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1 Biotransformation, antioxidant and histopathological biomarker responses to contaminants in
2 European and American Yellow Eels from the Gironde and St. Lawrence Estuaries

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14

15 **Abstract**

16 Since the early 1980s, populations of American (*Anguilla rostrata*) and European eels (*Anguilla*
17 *anguilla*) have suffered a sharp decline. The causes of their decline are likely multifactorial and
18 include chemical pollution. A field study was conducted in eight sites varying in organic and
19 metal contamination along the St. Lawrence (Eastern Canada) and Gironde (France) systems to
20 investigate the relationships among contaminants, biological characteristics and
21 biotransformation, antioxidant and histopathological biomarkers in eels from both species. For *A.*
22 *rostrata*, no major influences of persistent organic contaminants on biomarkers were identified.
23 For *A. anguilla*, eels from the most contaminated site expressed higher surface of
24 MelanoMacrophage Centers (MMCs) and eels from another contaminated site expressed higher
25 amount of lipofuscin pigment. These two histopathological biomarkers were also associated with
26 aging. Compared to eels from the cleanest French site, higher hepatic catalase activity and
27 density of MMC in eels from contaminated sites was related to higher concentration of organic
28 (DDT and metabolites, sum of PCBs, sum of PBDEs) and inorganic (Hg and Cd) contaminants.
29 In both species, a higher deposition of spleen hemosiderin pigment was measured in eels from
30 the most brackish sites compared to eels living in freshwater environments. Our results suggest
31 an association between higher hemosiderin pigment and metal contamination (As for *A. anguilla*
32 and Pb for *A. rostrata*). Parasitism by *A. crassus* was observed in European eels from freshwater
33 sites but not in eels from brackish habitats. Overall, contamination may pose a greater risk for
34 the health of European compared to American eels.

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39 **Keywords:** EROD activity, Antioxidant enzymes, Histopathological markers, Pollution, Atlantic
40 Eels

41 1. Introduction

42 In the Northern hemisphere, the abundance of European eel (*Anguilla anguilla*) and
43 American eel (*Anguilla rostrata*), here called “Atlantic eels”, has severely declined since the
44 early 1980’s (FAO/ICES, 2009; Cosewic, 2012). The causes of their decline are likely
45 multifactorial, including variables such as oceanographic and climate changes, overfishing,
46 barriers to migration, habitat degradation, parasites and chemical pollution (Castonguay et al.,
47 1994; Geeraerts and Belpaire, 2010). Although management and restoration programs have been
48 conducted in several hydrosystems in Europe and North America, eel stocks have not recovered
49 to levels of the past (ICES 2016). An increasing number of recent reports support that pollutants
50 might be among the possible synergistic causes contributing to the collapse of eel stocks
51 (Belpaire et al. 2016). Due to their unique ecological and physiological traits, eels are
52 particularly prone to bioaccumulation of contaminants through gills, skin and contaminated food
53 during their feeding and growth stage (yellow eel). Eels are facultative catadromous fish that can
54 be in contact with contaminated sediments for extended periods, leading to the accumulation of
55 high concentrations of organic contaminants due to their high lipid content (Robinet and
56 Feunteun, 2002; Daverat and Tomás, 2006; Thibault et al., 2007). Finally, eels are considered
57 good bio-indicators of contaminated habitats and are expected to reflect the impacts of local
58 pollutants (Belpaire and Goemans, 2007).

59 Recently, decreasing muscle concentrations of organic contaminants such as
60 PolyChlorinated Byphenyls (PCBs) and OrganoChlorine Pesticides (OCPs) have been reported
61 in eels from some historically contaminated habitats since their ban 40 years ago (Belpaire et al.,
62 2016). From 1988 to 2008, a decreasing trend in PCB and OCP concentrations has been reported
63 in the muscle of *A. rostrata* from the historically contaminated Lake Ontario (Byer et al. 2013).
64 This trend was confirmed by a significant decrease in the embryotoxic potential of organic
65 mixtures extracted from Lake Ontario eels captured in 1988, 1998 and 2008 for developing
66 *Fundulus heteroclitus* (Rigaud et al. 2016). However, these contaminants are persistent in
67 sediments and levels of PCBs, OCPs and PolyBrominated Diphenyl-Ethers (PBDEs) still remain
68 high and preoccupying in yellow eels, for example in eels from the Gironde system (Tapie et al.,
69 2011; Guhl et al., 2014). In addition to organic contaminants, metals such as mercury (Hg) and
70 cadmium (Cd) show no trend of decrease and persist in contaminated areas (Maes et al., 2008).

71 Consequently, eels inhabiting contaminated areas could be still affected by a combination of
72 contaminants acting synergistically.

73 Several field and laboratory studies have reported effects of contaminants in eels at several
74 levels of biological organization from the cellular, molecular and tissue levels to the level of
75 whole organisms using various biological biomarkers to investigate their health status. They
76 have shown that contaminants can delay growth, reduce lipid storage efficiency, cause oxidative
77 stress, induce DNA damage and histopathological lesions, which can ultimately negatively affect
78 migratory and reproductive capacities (Couillard et al., 1997; Robinet and Feunteun, 2002;
79 Palstra et al., 2006; Pierron et al., 2007; Geeraerts and Belpaire, 2010, Gravato et al. 2006).
80 Others effects such as disturbances of the immune and endocrine systems and a reduction of
81 migratory and reproductive capacities have been also reported (Belpaire et al. 2007). In addition,
82 ecological gradients such as latitudinal cline, longer distances to the spawning site and
83 differences in salinity directly or indirectly affect growth rate, body condition, size at
84 metamorphosis and lipid storage, which in turn could affect contaminant uptake (Vøllestad,
85 1992; Edeline et al., 2007; Thibault et al., 2007; Belpaire et al., 2009; Jessop, 2010; Daverat et
86 al., 2012). For *A. anguilla*, studies have suggested that the nematode *Anguillicola crassus* in the
87 swimbladder of yellow eels could affect their swimming performance, reduce their resistance to
88 other stressors such as contamination and increase spleen size, macrophage and lymphocyte
89 production to combat parasites (Kirk et al., 2000; Lefebvre et al., 2004).

90 In a series of related studies performed by our team along the St. Lawrence and Gironde
91 hydrosystems (Eastern Canada and Southwest France, respectively), we have identified that
92 certain metals and organic contaminants could disturb physiological functions in yellow Atlantic
93 eels (Baillon et al., 2015a, 2015b, 2016; Caron et al., 2016; Pannetier et al., 2016). In Baillon et
94 al. (2015a, 2015b, 2016), we conducted a large scale and without *a priori* transcriptomic based
95 approach. The global hepatic transcriptome of animals was determined by RNA-Seq. Genes that
96 most likely responded to single factors were identified. We detected transcriptomic responses
97 typical of PCB-170 and Cd exposure in wild *A. anguilla* collected in the most contaminated site
98 in the Gironde Estuary, associated with changes in the transcription levels of genes involved in
99 hepatic energy metabolism. In another paper (Caron et al. 2016), we focused on biomarkers
100 involved in glycolytic, aerobic and anaerobic capacities and lipid metabolism. We reported an
101 association between silver (Ag), lead (Pb) and arsenic (As) contamination in wild *A. rostrata*
102 from the St. Lawrence River system and inhibition of the enzyme G6PDH, involved in lipid

103 metabolism in liver and which also plays a role in oxidative stress response. This study has also
104 shown an association between zinc (Zn), copper (Cu) and Pb contamination and altered
105 glycolytic and anaerobic capacities in wild *A. anguilla* from the Gironde system. More recently,
106 we have demonstrated a significant difference in early size-at-age (a proxy of early growth rate)
107 between yellow eels captured upstream compared to downstream sites of the St. Lawrence and
108 the Gironde systems (Patey et al., submitted). These differences were present as early as 1 year
109 old and could influence contaminant accumulation.

110 In the present study, we investigated the relationships among tissue concentrations of
111 inorganic and organic contaminants, biological characteristics such as size, age, condition factor
112 and muscle lipids and a set of complementary biomarkers such as hepatic biotransformation
113 enzymes, antioxidant enzymes and splenic histopathological measurements in Atlantic eels. The
114 selected biomarkers have been successfully used, alone or in combination, in previous
115 biomonitoring studies on eels (Couillard et Hodson, 1996; Pacheco et Santos, 2002; Van der
116 Oost et al., 2003; Buet et al., 2006) and are among the list of recommended biomarkers in the
117 Report of the Workshop of the Working Group on Eels and the Working Group on Biological
118 Effects of Contaminants (ICES 2016). First, we examined the induction of Cytochrome P450-
119 dependent monooxygenase, measured as ethoxyresorufin *O*-deethylase (EROD) in eel livers.
120 This biotransformation marker is involved in a well-documented biochemical reaction of fish to
121 organic planar contaminants such as PCBs, polycyclic aromatic hydrocarbons (PAHs), dioxins
122 and furans and may be related to other effects such as oxidative stress (Van der Oost et al., 2003;
123 Whyte et al., 2008). Secondly, we measured hepatic activities of two antioxidant enzymes,
124 catalase (CAT) and superoxide dismutase (SOD), commonly used as biomarkers of antioxidant
125 capacities in fish and induced by several contaminants such as PCBs, PAHs, organochlorine
126 pesticides (OCPs) and metals (Van der Oost et al., 1996; Livingstone, 2001). In addition, we
127 examined a suite of histopathological biomarkers induced by chronic exposure to contaminants,
128 including density and surface of melano-macrophage centers (MMCs) and deposition of
129 lipofuscin and hemosiderin pigments in spleen (Wolke, 1992; Couillard et al., 1997; Fournie et
130 al., 2001; Au, 2004). In general, biomolecular biomarkers such as EROD, CAT and SOD are
131 more sensitive and specific short-term responses (days to weeks), whereas histopathological
132 biomarkers are responses to chronic exposure to contaminants and reveal more persistent
133 damage.

134 In this study, yellow eels were sampled in four sites varying in anthropogenic pressure and
135 contamination level in each of the Saint Lawrence System (SLS, Quebec, Canada) and the
136 Gironde System (GS, France), for two consecutive years. The objectives of this study were: (1)
137 to examine variations in responses of biomarkers in wild American and European eels among
138 sampling sites varying in contamination levels and between sampling years within a species; (2)
139 to investigate the relationships between biological characteristics, contaminants and biomarker
140 responses; and (3) finally, to generate a database to better target and prioritize future studies on
141 the impacts of contaminants on eel health in the Gironde and St. Lawrence River basins.

142

143 2. Materials and methods

144

145 2.1 Sampling sites

146 A total of 240 yellow eels (15 yellow eels/site/year/species) were captured in eight
147 different sites selected according to salinity, metal and organic contamination gradients found in
148 sediments and biota along the SLS and the GS in May and June 2011 and 2012 as previously
149 described by Baillon et al. (2015b) and Laporte et al. (2016). For *A. rostrata*, each year, 60
150 yellow eels were sampled in two reference sites, St Jean River (SJR, freshwater tributary of the
151 Gulf of the SLS) and Sud-Ouest River (SOR, freshwater tributary of the estuary of the SLS) and
152 two fluvial contaminated sites, St. Pierre Lake (SPL) and St. François Lake (SFL). Yellow eels
153 from SOR and SJR were captured in running freshwater habitats, which are tributaries opening
154 into a brackish estuary suggesting an influence of marine water (Thibault et al., 2007).

155 For *A. anguilla*, each year, 60 yellow eels were sampled in two fluvial sites, the
156 Dordogne river (DOR, considered as the least contaminated site in the watershed of the Gironde)
157 and the Garonne river (GAR, located near the city of Bordeaux and contaminated), one estuarine
158 site, Gironde (GIR, contaminated by metals and organic contaminants, (Kessaci et al., 2014))
159 and finally, one salt marsh site in the Arcachon Bay, Certes (CER) considered as a reference site.

160

161 2.2 Wild eels sample collection

162 All yellow eels were processed using the same standardized protocols in France and in
163 Québec. European and American yellow eels were captured using fyke nets (mesh size of 6.4
164 mm), trawl (mesh size of 31 mm) and electro-fishing by professional fishers or technical staff.
165 Given the substantial variability in the size of eels growing in different habitats, it was not

166 possible to collect same-sized eels in each sampling site. Body Length (BL) (± 1 mm) and Body
167 Mass (BM) (± 0.1 g) were measured for each fish in order to estimate the condition factor K
168 ($K=[(BM \text{ (g)}/(BL \text{ (cm)})^3]$). We also determined the ocular index (OI) expressed in % which
169 reflects changes in eye diameter, a useful criterion for assessing the degree of sexual
170 maturation/silvering in eels. Ocular enlargement occurs early in the sequence of sexual
171 maturation (Acou et al., 2005; Durif et al., 2005) and is thought to be a requirement for migration
172 at depth in the ocean. Then, eels were killed as soon as possible by decapitation and rapidly
173 dissected to collect parts of liver, caudal muscle and spleen for biochemical and histological
174 analysis. Samples of muscle were dissected in a standardized area situated at 2 cm posterior to
175 the anus and above the lateral line and kept in aluminium foil at -20°C . Samples of liver were
176 stored in cryogenic tubes at -80°C and samples of spleen were fixed in bottles containing 10%
177 phosphate-buffered formalin. Age was determined from transverse sections of sagittal otoliths as
178 described previously by Verreault et al. (2009). Muscle lipid content was determined by
179 gravimetry after filtration and evaporation of an aliquot of the DiChloroMethane (DCM) extract
180 from the same standardized area of muscle described above and was expressed in percentage of
181 dry-basis lipid content. The moisture content was determined as described by Tapie et al., (2011)
182 and expressed in percentage of wet-basis moisture content. Finally, each swimbladder was
183 dissected and examined macroscopically to determine the presence of the parasite *A. crassus*
184 (Girard and Elie, 2007). Parasite identification followed descriptions of Moravec and
185 Taraschewski (1988).

186 Metal analyses (Ag, As, Cd, Cr, Cu, Ni, Pb, Se, Zn, Hg) were carried out in liver and
187 muscle samples as previously described by Pannetier et al. (2016). The concentrations of seven
188 indicators of PCBs (CB50+28, CB 52, CB 101, CB 118, CB 138, CB 153 and CB 180), eight
189 OCPs (2,4'-DDE, 4,4'-DDE, 2,4'-DDD, 4,4'-DDD, 2,4'-DDT and 4,4'-DDT, 2,4'-DDT,
190 hexachlorobenzene (HCB) and lindane) and eight PBDEs (BDE 28, 47, 49, 99, 100, 153, 154,
191 183, 209), were performed on muscle samples following the procedures described by Caron et al.
192 (2016).

193

194 2.3 EROD activity

195 Supplies and chemicals for EROD assay were purchased from Sigma–Aldrich (St-Louis,
196 MO, USA). EROD activity was measured using a spectrofluorimetric method adapted for
197 microplates by Fragoso et al. (1998). This measure is based on the increase of the fluorescence in

198 the reaction medium due to the transformation of 7-ethoxyresorufin (7-ER) into resorufin. EROD
199 activity was measured in the post mitochondrial supernatant prepared from homogenates of 50
200 mg crushed liver in 1 ml of N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid (HEPES) (0.1
201 M, pH = 7.8) grinding buffer and centrifuged at 10 000 g during 20 min at 4°C. Reaction
202 mixtures contained 50 µl of supernatant in HEPES buffer, 10 µl of nicotinamide adenine
203 dinucleotide phosphate (NADPH, 20 mg/ml), and 50 µl of 7-ethoxyresorufin (0.024 mg/ml) in
204 DiMethylSulfOxide (DMSO). The activity was quantified in 96-well microplates using a
205 Cytofluor II® plate reader (excitation 530 nm, emission 590 nm, PerSeptive Biosystems,
206 Framingham, MA, USA). The fluorescence of resorufin was measured at 60 s intervals during 13
207 min. Fluorescence readings were compared to a resorufin standard curve. All samples were
208 assayed in triplicate. Total protein concentration of the supernatant was measured with Bradford
209 reagent, using bovine serum albumin as a protein standard (BIO-RAD Laboratories, Hercules,
210 CA, USA). EROD activity was expressed as pmoles resorufin/min/mg protein. In addition to
211 randomized samples from different sites on each microplate, a positive control was added to
212 assess the quality and repeatability of measurements. Liver homogenates from pooled livers of
213 15 Atlantic tomcod which were injected 20 mg/kg benzo[a]pyrene (10 mg/ml in corn oil)
214 intraperitoneally and sampled 48 h after treatment were used as positive controls (Couillard et
215 al., 2004). Coefficients of variation among samples and positive control replicates were < 20%,
216 and the coefficient of variation of the sample means was < 25%.

217

218 2.4 Antioxidant enzymes

219 Catalase (CAT) and superoxide dismutase (SOD) enzymes were measured in cytosolic
220 supernatant prepared from homogenates of 50 mg crushed liver in HEPES buffer. Liver samples
221 were constantly kept on ice during homogenization. An aliquot of the homogenate was
222 centrifuged at 1500 g for 5 min at 4°C and the supernatant was stored at – 80°C for SOD assays
223 whereas a second aliquot was centrifuged at 10 000 g for 15 min at 4°C and the supernatant was
224 stored at – 80°C for CAT assays. Catalase and SOD activities were measured using commercial
225 kits from Cayman Chemical Company Inc. (Ann Arbor, MI, USA) in 96 well microplates with a
226 UV/visible spectrophotometer (Cary 50, Varian inc.). The CAT assay is based on the reaction of
227 the enzyme with methanol in the presence of an optimal concentration of H₂O₂. The generated
228 formaldehyde is allowed to react with 4-amino-3-hydrazine-5-mercapto-1,2,4-triazole (as the
229 chromogen). The aldehydes, upon oxidation, change from colorless to a purple color that is

230 measured spectrophotometrically at 540 nm. One unit (U) of CAT activity is defined as the
231 amount of enzyme that causes the formation of 1 nmol of formaldehyde per minute at 25°C. The
232 SOD assay utilizes a tetrazolium salt for detection of superoxide anion radicals generated by
233 xanthine oxidase and hypoxanthine measured spectrophotometrically at 450 nm. One unit of
234 SOD is defined as the amount of enzyme needed to exhibit 50% dismutation of the superoxide
235 radical. All samples were tested in triplicate and each absorbance reading was compared to a
236 standard curve and a positive control. Total protein concentration of the supernatant was
237 determined using the Bradford (1976) method using bovine serum albumin as standard and CAT
238 and SOD activities were expressed as $\mu\text{mol}/\text{min}/\text{mg}$ protein.

239

240 2.5 Histopathological biomarkers

241 The spleen was selected as a tissue for the examination of the impacts of contaminants on
242 histopathological biomarkers in fish (Au, 2004). A transversal section from the central part of
243 spleen was fixed in 10% phosphate-buffered formalin within minutes after death. Samples were
244 then embedded in paraffin and 6 μm thick sections were stained with Hematoxylin and Eosin
245 (HE) (Luna, 1968). Two additional sections of spleen were stained with Perls' Prussian blue
246 staining for ferric iron associated with hemosiderin deposits or with Schmorl staining for
247 lipofuscin deposits (Luna, 1968).

248 To be categorised as a MMC, a MMC had to be composed of three or more macrophages
249 containing yellow to dark brown pigments within their cytoplasm (Couillard and Hodson, 1996).
250 In sections of spleen stained with HE, the density of MMCs was evaluated by counting the
251 number of MMCs at 100X magnification in three randomly selected microscopic fields and by
252 dividing this number by the area of the fields examined (2.86 mm^2). The surface areas of 10
253 randomly selected MMC were measured with an image analysis system (Image-Pro Express
254 software, version 6.0) and mean of surface area of MMCs were calculated for each fish. The
255 proportion of 25 randomly selected MMCs that stained positively with Perls' Prussian Blue for
256 hemosiderin and Schmorl for lipofuscin was evaluated at 100X magnification. The intensity of
257 staining was graded from 1 to 5 (1: no macrophage stained; 2: 0-25% stained; 3: 25-50% stained;
258 4: 50-75% stained and 5: > 75%). The reproducibility of the methods was assessed by repeated
259 measurements of a subsample ($\approx 10\%$) of slides. The coefficient of variation among the mean of
260 the third quartile (Q3) for the density and for the surface of MMCs separately in twelve eels were
261 < 25% (repeated 3 times by two different observers for each eel). All tissues were examined

262 blindly and randomly without the observer knowing the site of origin of eels. Density was
263 expressed as number of MMC/mm² and surface as mm²/MMC.

264

265 2.6 Statistical analyses

266 Analyses of variance were conducted using Prism v. 7.0 (GraphPad Software Inc., San
267 Diego, CA, USA) and multivariate analyses were conducted using XLSTAT-ADA (Addinsoft,
268 USA). Male eels were removed due to a small sample size (N=3 in the GS and N=0 in the SLS).
269 Eels ≤ 2 years old and ≥ 20 years old were also removed, because they were restricted to a small
270 number of sites/sampling year. The total number of eels retained for statistical analyses of
271 variance were 105 in *A. rostrata* and 111 in *A. anguilla* (Table 2).

272 Before performing the tests, homoscedasticity and normality of data were verified using
273 the Bartlett and the Shapiro-Wilk tests respectively. Biological characteristics, contaminant
274 concentrations and biomarker data were first compared among sampling years within sites using
275 the student t-test ($p \leq 0.05$). Then, data were compared among sampling sites within species
276 using a one-way ANOVA test followed by Tukey-Kramer HSD test to identify significant
277 differences between pairs ($p \leq 0.05$). A non-parametric Kruskal-Wallis test (KW) followed by
278 Dunn's test was performed when the ANOVA's assumptions were not met.

279 Principal Component Analyses (PCAs) and Kendall rank correlation were performed to
280 explore the relationships among biomarker responses, contaminants and biological
281 characteristics for the two years combined in each species. Prior to PCA analyses, correlation
282 matrices were performed by category of contaminants for PCBs, DDTs, and for PBDEs in order
283 to group contaminant variables correlated with each other from the same category and improve
284 the quality of PCA (data not shown). Due to very strong correlations among all PCB congeners,
285 the sum of 7 PCBs was used. For contaminants of the DDT family, only the muscle
286 concentrations of 4,4'-DDD, 2,4'-DDE and 4,4'-DDE were retained. For contaminants of the
287 PBDE family, only PBDEs 47, 100, 153 and 154 were included. Finally, essential metals (Cu, Se
288 and Zn) in liver and muscle were removed from the analysis since their concentrations were
289 unlikely to pose a risk to the health of eels (Pannetier et al., 2016). Consequently, the non-
290 essential metals Ag, Cd, Pb, Hg and As in muscle and in liver (except for liver Hg due to high
291 proportion of missing values) were kept for PCAs. Only individuals for which measurements
292 were available for each variable were retained for the PCAs (N=82 for *A. rostrata* and N=87 for
293 *A. anguilla*).

294 In summary, a set of 17 variables including 9 categories of contaminants [sum of PCBs,
295 sum of DDT and metabolites, hexachlorobenzene, lindane, sum of PBDEs, muscle Ag, Cd, Pb,
296 Hg and As, liver Ag, Cd, Pb and As], 5 biological variables (BL, BM, lipids, age, number of *A.*
297 *crassus*) and 7 biomarker variables (EROD, CAT, SOD, density and surface of MMCs,
298 lipofuscin and hemosiderin pigments) were considered in a total of 82 yellow eels for *A. rostrata*
299 and 87 yellow eels for *A. anguilla* captured in both years. For contaminants, non-detectable data
300 were replaced by $\frac{1}{2}$ of their Limits of Detection (LoDs). Kendall rank correlations were used to
301 assess the strength relationships among biomarkers, contaminants and biological characteristics.
302 The significance level (p value) of the Kendall's correlations was fixed at 0.05.

303

304 **3. Results and discussion**

305 3.1 Morphometry and level of contamination in *A. rostrata* and *A. anguilla*

306 Since contamination and biological characteristics of eels from this study have been
307 reported in companion studies (Baillon et al., 2015b; Baillon et al., 2016; Caron et al., 2016;
308 Laporte et al., 2016; Pannetier et al., 2016), they will only be briefly summarized here.

309 For *A. rostrata*, fish from the fluvial lakes SPL and SFL were generally larger, older and fatter
310 than fish from SOR and SJR (significantly in 2011 but not in 2012) and no *A. crassus* was
311 detected in American eels (Tables 2 and S.1). As reported by Baillon et al. (2015b), eels from the
312 fluvial lakes SPL and SFL were more contaminated with organic chemicals such as PCBs,
313 PBDEs and DDT and metabolites than fish from the downstream sites SOR and SJR (Table S.2).
314 For non-essential metals, significantly higher concentrations of Cd and Pb were observed in eels
315 sampled in the downstream site SOR compared to the upstream site SFL (Table S.3 and S.4).
316 Muscle Hg was significantly higher in eels from upstream sites SPL and SFL compared to SJR
317 (Pannetier et al., 2016). For *A. anguilla*, in 2011, fish from GIR were significantly longer,
318 heavier and fatter than eels from all other sites and in 2012, the size of eels from GIR was
319 comparable to that of fish from GAR and CER (Table S.1). In both years, eels from GAR were
320 significantly older than eels from DOR and CER and eels from DOR were significantly smaller
321 and lighter than eels from the other sites. A concentration gradient of increasing organic
322 contamination (PCBs, PBDEs and DDT and metabolites) was observed along the French sites in
323 the order CER-DOR-GAR-GIR (Table S.2). High tissue concentrations of Cd (muscle and liver)
324 and Pb (liver) were detected in eels from the most contaminated sites GIR and GAR and

325 surprisingly also in eels from DOR, a site originally considered as clean (Pannetier et al., 2016).
326 In both species, higher As concentrations were found in fish from the reference sites CER and
327 SJR and from the contaminated site GIR which had in common an elevated salinity (Pannetier et
328 al., 2016). For *A. anguilla* from the most contaminated site GIR, muscle PCBs, DDT and
329 metabolites and Cd concentrations were 10 and 11-folds higher than in *A. rostrata* from the
330 contaminated sites SPL (Tables S.2 and S.4). Eels from SPL had two-fold higher muscle PBDE
331 concentrations and three-fold higher muscle Hg concentrations than eels from GIR (Tables S.2
332 and S.3). Eels from CER had three-fold higher concentrations of muscle As than eels from SJR
333 (Tables S.2 and S.4). Finally, the ocular index (OI) reached $2.7 \pm (2.0, 3.8)$ (median \pm (Q1, Q3);
334 $n=108$) and $3.9 \pm (3.3, 4.8)$ (median \pm (Q1, Q3); $n=111$) for *A. rostrata* and for *A. anguilla*
335 respectively. In pre-migrating silver eels, this index should be > 6.5 (Cottrill, 2002; Pankhurst,
336 1982). Hence, our eels were not pre-migrating and were all yellow eels.

337

338 3.2 Patterns of biological characteristics, contaminants and biomarker responses in *A.*
339 *rostrata*

340 The first 3 principal components of the PCA (PC1, PC2 and PC3) covered 37 % of the
341 total variability of the overall data (Table 1). Principal component 1 was mainly composed of
342 biological variables and organic and inorganic contaminants and it separated in two distinct
343 clusters eels from the contaminated sites SFL and SPL from the reference sites SOR and SJR
344 (Figs. 2 a and b). The previously described associations between high concentrations of
345 persistent organic contaminants, higher fat content and older age (Larsson et al, 1990) are
346 illustrated on this axis. No significant association between muscle lipids and biomarker
347 responses were observed although muscle lipids were positively correlated to concentrations of
348 lipophilic organic contaminants (Table S.6). Thus, there is no evidence of a detrimental effect of
349 these compounds on lipid accumulation in yellow eels. Principal component 2 was mainly
350 composed of metal contaminants (Figs. 2 a and b). The majority of eels from SJR were
351 associated with muscle and liver As in agreement with higher As levels (although non-
352 significant, Tables S.3 and S.4) compared to the group of eels from SOR which was associated
353 with liver Cd. Higher As concentrations in fish from SJR are typical of fish living in marine
354 environments (Pannetier et al., 2016) and could be explained by differences in migratory
355 behavior, spatial distribution and feeding. Further studies combining chemical analyses of
356 otoliths with biomarker and contaminant measurements are recommended to verify this

357 hypothesis. No biomarker variables were distributed along the PC1-axis or the PC2-axis (Table 1
358 and Fig. 2 a). Since PC1 and PC2 discriminated sampling sites, our results are consistent with the
359 absence of significant differences for most biomarkers (EROD activity, density, surface and
360 accumulation of pigments in splenic MMC) among sampling sites and between years presented
361 in Table 2.

362 The density and surface of MMCs and the deposition of hemosiderin pigments were
363 distributed along the PC3-axis (Table 1). Hemosiderin pigments in spleen MMC were associated
364 to a group of eels from SOR and SJR sites (Fig. 2 a). Moreover, they were negatively correlated
365 with size and organic contaminants and weakly positively correlated with muscle and liver Pb
366 (Table S.5). An experimental study using common carp (*Cyprinus carpio* L.) has already
367 demonstrated hemosiderin deposition in fish exposed to a mixture of metals (Cd, Pb, Cr and Ni)
368 compared to controls (Vinodhini and Narayanan, 2008). However, given the weak correlation
369 and the lack of significant difference among sites (Fig. 2 a), it seems difficult to conclude for an
370 effect of Pb on deposition of hemosiderin pigment. Note however that in a companion study
371 from the same research project, Caron et al. (2016) identified an association between Ag, Pb and
372 As contamination and an inhibition of the hepatic enzyme G6PDH in the same eels as those used
373 in this study, suggesting an effect of metal contamination in these eels. This enzyme is involved
374 in lipid metabolism and is closely related to antioxidant response (Martinez-Alvarez et al., 2005).
375 Differences in environmental conditions in estuarine compared to fluvial sites could contribute to
376 increase hemosiderin deposition in SJR and SOR sites through exposure to an unknown
377 biological or chemical hemolytic agent.

378 Catalase and SOD activities were not significantly correlated with PC1, PC2 or PC3 axes
379 and their explained variance (%) was low (Table 1). Although CAT activity was not correlated
380 with biological characteristics or contaminants (Table S.5), eels from SOR expressed
381 significantly higher activities of hepatic CAT than fish from other sites in 2011 (but not in 2012)
382 and a significant inter-annual difference was observed, with higher CAT activity in 2011 in SOR
383 (Table 2). SOD activity was negatively correlated with size, muscle lipid content and PCBs,
384 more weakly with age, PBDEs and DDT and metabolites and was associated with a group of eels
385 from SOR and with some fish from SPL (Fig. 2 and Table S.5). This observation is consistent
386 with significantly higher liver SOD activity found in eels from SOR in both years and with the
387 significant inter-annual difference observed for SOD, higher in 2012 in SPL fish (Table 2). SOD
388 activity was weakly positively correlated to liver Cd which could possibly explain the high level

389 of SOD activity in SOR and SPL fish (Table S.5). As shown in Fig. 2, a majority of eels from
390 SOR and some eels from SPL were associated with the liver Cd vector in agreement with the
391 significantly higher liver Cd observed in eels from SOR compared to eels from SJR and SPL
392 (Table S.4). Similar results were observed by Martinez-Alvarez et al. (2005), who reported that
393 Cd exposure could cause an increase in antioxidant enzymes by competing with essential metals
394 on protein-binding sites. Further experimental studies are needed to confirm the potential cause-
395 effect relationship behind this weak association.

396 Although EROD activity did not vary significantly among sites and years, in 2011
397 however, EROD activity was highly variable in eels from SPL compared to the other sites (with
398 a high 75th percentile of 31.6 compared to 15.7 and 10.3 for SOR and SJR sites respectively,
399 Table 2).

400 Previous studies conducted in the Gironde and St. Lawrence River systems have reported
401 that some eels during their growth phase migrate from the upstream to the downstream sections
402 and inversely (Daverat and Tomás, 2006; Thibault et al., 2007). Inter-individual variations in
403 migratory behaviour and growth rates (Patey et al., submitted), in addition to variations in diet
404 selection (Pegg et al., 2015) could affect contaminant uptake. Although a wide range of
405 persistent contaminants (PCBs, PBDEs, pesticides and metals) were measured in eels in this
406 study, other contaminants could influence the biomarker responses. PAHs are known to induce
407 both EROD and antioxidant enzyme activities (Martinez-Alvarez et al., 2005; Whyte et al.,
408 2008). For example, in field and experimental studies, Gravato et al. (2006, 2010) reported
409 significant increases in liver EROD activity and antioxidant capacities in European eels
410 contaminated by PAHs. Seasonal or inter-annual variations in natural factors such as temperature
411 and salinity could also influence antioxidant defense and biotransformation enzyme activities in
412 fish, even if exposure remains constant (Martinez-Alvarez et al., 2005; Whyte et al., 2008;
413 Chainy et al., 2016). Eels sampled in tributaries from the downstream St. Lawrence system (SOR
414 and SJR), exposed to a strong salinity gradient over a short distance, are possibly more prone to
415 variations in salinity, metal contaminants and temperature which may possibly cause inter-
416 annual variations in antioxidant enzymes (Thibault et al., 2007; Pannetier et al., 2016).

417

418 3.3 Patterns of biological characteristics, contaminants and biomarker responses in *A.*
419 *anguilla*

420 The first 3 principal components of the PCA covered 39 % of the variability of the
421 overall data (Table 1). PC1 was mainly composed of age, organic and inorganic contaminants
422 and mainly contributed to separate the most contaminated sites GIR and GAR from the reference
423 site CER (Fig. 3 and Table 1). Like for *A. rostrata*, *A. anguilla* muscle lipids were significantly
424 correlated with morphometric and organic contaminants and no strong correlations were
425 observed among biomarker responses and muscle lipids (Kendall rank correlation < 0.200 , Table
426 S.6) suggesting no impairment of contaminants on fat reserves.

427 Hemosiderin pigment deposition in splenic MMC was distributed along the PC2-axis
428 (Table 1 and Fig. 3 a). Hemosiderin pigment, muscle and liver As scored positively on PC2 and
429 were significantly and positively correlated with each other (Fig. 3 a and Table S.6). Moreover,
430 vectors of these variables are oriented between eels from CER and GIR (Fig. 3 b). This
431 observation is consistent with significantly higher values in fish from the salt marsh site CER
432 and intermediate values in fish from the estuarine site GIR observed in both years (Fig. 1 a).
433 Moreover, it was in agreement with the observation that eels living in brackish or saltwater
434 environments accumulate more As than their freshwater counterparts (Pannetier et al., 2016).
435 Sorensen and Smith (1981) demonstrated the presence of As in hemosiderin granules in the liver
436 of adult channel catfish (*Ictalurus punctatus*) exposed experimentally to As and suggested that
437 this deposit constituted a mechanism for storage and detoxification. The presence of hemosiderin
438 pigments is linked to a breakdown of red blood cells which could cause hemosiderosis,
439 especially in the spleen, and increase the size and the number of MMCs in fish from
440 contaminated areas (Khan and Kiceniuk, 1984). Further studies are needed to determine if eels
441 living in saltwater environments are exposed, in addition to As, to other chemical or biological
442 hemolytic agents leading to increased deposition of hemosiderin in their spleen.

443 Although other biomarkers were not significantly correlated with PC1, PC2 and PC3 axes
444 and their explained variance (%) were low (Table 1), contrary to what was observed for *A.*
445 *rostrata*, in *A. anguilla* most biomarkers displayed significant spatial and inter-annual variations
446 (Table 2). Variations in lipofuscin pigments showed significant differences among sites in 2011,
447 with higher values in fish from GAR compared to fish from DOR (Fig. 1 b). Lipofuscin
448 deposition in splenic MMC was more associated with age and GAR fish (Table S.6 and Fig. 3 b).
449 Moreover, eels from GAR were older than fish from other French sampling sites (Table S.1).
450 Lipofuscin is known as an “aging pigment” and an increase in lipofuscin pigments in MMCs is
451 generally found in older fish (Wolke, 1992; Agius and Roberts, 2003). Therefore, in *A. anguilla*,

452 lipofuscin pigments globally increased in older and more contaminated fish, mostly found in
453 GAR, in contrast with *A. rostrata* in which lipofuscin deposition was not significantly correlated
454 to either of these variables.

455 Values of density of MMCs were significantly higher in GAR compared to DOR, with
456 intermediate values for fish from GIR and CER in 2012 (Table 2). In 2011, the surface of MMCs
457 expressed significantly higher values in eels from GIR compared to DOR. The PCA analysis
458 showed that the strongest correlations were between density of MMC and muscle Hg and Cd
459 (Kendall's tau = 0.281 and 0.235 respectively) and between surface of MMC and age, BL and
460 BM (Kendall's tau = 0.384, 0.337 and 0.317, respectively, Table S.6). Several studies have
461 reported on correlations between increased number of MMC in fish exposed to Hg and other
462 metals (Meinelt et al., 1997; Agius and Roberts, 2003; Schwindt et al., 2008). Khan et al. (1994)
463 observed an increase in the size of splenic MMCs with increasing length and weight of winter
464 flounder living in a contaminated area and Couillard and Hodson (1996) also reported an
465 increase of surface of MMCs in the spleen of old wild Atlantic tomcod collected from estuarine
466 sites receiving pulp and paper mill effluents. Supported by these studies, the increase in the
467 density of MMCs in eels from GAR with Hg and Cd contamination and the increase in the
468 surface of spleen MMCs in eels from GIR was possibly associated with aging. Since the
469 Kendall's tau of these correlations was weak, we do not exclude that other potential
470 contaminants and/or natural factors not measured could influence the response of these
471 biomarkers.

472 Catalase activity was more strongly correlated with higher organic contaminants such as
473 DDT and metabolites, PBDEs and PCBs and associated to eels from GAR and GIR (Fig. 3 and
474 Table S.6). In addition, fish from GIR, GAR and DOR expressed significantly higher liver CAT
475 activities compared to CER fish in 2011 (Table 2). In 2012, only fish from GIR had significantly
476 higher CAT activities compared to GAR fish (Table 2). Liver CAT activity was significantly
477 higher in 2012 than in 2011 but only in fish from CER, in contrast to liver SOD which remained
478 unchanged for these fish, but which decreased between 2011 and 2012 in GAR and GIR. To
479 support the possible effect of organic contaminants on hepatic CAT enzyme in GAR and GIR,
480 other field studies have reported significant increases in CAT activity in yellow *A. anguilla*
481 exposed to organic contaminants (Regoli et al., 2003; Ahmad et al., 2006; Buet et al., 2006).

482 No significant relationships between EROD activity and tissue contaminant
483 concentrations were observed indicating that this biomarker response was not associated with

484 exposure to the measured persistent contaminants in the Gironde system (Tables S.6). Although
485 concentrations of organic contaminants in eels from GAR and GIR were higher in 2011, EROD
486 activity was twice higher in 2012 in fish from GIR compared to fish from DOR and CER and
487 was also higher in fish from GAR compared to DOR (Table 2). We hypothesize that other
488 chemical factors not measured, such as PAHs, may have contributed to the higher EROD activity
489 found in eels from GAR and GIR in 2012. Elevated concentrations of PAHs and pyrene were
490 detected in the muscle tissue of European flounder (*Platichthys flesus*) in the Gironde Estuary
491 (Laroche et al. 2012). Further studies are needed to confirm the link between PAH exposure and
492 EROD induction in eels from GAR and GIR.

493 Finally, the parasite *A. crassus* was detected only in *A. anguilla* (Table 2). The number of
494 *A. crassus* was not correlated with any biological, contaminant or biomarker variables (Table
495 S.6). In 2011, fish from the freshwater sites DOR and GAR presented a significantly higher rate
496 of infection compared to samples from the brackish water site GIR and the salt marsh site CER,
497 consistent with field and experimental studies having demonstrated that infection levels of *A.*
498 *crassus* decrease with an increase in salinity (Kirk et al., 2000; Morrison et al., 2003; Lefebvre
499 and Crivelli, 2012).

500

501 4. Conclusions

502 Our study revealed similarities in the biomarkers examined between *A. anguilla* and *A.*
503 *rostrata*, supporting their use in comparative ecotoxicological studies. The basal levels of EROD
504 activity and of biomarkers of histopathology were similar in eels of both species from reference
505 sites. A high deposition of hemosiderin was observed in eels from the most brackish sites in both
506 species. However, the cause of this deposition is uncertain. In *A. anguilla*, it was strongly
507 associated to As accumulation whereas in *A. rostrata*, it was weakly associated to Pb. Further
508 studies are needed on the relationship between environmental variations, migration and feeding
509 patterns and accumulation of potentially hemolytic agents in Atlantic eels growing in saline
510 water, and on the potential impact of this accumulation on blood cells integrity and on the
511 immune function.

512 In comparison to its American cousin from the St. Lawrence system, *A. anguilla* from
513 Southwest France may be more affected by contaminants. Higher CAT activity, density of MMC
514 and hemosiderin pigment were observed in eels from the contaminated sites GAR and GIR and
515 were related to organic and metal contaminants such as DDTs, PCBs, PBDEs and muscle Hg and

516 Cd. This, in association with the higher concentrations of PCBs, DDTs and metals observed in
517 the tissues of *A. anguilla* from GAR and GIR compared to *A. rostrata*, suggests a more important
518 risk of impacts of these contaminants in these eels compared to American eels from the more
519 contaminated sites SPL and SFL. For *A. rostrata*, in both study years, neither the PCA nor
520 ANOVA identified major influences of persistent organic or inorganic contaminants on EROD,
521 catalase or histopathological biomarkers.

522 Field studies have already reported that the sensitivity of biomarkers could vary from one
523 habitat to another in response to the complex mixtures of contaminants and other environmental
524 conditions (Wolke, 1992; Van der Oost et al., 2003). Contaminants such as PAHs, dioxins and
525 furans, not measured in our study, are known to induce biotransformation and antioxidant
526 enzymes in the short-term and histopathological biomarkers over longer periods of exposure
527 (Van der Oost et al., 2003). The yellow stage of both species are not strictly catadromous and
528 perform short migrations in order to forage food in more productive environments (Daverat and
529 Tomás, 2006; Thibault et al., 2007). Even if these migrations are short (≈ 10 km) compared to
530 the reproductive migration of silver eels, they are of considerable importance because they occur
531 during the prolonged growth stage and involve switching between environments varying widely
532 in contaminants and other abiotic factors such as temperature and salinity (Thibault et al., 2007).
533 Finally, the small number of eels that could be collected in this study and the wide range of
534 estimated age (2 to 18 years old) may constitute biases and impose a requirement to be prudent
535 regarding the overall conclusions. The peculiar biology of these species, which grow at widely
536 different rates depending on their habitat (Patey et al, submitted), makes it impossible to
537 eliminate biases such as size and age among study sites. For that purpose, complementary studies
538 with larger sample size and a greater number of sites are recommended. Nevertheless, this study
539 provides a database to better target and prioritize future studies on the impacts of contaminants
540 on eel health in the Gironde and St. Lawrence River basins. Combined with others (Pierron et al.,
541 2013; Baillon et al., 2015a; Baillon et al., 2015b; Baillon et al., 2016; Caron et al., 2016) it
542 supports that contamination represents a risk to the health of European eels from the Gironde
543 system that could be contributing to the lack of recovery of these fish.

544

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- 1 **Table 1:** Explained variance (%) for each significant principal component (PC) of the three different PCA followed by the significant variables in each PC.
 2 Variables significantly correlated with the principal components (PC1, PC2, PC3) in each PC are noted in bold.

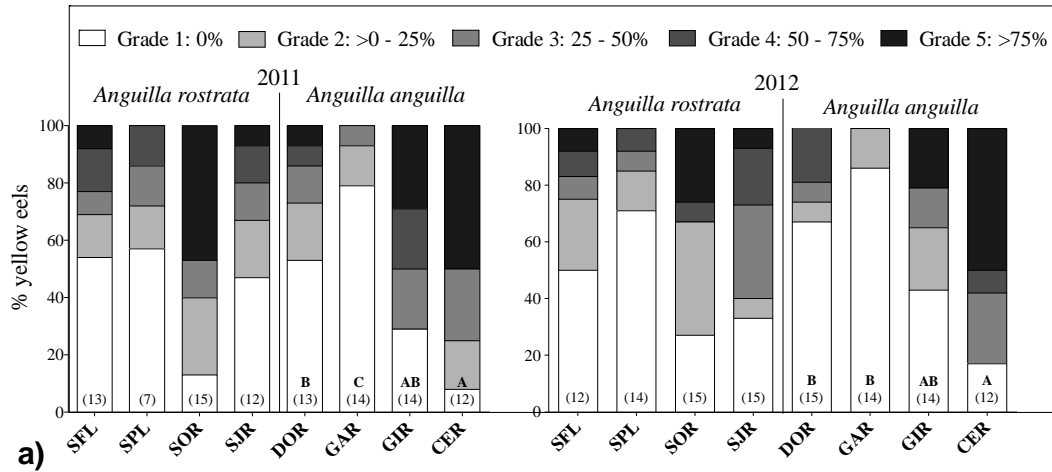
<u>Species</u>	<u>Anguilla rostrata</u>			<u>Anguilla anguilla</u>		
	<u>2011-2012</u>			<u>2011-2012</u>		
	<u>PC1</u>	<u>PC2</u>	<u>PC3</u>	<u>PC1</u>	<u>PC2</u>	<u>PC3</u>
Explained variance (%)	22.21	8.07	6.63	19.72	11.05	8.01
Variables						
Body Length (mm)	11.39	2.64	0.09	5.37	9.82	3.36
Body Mass (g)	11.56	3.67	0.15	4.67	11.39	3.81
Lipid content (%)	8.36	5.48	0.01	4.44	5.01	2.22
Age (year)	2.29	6.38	12.09	8.61	0.02	7.45
Number <i>A. crassus</i>	0.00	0.00	0.00	0.07	9.26	1.02
EROD (pmol/min/mg protein)	0.67	0.18	4.88	0.36	0.03	9.66
Catalase ($\mu\text{mol}/\text{min}/\text{mg}$ protein)	0.07	1.66	1.71	3.58	0.01	1.00
Superoxide dismutase ($\mu\text{mol}/\text{min}/\text{mg}$ protein)	2.45	0.72	1.23	0.09	0.11	4.19
Density of MMCs (number/ mm^2)	0.04	0.38	16.08	1.37	1.06	3.28
Surface of MMCs (MMC/ mm^2)	0.08	1.95	21.72	3.54	2.09	5.45
Hemosiderin pigment (median of grades)	1.68	1.58	13.77	0.00	11.29	5.24
Lipofuscin pigment (median of grades)	0.02	0.01	0.01	1.91	0.00	0.59
Hexachlorobenzene (ng/g dw)	6.64	6.51	2.11	5.72	0.32	6.12
Lindane (ng/g dw)	0.04	0.10	0.14	0.18	7.67	9.81
Sum of DDT and metabolites (ng/g dw)	10.57	1.93	0.00	12.08	1.68	1.12
Sum of PBDEs (ng/g dw)	8.27	0.35	0.35	9.33	3.60	1.07
Sum of PCBs (ng/g dw)	11.30	0.14	0.01	10.77	1.10	3.59
Muscle As ($\mu\text{g}/\text{g}$ dw)	4.87	14.33	0.00	0.39	12.52	4.64
Muscle Ag ($\mu\text{g}/\text{g}$ dw)	0.32	7.54	0.02	0.03	0.84	2.60
Muscle Cd ($\mu\text{g}/\text{g}$ dw)	0.65	0.79	3.71	0.19	2.04	5.95
Muscle Pb ($\mu\text{g}/\text{g}$ dw)	0.40	2.27	0.01	0.07	2.32	0.44
Muscle Hg ($\mu\text{g}/\text{g}$ dw)	4.76	8.01	4.39	7.57	1.89	2.59
Liver As ($\mu\text{g}/\text{g}$ dw)	3.96	13.44	0.71	0.17	12.50	3.98
Liver Ag ($\mu\text{g}/\text{g}$ dw)	3.13	7.67	0.00	6.08	0.65	0.13
Liver Cd ($\mu\text{g}/\text{g}$ dw)	0.50	8.49	13.33	7.03	2.05	0.36
Liver Pb ($\mu\text{g}/\text{g}$ dw)	5.99	3.77	3.50	6.39	0.74	10.34

4 **Table 2:** Spatial and temporal variations of hepatic EROD activity, antioxidant enzymes, splenic melanomacrophage centers (MMCs) and number of *A. crassus*
 5 (median (Q1; Q3) (n)) of yellow *A. rostrata* in the St. Lawrence system and *A. anguilla* in the Gironde system in 2011 and 2012. Different letters indicate
 6 significant differences among sites within species and within sampling years (ANOVA or KW, $p \leq 0.05$). An asterisk * indicates a significant difference between
 7 2011 and 2012 (T-test, $p < 0.05$). DOR= Dordogne; GAR= Garonne; GIR= Gironde; CER= Certes; SFL = Saint-François Lake; SPL = Saint-Pierre Lake; SOR =
 8 Sud-Ouest River; SJR = Saint-Jean River; NA= Absent.

Species	<i>Anguilla rostrata</i>				<i>Anguilla anguilla</i>				
	Biomarkers	Years	SFL	SPL	SOR	SJR	DOR	GAR	GIR
Liver EROD (pmol/min/mg protein)	2011	7.1 (5.7; 8.8) (15)	8.9 (5.2; 31.6) (7)	10.5 (7.9; 15.7) (15)	8.7 (7.6; 10.3) (12)	11.9 (6.3; 13.9) (13)	7.2 (6.1; 14.2) (14)	10.4 (6.9; 12.4) (15)	8.8 (5.1; 11.2) (11)
	2012	5.1 (3.7; 11.7) (12)	7.1 (6.1; 9.9) (14)	11.6 (8.9; 15.9) (15)	11.4 (8.5; 22.5) (15)	9.9 (6.2; 11.0) (15) C	18.6 (13.9; 25.7) (15) AB	19.7 (16.9; 40.2) (14) A	13.3 (8.9; 16.3) (14) BC
Liver Catalase ($\mu\text{mol}/\text{min}/\text{mg}$ protein)	2011	2368 (1827; 2896) (14) BC	4078 (2657; 5638) (7) AB	3991 (3102; 4312) (12) A*	2246 (1971; 2647) (15) C	2319 (1902; 2581) (8) A	2263 (1455; 2904) (14) A	3515 (2495; 4931) (14) A	893(631; 984) (10) B
	2012	2168 (1323; 3319) (12)	2260 (1534; 3598) (14)	1953 (1663; 2641) (13)	2350 (1523; 2887) (15)	1961 (1567; 2408) (12) AB	1834 (1606; 2039) (13) B	2768 (2296; 3589) (14) A	1919 (1544; 3148) (12) AB*
Liver superoxide dismutase ($\mu\text{mol}/\text{min}/\text{mg}$ protein)	2011	5322 (2635; 7629) (12) A	962 (619; 2044) (5) B	5404 (4627; 7261) (12) A	5058 (3793; 5850) (14) A	3910 (3061; 4694) (8)	4281 (2933; 4901) * (14)	3651 (3161; 5261) * (14)	3857 (2762; 4482) (10)
	2012	3320 (2092; 5498) (12) B	3851 (3346; 5305) (14) AB*	6275 (4605; 7646) (14) A	4345 (3008; 6620) (15) AB	3203 (2592; 4175) * (12)	2766 (1775; 3623) (12)	2398 (1827; 3155) (14)	3037 (1957; 3838) (12)
Density of MMCs (number/ mm^2)	2011	2.1 (1.4; 2.8) (15)	2.8 (1.7; 4.5) (7)	2.4 (1.7; 3.1) (15)	2.4 (1.7; 3.1) (15)	4.2 (2.8; 5.1) (13)	2.8 (2.1; 3.9) (14)	3.8 (2.4; 4.5) (15)	2.8 (2.4; 4.5) (11)
	2012	2.8 (2.4; 3) (12)	2.6 (1.7; 3.2) (14)	3.5 (2.4; 3.8) (15)	2.8 (2.1; 3.5) (15)	2.8 (2.1; 4.2) (15) AB	4.5 (2.4; 6.3) (15) A	2.8 (2.3; 5.8) (14) AB	1.9 (1.7; 2.3) (12) B
Surface of MMCs (MMC/ mm^2)	2011	0.003 (0.002; 0.005) (15)	0.005 (0.003; 0.006) (7)	0.005 (0.003; 0.007) (15)	0.002 (0.002; 0.004) (15)	0.003 (0.002; 0.005) (13) B	0.004 (0.003; 0.007) (14) AB	0.007 (0.006; 0.01) (15) A*	0.003 (0.002; 0.007) (11) AB
	2012	0.004 (0.004; 0.006) (12)	0.003 (0.002; 0.004) (14)	0.005 (0.003; 0.007) (15)	0.003 (0.001; 0.006) (15)	0.005 (0.004; 0.008) * (15)	0.006 (0.004; 0.008) (15)	0.005 (0.003; 0.005) (14)	0.005 (0.003; 0.008) (12)
Number of <i>A. crassus</i>	2011	NA	NA	NA	NA	1 (0.5; 3) (13) A*	2.5 (0; 4.3) (14) A	0 (15) B	0 (11) B
	2012	NA	NA	NA	NA	0 (0; 0) (15)	0 (0; 5) (15)	0 (0; 1) (14)	0(0; 0.3) (14)

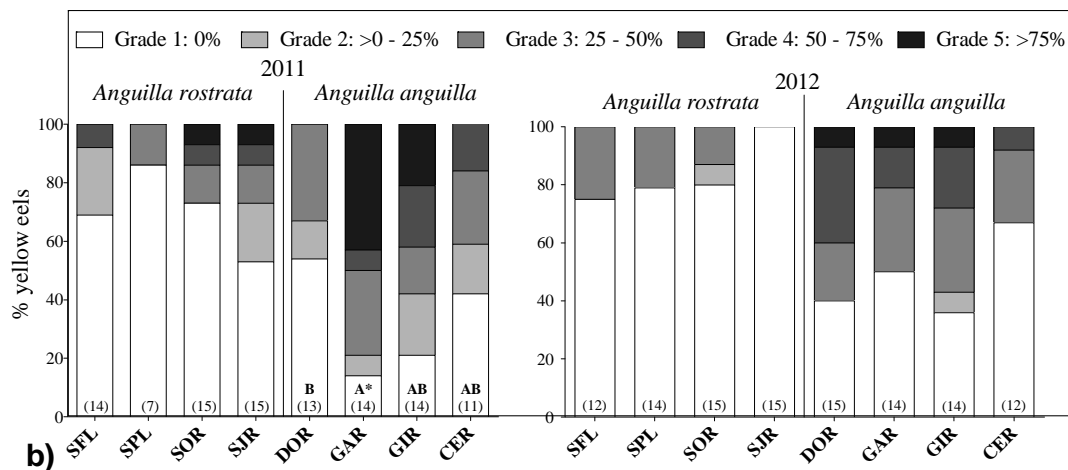
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Hemosiderin pigment



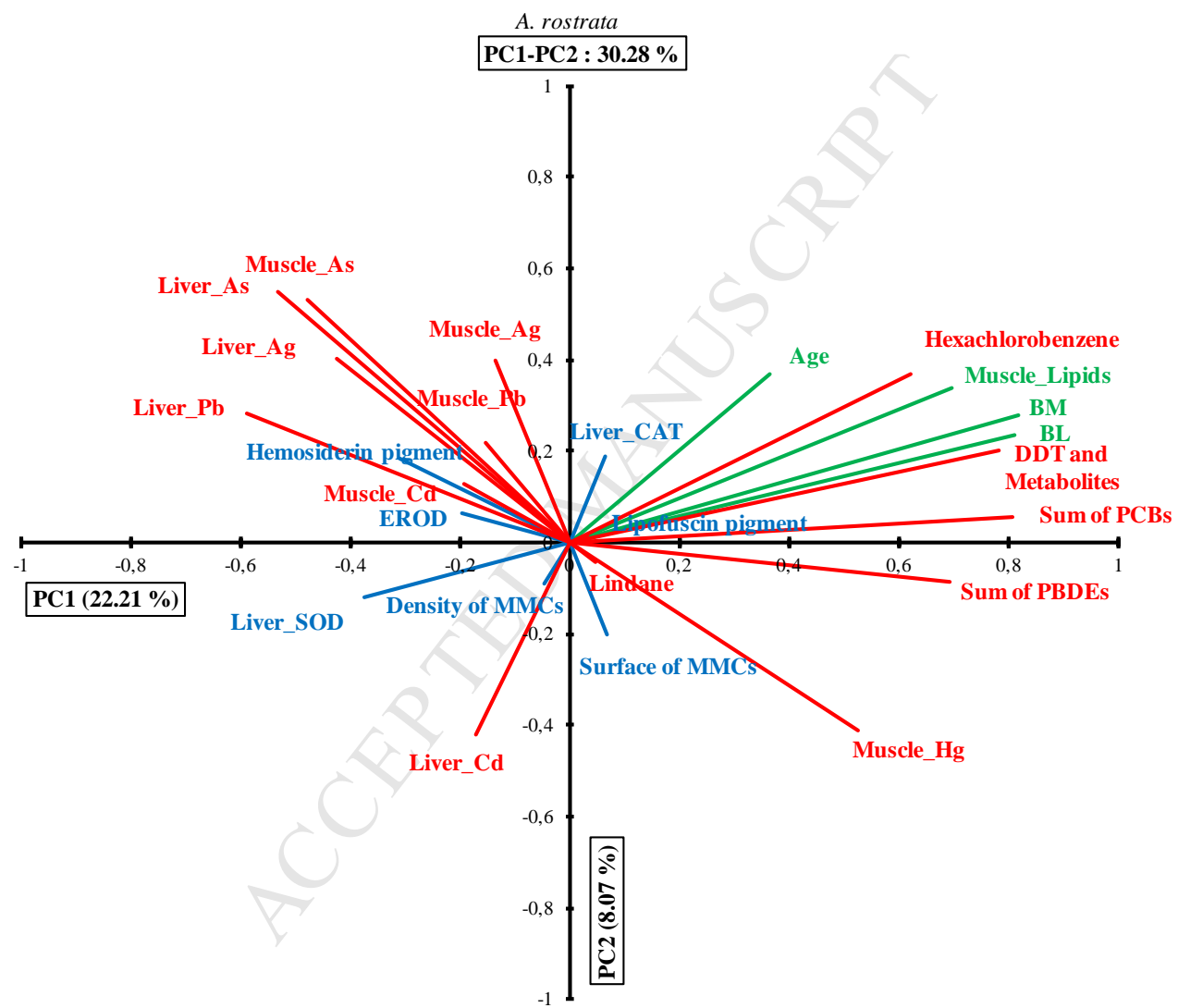
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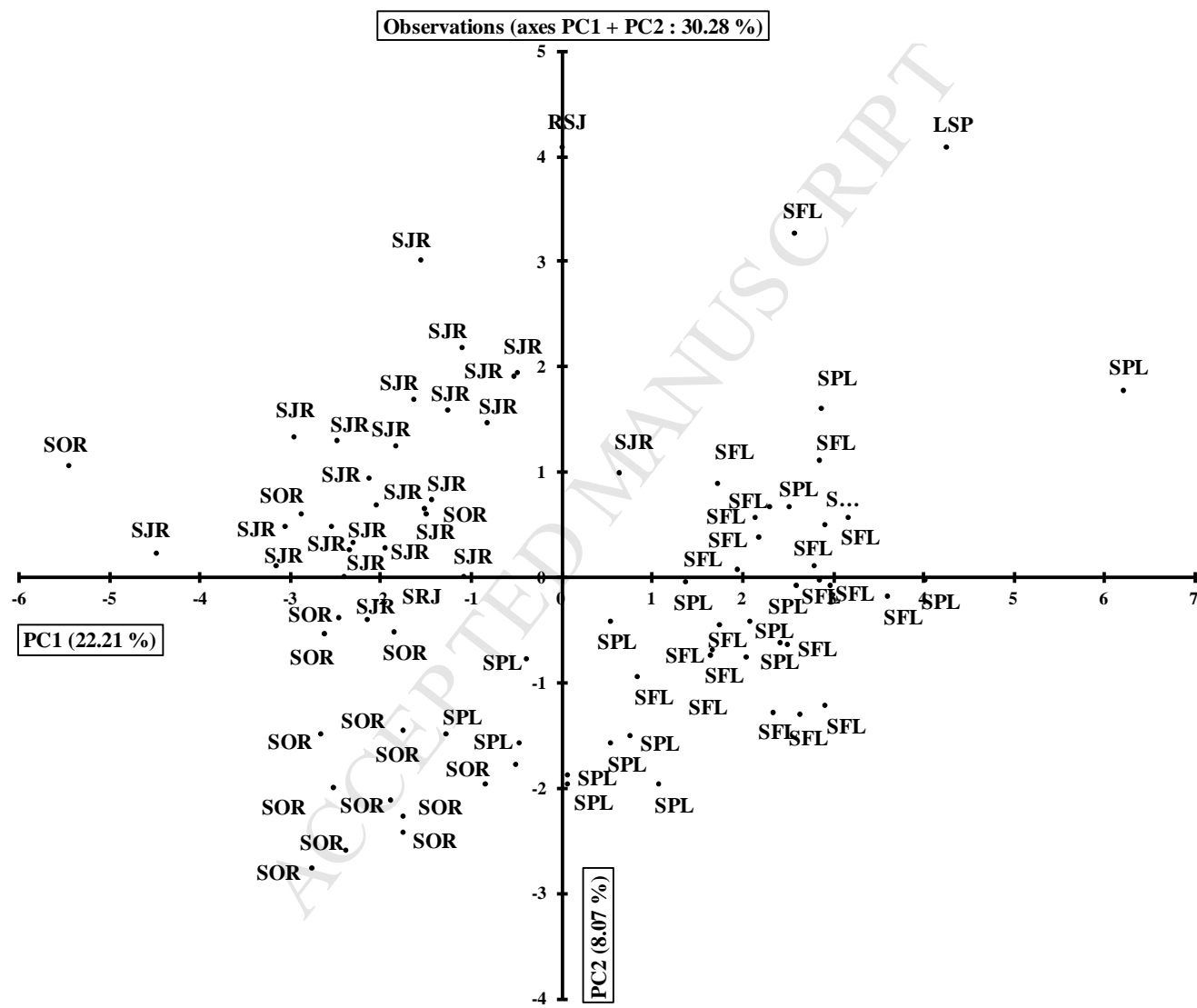
Lipofuscin pigment



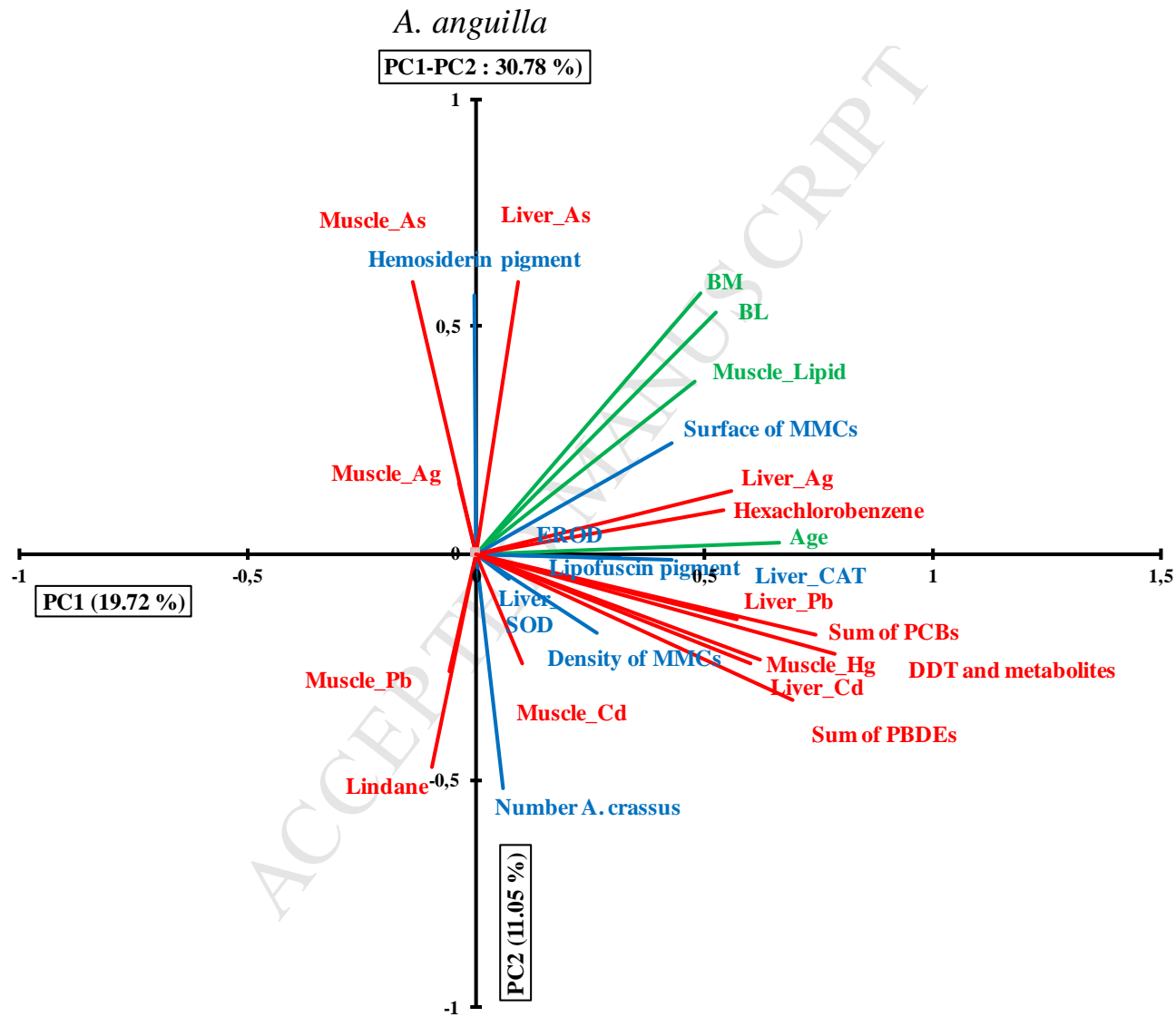
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4 **Figure 1.** Spatio-temporal variations of the relative proportions of *A. rostrata* in the St.
 5 Lawrence System and *A. Anguilla* in the Gironde System with different grades of a)
 6 hemosiderin and b) lipofuscin pigment depositions in spleen MMCs in 2011 and 2012.
 7 Sample size (n) is indicated in parentheses. Different letters indicate significant differences
 8 among sites within species and within a sampling year (Kruskal-Wallis test, $p \leq 0.05$).

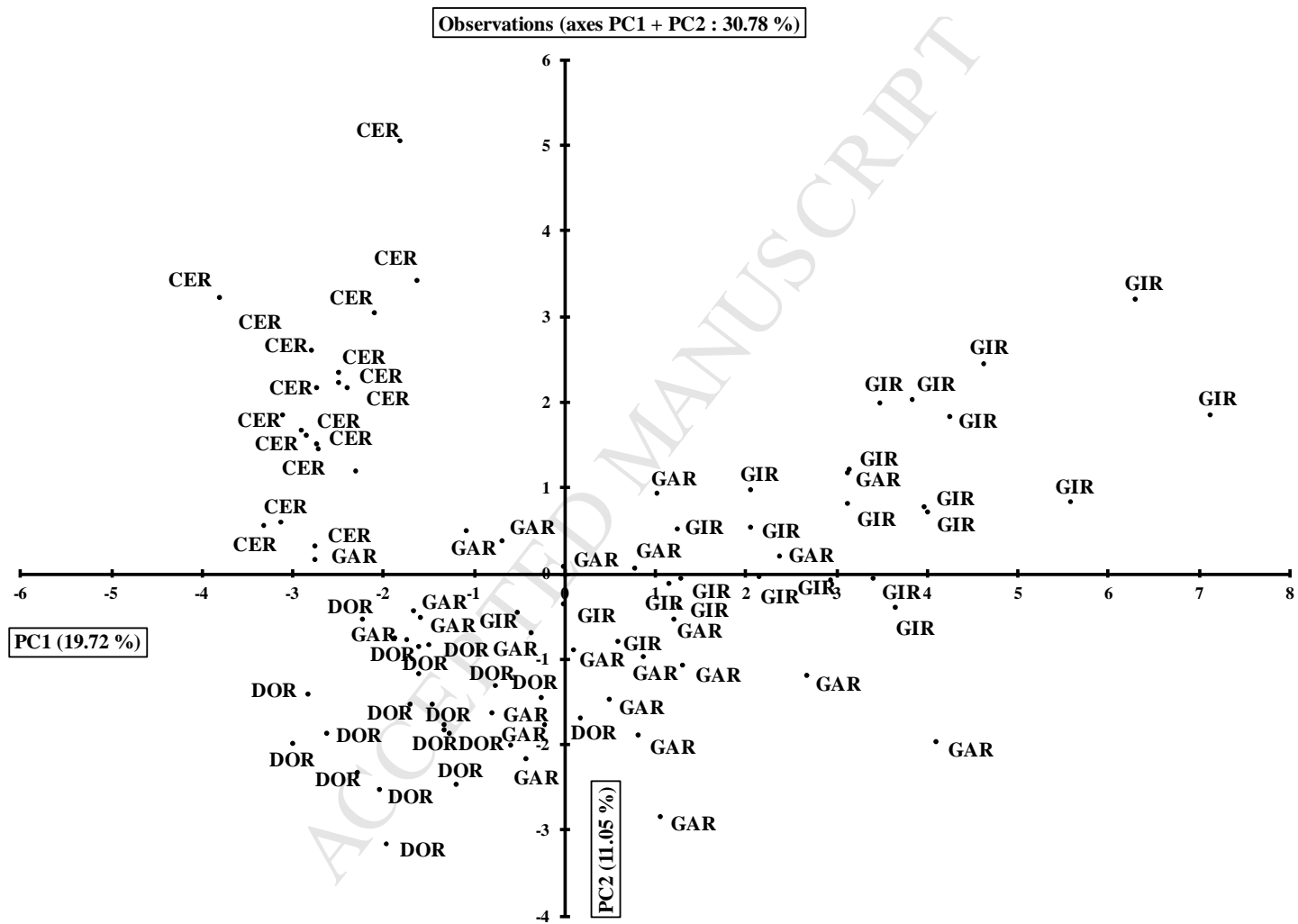




11 **Figure 2.** Principal component analysis (PCA) (n=82) illustrating: a) Relationships between patterns of contamination (red),
12 biomarker response (blue) and biological characteristics (green) in *A. rostrata* in 2011 and 2012 and b) Screening of eels from two
13 fluvial contaminated sites, Lake St. Pierre (LSP) and Lake St. François, (LSF) and two references sites, St Jean River (SJR) and Sud-
14 Ouest River (SOR). Scores for the principal components 1 and 2 were 22.21% and 8.07%, respectively. Sites of capture are identified
15 for each individual. BM: Body Mass, BL: Body Length, MMCs: Melano-Macrophage centers, SOD: superoxide dismutase and CAT:
16 catalase.



18



19

b)

20

21 **Figure 3.** Principal component analysis (PCA) (n=87) illustrating: a) Relationships existing between the pattern of contamination (red),
22 biomarker responses (green) and biological characteristics (blue) in *A. anguilla* in 2011 and 2012 and b) Screening of eels from
23 contaminated sites, Gironde estuary (GIR) and Garonne River (GAR) and two references sites, Dordogne River (DOR) and the salt marsh
24 Certes (CER). Scores for the principal components 1 and 2 were 19.72% and 11.05%, respectively. Sites of capture are identified for each
25 individual. BM: Body Mass, BL: Body Length, MMCs: Melano-Macrophage centers, SOD: superoxide dismutase and CAT: catalase

Highlights

- Patterns of contaminants and biological characteristics differed among sites and species.
- In American eels, biomarker responses did not differ clearly among sites and years.
- In European eels, biomarker responses were higher in contaminated sites.
- High spleen hemosiderin deposition was observed in eels from the most brackish sites.
- European eels may be more affected by contaminants than American eels.