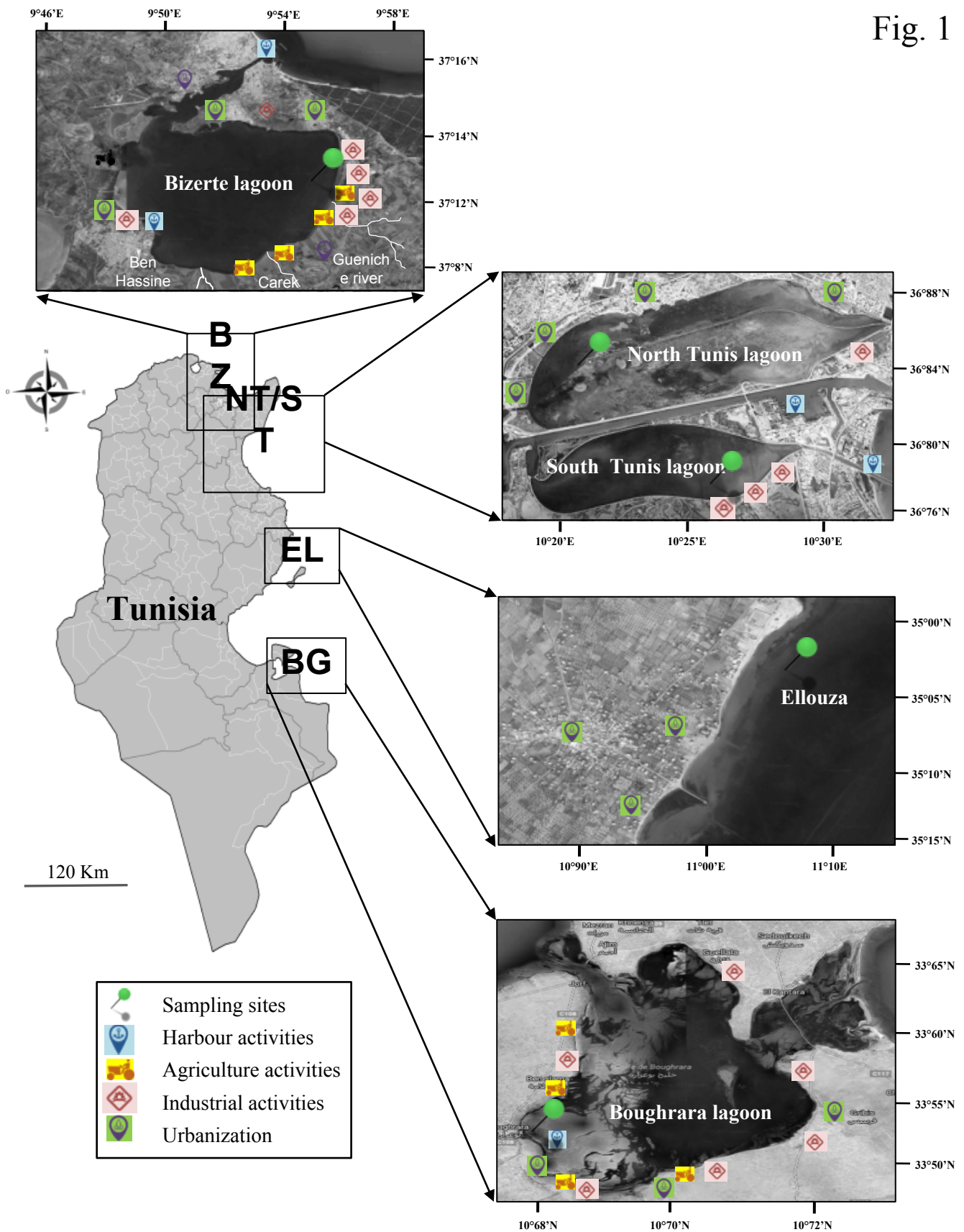
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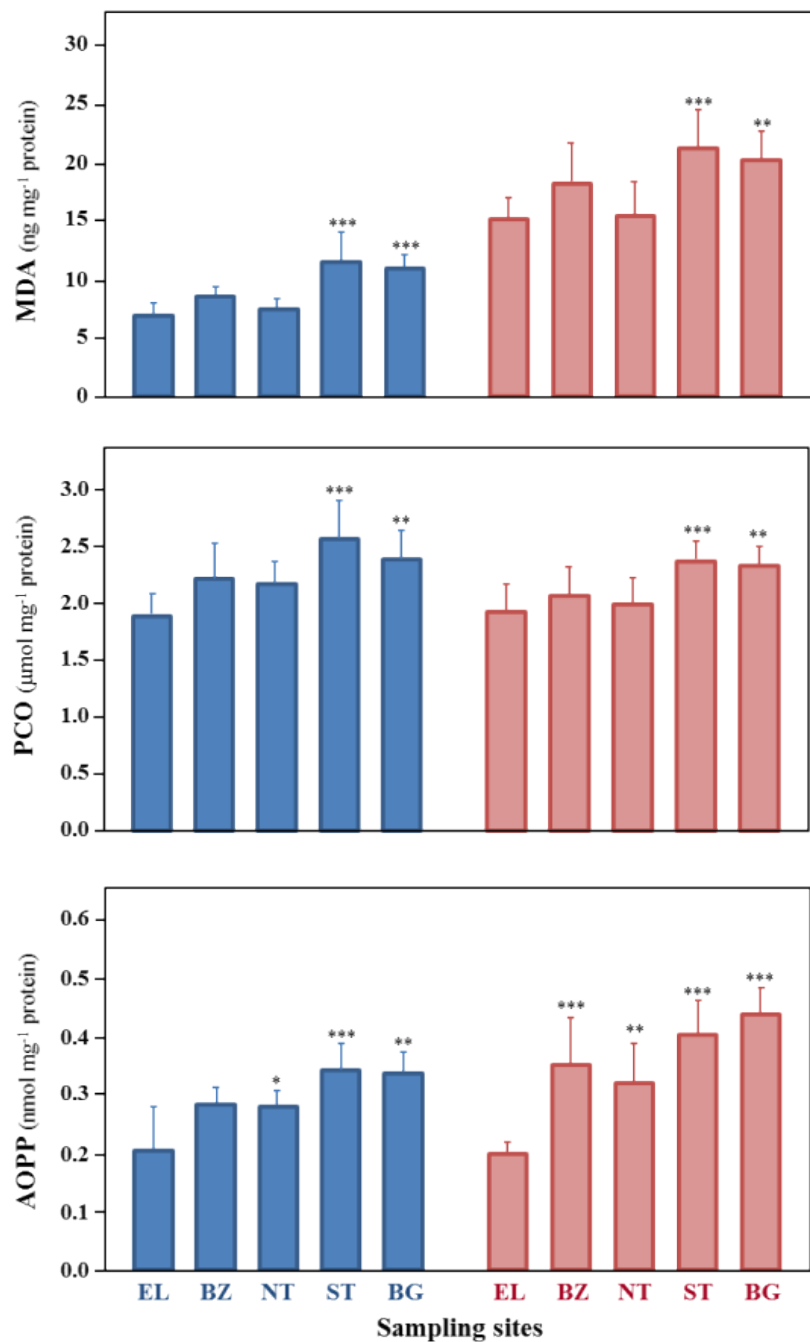
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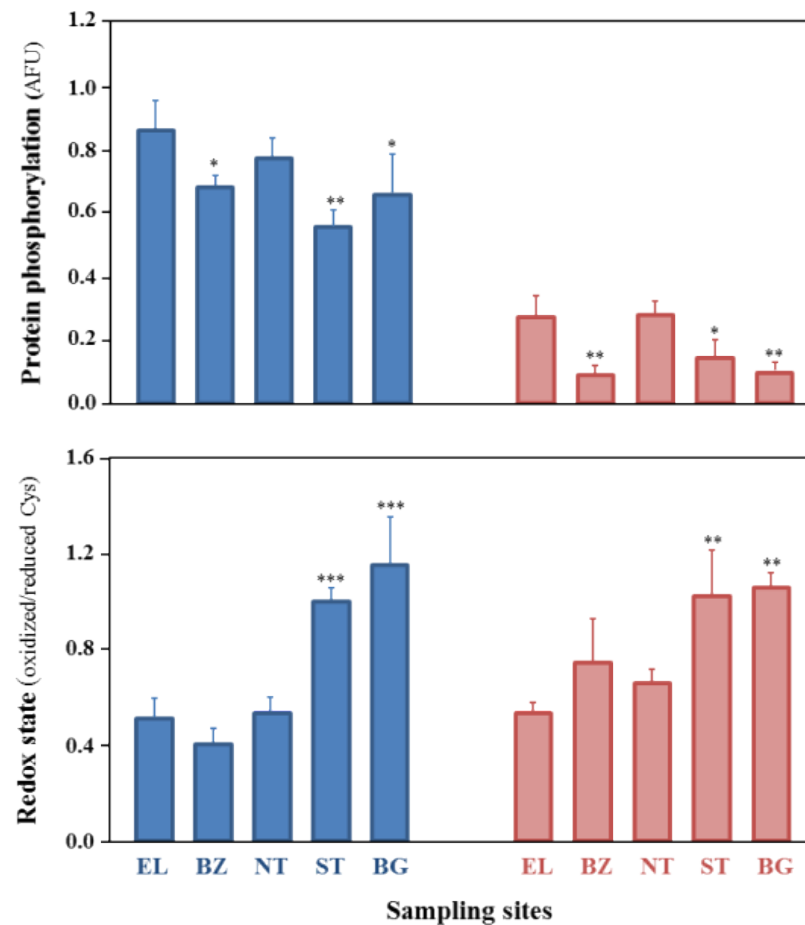
Fig. 1



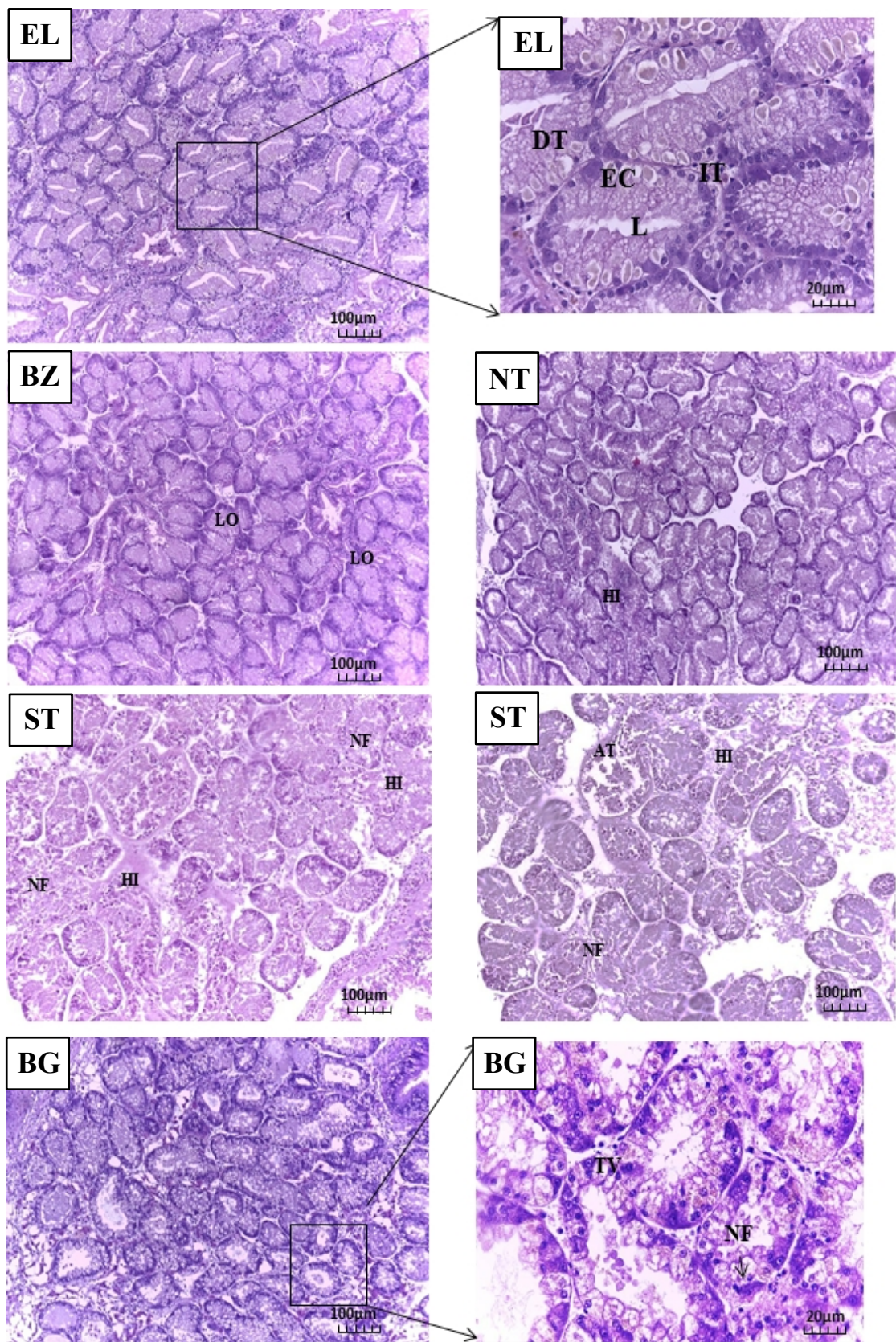
(A)



(B)



(A)



(B)

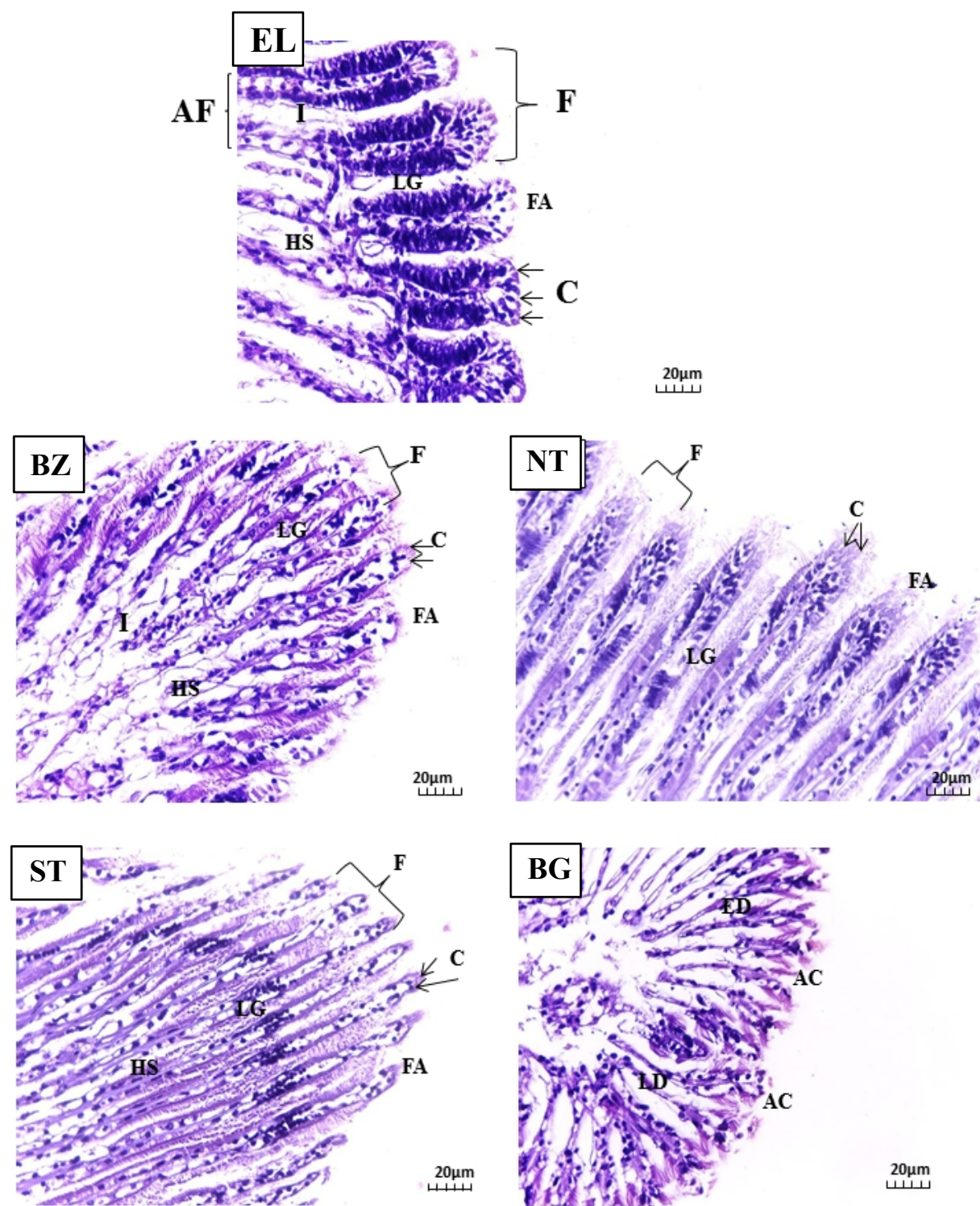


Fig. 4

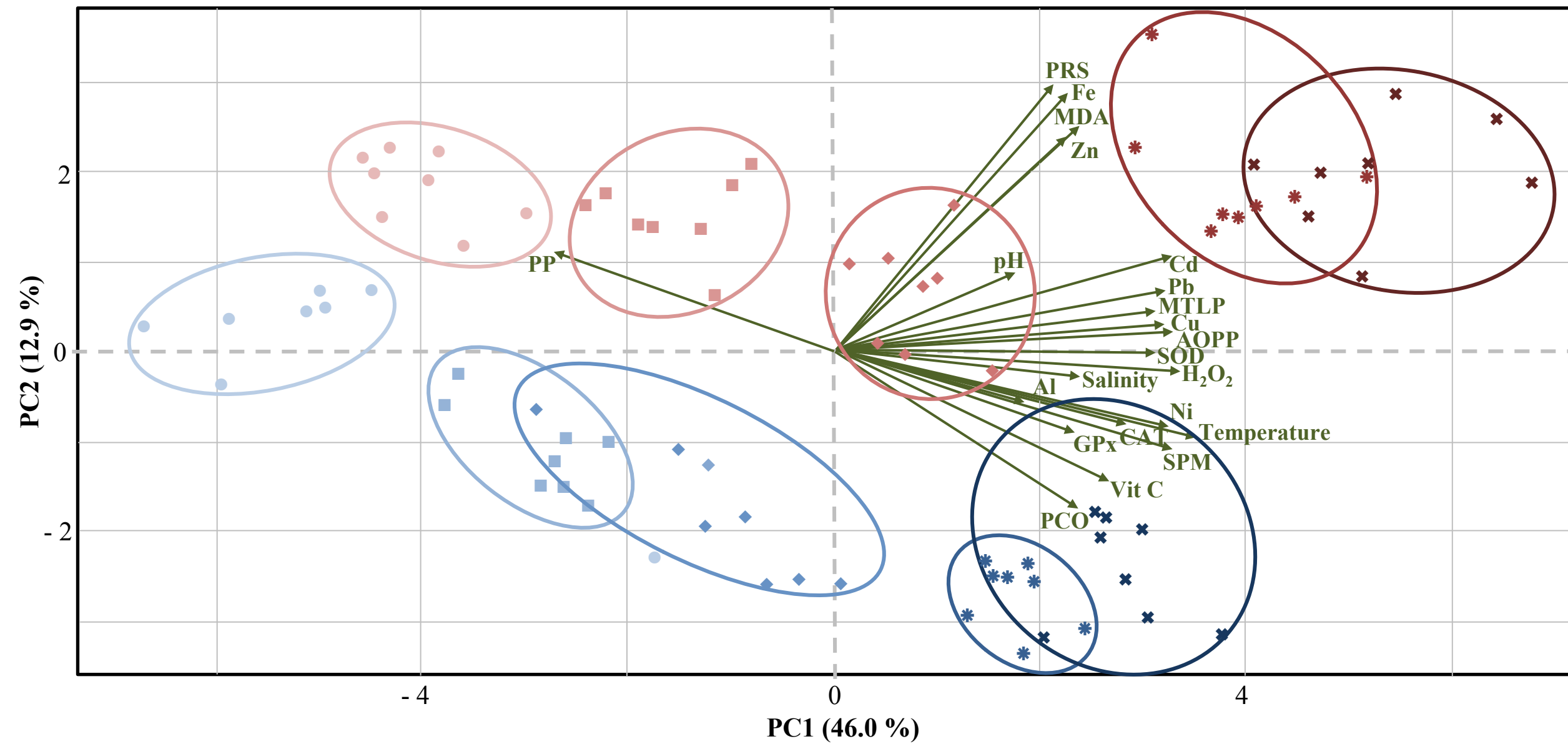


Table 1. Physico-chemical parameters determined in water at the different studied locations along the Tunisian coast.

Sites ^a	Temperature (°C)	Salinity (psu)	pH	Suspended Particulate Matter (mg L ⁻¹)
EL	23.36 ± 2.82 ^b	37.91 ± 2.10	8.15 ± 0.02	112.51 ± 6.78
BZ	28.06 ± 2.12	37.33 ± 1.63	8.00 ± 0.03	200.45 ± 14.36***
NT	26.41 ± 1.53	37.22 ± 2.62	8.01 ± 0.01	163.22 ± 10.32***
ST	29.17 ± 2.05*^c	38.01 ± 1.11	8.10 ± 0.02	198.66 ± 11.75***
BG	31.47 ± 1.99**	39.11 ± 1.39	8.00 ± 0.11	231.22 ± 9.47***

^a Sampling sites along the Tunisian coast are as follows: Ellouze (EL) and Bizerte (BZ), North (NT) and South Tunis (ST), and Boughrara (BG) lagoons.

^b Results represent the mean ± SD from the analysis of three independent samples. Determinations were recorded during February 2017.

^c Statistical significance was compared with the values at EL, the reference site: *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$. Statistically significant results are highlighted in bold.

Table 2. Trace metals concentrations in the gills and the digestive gland of *Venerupis decussata* clams from different locations along the Tunisian coast.

Tissues	Sites ^a	Metals ($\mu\text{g g}^{-1}$ dry weight)						
		Cd	Pb	Cu	Zn	Fe	Al	Ni
Gills	EL	0.85 \pm 0.09 ^b	0.99 \pm 0.17	4.10 \pm 1.11	25.73 \pm 5.35	367.39 \pm 7.41	19.84 \pm 0.50	0.32 \pm 0.05
	BZ	0.92 \pm 0.12	1.29 \pm 0.10*	4.89 \pm 0.83	30.51 \pm 2.52	369.35 \pm 6.22	21.47 \pm 2.88	0.44 \pm 0.03*
	NT	0.90 \pm 0.18	1.38 \pm 0.27	4.69 \pm 1.04	30.83 \pm 3.95	345.27 \pm 6.94	21.32 \pm 2.76	0.59 \pm 0.03***
	ST	1.26 \pm 0.18***^c	1.68 \pm 0.17***	6.87 \pm 1.02***	32.02 \pm 6.88	373.74 \pm 10.97	23.83 \pm 1.43***	0.77 \pm 0.10***
	BG	1.42 \pm 0.25***	1.91 \pm 0.21***	8.77 \pm 1.13***	34.19 \pm 6.95	377.37 \pm 7.32	22.49 \pm 0.51*	1.06 \pm 0.10***
Digestive gland	EL	0.87 \pm 0.10	1.13 \pm 0.33	5.39 \pm 0.97	36.08 \pm 7.49	395.92 \pm 15.03	16.90 \pm 4.25	0.42 \pm 0.07
	BZ	1.03 \pm 0.10	1.66 \pm 0.46*	5.54 \pm 0.95	36.52 \pm 7.02	406.36 \pm 12.66	20.96 \pm 4.16	0.54 \pm 0.09
	NT	1.01 \pm 0.07	1.82 \pm 0.31**	6.31 \pm 0.65	37.17 \pm 7.33	402.92 \pm 8.97	20.25 \pm 3.98	0.58 \pm 0.12
	ST	1.32 \pm 0.10**	2.33 \pm 0.31***	8.42 \pm 1.06***	50.44 \pm 6.36**	433.83 \pm 13.51***	21.87 \pm 4.25	0.82 \pm 0.18***
	BG	1.87 \pm 0.38***	2.31 \pm 0.32***	8.13 \pm 1.17***	50.08 \pm 6.23**	435.22 \pm 13.07***	22.99 \pm 4.32	0.84 \pm 0.09***

^a Sampling site along the Tunisian coast are as follows: Ellouze (EL) and Bizerte (BZ), North (NT) and South Tunis (ST), and Boughrara (BG) lagoons.

^b Results represent the mean \pm SD of six independent replicates of twenty pooled clams.

^c Statistical significance was compared with the values at EL, the reference site: *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$. Statistically significant results are highlighted in bold.

Table 3. Hydrogen peroxide levels and antioxidant responses in the gills and in the digestive gland of *V. decussata* clams from different locations along the Tunisian coast.

Tissues	Sites ^a	H ₂ O ₂ ($\mu\text{mol mg}^{-1}$)	Enzymatic antioxidants			Non-enzymatic antioxidants		
			CAT ^b (U mg^{-1})	GPx (U mg^{-1})	SOD (U mg^{-1})	MTLPs ($\mu\text{mol g}^{-1}$)	NPSH ($\mu\text{mol mg}^{-1}$)	Vit C (nmol mg^{-1})
Gills	EL	1.08 \pm 0.37 ^c	10.68 \pm 1.82	33.57 \pm 5.71	14.08 \pm 4.52	0.23 \pm 0.08	0.23 \pm 0.06	11.27 \pm 2.95
	BZ	1.54 \pm 0.18 ^{**d}	13.13 \pm 2.21	38.58 \pm 1.72 [*]	16.43 \pm 2.10	0.34 \pm 0.10	0.30 \pm 0.06	13.42 \pm 1.54
	NT	1.15 \pm 0.15	10.95 \pm 2.83	37.77 \pm 1.90	14.07 \pm 1.27	0.28 \pm 0.026	0.26 \pm 0.07	12.37 \pm 1.96
	ST	1.73 \pm 0.18 ^{***}	18.05 \pm 3.58 ^{**}	44.75 \pm 2.14 ^{***}	19.21 \pm 2.47 ^{**}	0.40 \pm 0.07 ^{**}	0.35 \pm 0.08 [*]	15.69 \pm 1.77 ^{***}
	BG	1.74 \pm 0.10 ^{***}	17.74 \pm 3.20 ^{**}	43.26 \pm 2.26 ^{***}	18.82 \pm 0.88 ^{**}	0.44 \pm 0.09 ^{**}	0.32 \pm 0.07	14.44 \pm 1.72 [*]
Digestive gland	EL	1.30 \pm 0.08	10.30 \pm 1.65	28.35 \pm 2.76	15.74 \pm 3.34	0.31 \pm 0.05	0.15 \pm 0.02	11.39 \pm 2.06
	BZ	1.69 \pm 0.14 ^{***}	17.03 \pm 4.18 ^{**}	28.17 \pm 3.53	18.60 \pm 1.81	0.44 \pm 0.07 ^{**}	0.17 \pm 0.01	13.05 \pm 0.98
	NT	1.46 \pm 0.17	11.15 \pm 2.85	27.27 \pm 4.95	15.80 \pm 2.88	0.36 \pm 0.06	0.15 \pm 0.01	11.39 \pm 1.51
	ST	1.84 \pm 0.27 ^{***}	17.18 \pm 1.83 ^{**}	41.46 \pm 8.99 ^{***}	21.22 \pm 1.07 ^{***}	0.48 \pm 0.08 ^{***}	0.24 \pm 0.03 ^{***}	14.88 \pm 1.26 ^{***}
	BG	1.92 \pm 0.08 ^{***}	17.73 \pm 2.53 ^{***}	52.07 \pm 4.95 ^{***}	20.11 \pm 1.66 ^{**}	0.47 \pm 0.06 ^{***}	0.21 \pm 0.01 [*]	14.27 \pm 1.10 ^{**}

^a Sampling site along the Tunisian coast are as follows: Ellouze (EL) and Bizerte (BZ), North (NT) and South Tunis (ST), and Boughrara (BG) lagoons.

^b Abbreviations of the different antioxidants are as follows: Catalase (CAT), glutathione peroxidase (GPx), superoxide dismutase (SOD), metallothionein-like proteins (MTLPs), non-protein sulfhydryls (NPSH), and ascorbic acid (Vitamin C, Vit C).

^c Results are given as the mean \pm SD of six (H₂O₂) or eight (antioxidant responses) replicates of twenty pooled clams.

^d Statistical differences between sites, compared to EL, are represented as: *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$. Statistically significant results are highlighted in bold.

Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

Author statement:

Safa Bejaoui: Conceptualization, Investigation, Resources, Writing, Funding acquisition;
Carmen Michán: Conceptualization, Formal analysis, Resources, Writing, Visualization, Supervision; **Khaoula Telahigue:** Investigation; **Salwa Nechi:** Investigation; **Mhamed el Cafsi:** Supervision, Funding acquisition; **Nejla Soudani:** Investigation; **Julián Blasco:** Writing, Funding acquisition; **Pedro M. Costa:** Writing, supervision, Funding acquisition; **José Alhama:** Conceptualization, Formal analysis, Resources, Writing, Visualization, Supervision, Funding acquisition

Supplementary Table 1. Biometric parameters determined in *Venerupis decussata* clams collected at different locations along the Tunisian littoral zone.

Sites ^a	Shell length (mm)	Shell width (mm)	Weight ^b (g)
EL	39.5 ± 2.5	16.3 ± 2.0	13.3 ± 2.9
BZ	40.2 ± 1.5	17.9 ± 0.5	13.6 ± 1.7
NT	42.2 ± 1.0	16.6 ± 1.2	12.9 ± 2.4
ST	41.2 ± 2.0	18.3 ± 0.4	12.7 ± 0.5
BG	43.3 ± 1.3	19.3 ± 1.1	14.2 ± 1.8

^a Sampling sites along the Tunisian coast are as follows: Ellouze (EL) and Bizerte (BZ), North (NT) and South Tunis (ST), and Boughrara (BG) lagoons.

^b Weight values of the whole clams before dissection.

Forty clams were analyzed for each location. Statistical differences among clams from the different locations and those at EL reference site were not significant.

Supplementary Table 2. Spearman correlation coefficients between the physico-chemical parameters determined in water, and metal concentrations and biochemical determinations measured in clam tissues (digestive gland and gills) along the Tunisian littoral zone.

	Temperature	Salinity	pH	SPM	Cd	Pb	Cu	Zn	Fe	Al	Ni	H ₂ O ₂	CAT	GPx	SOD	MTLP	NPSH	Vit C	MDA	PCO	AOPP	PP	PRS		
Temperature	1.000																								
Salinity	0.700 *	1.000																							
pH	-0.616	0.051	1.000																						
SPM	0.900 **	0.500	-0.872 **	1.000																					
Cd	0.960 ****	0.665 *	-0.568	0.837 **	1.000																				
Pb	0.812 **	0.492	-0.391	0.615	0.915 ***	1.000																			
Cu	0.837 **	0.689 *	-0.328	0.640	0.915 ***	0.903 ***	1.000																		
Zn	0.419	0.271	-0.164	0.295	0.636	0.782 *	0.697 *	1.000																	
Fe	0.468	0.394	-0.253	0.419	0.661 *	0.697 *	0.673 *	0.939 ***	1.000																
Al	0.911 ***	0.615	-0.442	0.739 *	0.818 **	0.673 *	0.661 *	0.188	0.200	1.000															
Ni	0.886 **	0.591	-0.455	0.689 *	0.903 ***	0.891 **	0.855 **	0.503	0.406	0.818 **	1.000														
H ₂ O ₂	0.935 ***	0.714 *	-0.492	0.812 **	0.964 ****	0.867 **	0.891 **	0.661 *	0.721 *	0.794 **	0.806 **	1.000													
CAT	0.935 ***	0.640	-0.480	0.788 **	0.891 **	0.733 *	0.794 **	0.309	0.358	0.927 ***	0.806 **	0.855 **	1.000												
GPx	0.788 *	0.788 *	-0.227	0.591	0.661 *	0.491	0.527	0.055	0.103	0.891 **	0.697 *	0.684 *	0.733 *	1.000											
SOD	0.862 **	0.714 *	-0.316	0.689 *	0.891 **	0.818 **	0.855 **	0.636	0.709 *	0.758 **	0.697 *	0.964 ****	0.842 **	0.648 *	1.000										
MTLP	0.840 **	0.580	-0.456	0.728 *	0.924 ***	0.900 ***	0.888 **	0.796 **	0.827 **	0.638	0.748 *	0.954 ****	0.766 *	0.468	0.936 ***	1.000									
NPSH	0.506	0.346	-0.177	0.383	0.279	0.134	0.209	-0.460	-0.474	0.711 *	0.426	0.255	0.596	0.687 *	0.267	0.079	1.000								
Vit C	0.877 **	0.617	-0.329	0.679 *	0.802 **	0.693 *	0.748 *	0.310	0.304	0.918 ***	0.766 *	0.839 **	0.912 ***	0.784 **	0.851 **	0.744 *	0.671 *	1.000							
MDA	0.443	0.295	-0.177	0.345	0.636	0.721 *	0.661 *	0.964 ****	0.964 ****	0.236	0.418	0.709 *	0.370	0.091	0.721 *	0.833 **	-0.413	0.389	1.000						
PCO	0.886 **	0.591	-0.379	0.689 *	0.794 **	0.685 *	0.733 *	0.212	0.176	0.952 ***	0.830 **	0.782 *	0.915 ***	0.818 **	0.758 *	0.657 *	0.754 *	0.973 ****	0.248	1.000					
AOPP	0.837 **	0.492	-0.505	0.739 *	0.903 ***	0.855 **	0.745 *	0.685 *	0.733 *	0.721 *	0.721 *	0.903 ***	0.806 **	0.515	0.891 **	0.930 ***	0.140	0.723 *	0.745 *	0.648 *	1.000				
PP	-0.369	-0.246	0.253	-0.369	-0.552	-0.539	-0.539	-0.867 **	-0.927 ***	-0.139	-0.285	-0.624	-0.309	0.006	-0.612	-0.748 *	0.486	-0.292	-0.939 ***	-0.139	-0.673 *	1.000			
PRS	0.812 **	0.739 *	-0.303	0.615	0.903 ***	0.855 **	0.939 ***	0.673 *	0.661 *	0.636	0.867 **	0.842 **	0.770 *	0.539	0.782 *	0.845 **	0.146	0.669 *	0.624	0.661 *	0.758 *	-0.567	1.000		

Number in bold indicate significant correlations (red, positive; green, negative). Asterisks indicate the signification of the correlations: *, $p < 0.0332$; **, $p < 0.0021$; ***, $p < 0.0002$; ****, $p < 0.0001$.

Abbreviations are as follows: Suspended particulate matter (SPM), catalase (CAT), glutathione peroxidase (GPx), superoxide dismutase (SOD), metallothionein-like proteins (MTLP), non-protein sulfhydryls (NPSH), vitamin C (Vit C), malondialdehyde (MDA), protein carbonyl levels (PCO), advanced oxidation protein products (AOPP), protein phosphorylation (PP) and protein redox state (PRS).