

An integrative perspective on fish health: Environmental and anthropogenic pathways affecting fish stress

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Quentin Schull, Anaïs Beauvieux, Vincent A Viblanc, Luisa Metral, Lina Leclerc, et al.. An integrative perspective on fish health: Environmental and anthropogenic pathways affecting fish stress. Marine Pollution Bulletin, 2023, 194 (Part B), pp.115318. 10.1016/j.marpolbul.2023.115318. hal-04265143

HAL Id: hal-04265143 https://hal.science/hal-04265143

Submitted on 30 Oct 2023

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21 ABSTRACT

Multifactorial studies assessing the cumulative effects of natural and anthropogenic stressors 22 23 on individual stress response are crucial to understand how organisms and populations cope 24 with environmental change. We tested direct and indirect causal pathways through which environmental stressors affect the stress response of wild gilthead seabream in Mediterranean 25 coastal lagoons using an integrative PLS-PM approach. We integrated information on 10 26 27 environmental variables and 36 physiological variables into seven latent variables reflecting lagoons features and fish health. These variables concerned fish lipid reserves, somatic 28 29 structure, inorganic contaminant loads, and individual trophic and stress response levels. This 30 modelling approach allowed explaining 30% of the variance within these 46 variables considered. More importantly, 54% of fish stress response was explained by the dependent 31 lagoon features, fish age, fish diet, fish reserve, fish structure and fish contaminant load latent 32 variables included in our model. This integrative study sheds light on how individuals deal 33 with contrasting environments and multiple ecological pressures. 34

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Key words: Ecosystem, PLS-PM, Eco-physiology, Direct and indirect effects, Fish health,Stress

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41 INTRODUCTION

In natural ecosystems, organisms are subject to a variety of environmental pressures 42 that act simultaneously and vary over time and space. Understanding the consequences of 43 44 environmental variation on individual health, and their long-term consequences on population health and dynamics, requires deciphering the complex interactions between environmental 45 pressures (natural and anthropogenic) and their causal effects on organisms (Doney et al., 46 47 2012; Portner and Farrell, 2008). In complex ecosystems, ecological studies often progress in small steps, identifying and testing the effects of a few handpicked variables on individual 48 organism health and fitness (Doney et al., 2012; Portner and Farrell, 2008). However, where 49 50 interactions among multiple ecological pressures (both intrinsic and extrinsic to the organism) may generate interactive effects, single-stressor based models reach a limit: the resulting 51 global impact of independent environmental pressures on individual organisms might either 52 exceed (i.e. synergism) or fall below (i.e. antagonism) their expected additive effects (Côté et 53 al., 2016; Crain et al., 2008). Therefore, there is an urgent need for multifactorial studies 54 55 where the cumulative effect of multiple environmental pressures on individual organisms is assessed (Brosset et al., 2021; Rosa and Seibel, 2008). This can be achieved through 56 integrative modelling approaches that consider both the putative causal direct and indirect 57 58 pathways through which environmental pressures may affect individual health and fitness.

59 Such approaches should be particularly useful in transitional ecosystems, such as 60 coastal lagoons, where organisms are subject to a high diversity of environmental pressures. 61 Together with estuaries, coastal lagoons form part of a continuum between continental and 62 aquatic ecosystems and are consequently exposed to highly variable hydrological conditions 63 (Pérez-Ruzafa et al., 2011). Situated at the interface between open sea and land, these areas of 64 high biological productivity receive high inputs of organic matter and nutrients of natural and 65 human origin (e.g. agriculture and urban run-offs), but also act as reservoirs for toxic

compounds and disruptors, such as pollutants (Losso and Ghirardini, 2010; Lotze, 2006; 66 Nixon et al., 2007). In such contexts, identifying favourable habitats for a given species 67 and/or life history stage, is faced with the challenging task of assessing how individual 68 organisms cope with the array of interacting environmental factors which they are subjected 69 to in contrasting environments. One way of doing so is to focus on the cumulative effects of 70 multiple natural and anthropogenic stressors on individual physiological processes (Todgham 71 72 and Stillman, 2013), as these are tightly related to individual performance and fitness (Bryndum-Buchholz et al., 2019; Purves et al., 2013; Williams et al., 2008). 73

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75 Stress response as an integrative measure of individual health

76 A key physiological system involved in regulating individual homeostasis is the hypothalamic-pituitary-adrenal (HPA; or interrenal, e.g. in fish) axis, a highly conserved 77 function across vertebrates (Denver, 2009). Subjected to a stressor (external - for instance 78 predation, or internal – for instance fasting), the activation of the HPA axis ultimately leads to 79 the secretion of glucocorticoid (GC) hormones by the adrenal cortex (Wingfield and Romero, 80 2011), whose actions mediate changes in individual energy balance (e.g. mobilisation of 81 energy stores; immunity; cognition; visual acuity; and behaviour, Munck et al. 1984, Johnson 82 et al. 1992). This initial cascade of physiological and behavioural changes enables the 83 organism to cope with acute stressors by mobilising adequate bodily functions, while 84 concurrently inhibiting non-essential functions (e.g. reproduction, digestion) (Wingfield and 85 Romero, 2011). Thus, a commonly used marker of individual stress response is the measure 86 of circulating GC concentrations (Goikoetxea et al., 2021; Harris, 2020; Sadoul and Geffroy, 87 2019). Yet, since the primary function of GCs is to promote adaptive responses to changing 88 environments, GC levels alone are not sufficient to determine whether individuals' coping 89

90 capacities have been overwhelmed by environmental stressors (Boonstra, 2013; Boonstra et
91 al., 2007; MacDougall-Shackleton et al., 2019).

Thus, in addition to increased glucocorticoid levels, physiologically stressed 92 93 individuals often exhibit downstream consequences associated with the long-term overstimulation of emergency responses (Romero, 2004). For instance, individual oxidative 94 stress increases under acute as well as chronic stress and has been related to glucocorticoid 95 96 levels in many animals (Costantini et al., 2011; Lemonnier et al., 2022). Changes in 97 individual immune profiles may also present other downstream metric useful to characterize chronic stress. Chronic stress loads that exceed the normal reactive scope of individuals are 98 99 often associated with immunosuppressive effects (see Martin 2009; Table 1 in Romero et al. 2009) such as down-regulation of inflammatory responses (Cohen et al., 2012) or changes in 100 leukocyte profiles (Davis et al., 2008). Taken together, glucocorticoid levels, oxidative stress 101 status and immune function may present an integrative view of individual exposure to acute 102 and chronic stress and provide a good overview of individual health and how individuals deal 103 104 with their environment.

105

106 Factors affecting individual health and the case of aquatic ecosystems

Extrinsic (e.g. predation, parasitism, microclimate, contaminant) and intrinsic (e.g. individual 107 age, body condition, life history stage or genetic) factors may affect organism stress through 108 both direct and indirect pathways. In aquatic ecosystems, limited oxygen (Wajsbrov et al., 109 1991), extreme osmotic (Laiz-Carrión et al., 2005; Tandler et al., 1995) or temperature 110 (Heather et al., 2018) conditions have direct impacts on organism's immune function (Cuesta 111 et al., 2005) and oxidative status (Madeira et al., 2016, 2013). Metal contaminants 112 accumulated in an organism also have direct long-term consequences on anti-oxidant 113 depletion (Médale et al., 2005; Saera-Vila et al., 2009), enhancing oxidative stress (Benhamed 114

et al., 2016; Mourente et al., 2002; Soares et al., 2008) and impairing immune function 115 (Cerezuela et al., 2016; Guardiola et al., 2016, 2015). Similarly, individual age or experience 116 also directly affects the way organisms cope with given stressors; older individuals often 117 coping more efficiently with stressors owing to past experience, until late life senescence 118 (Barcellos et al. 2012, Navaratnarajah and Jackson 2013). For instance, age- and species-119 specific changes in bio-accumulation or sensitivity to specific contaminants are thought to 120 121 evolve: metal contaminants such as mercury (Hg) shows bio-accumulative patterns over life within and across species (Ourgaud et al., 2018), whereas others are detoxified or diluted 122 through time (Lee et al., 1998). Indeed some contaminant are known to biodiminished during 123 ontogenetic development (e.g. passed within the eggs, Johnson et al., 2020) or due to a shift in 124 the diet while growing, therefore lowering exposition (Lee et al., 1998). Moreover 125 contaminants may induce reduced stress response often underlying difficulties to deal with 126 127 these stressors (Hontela and Fortin, 1995; Tan et al., 2009). In the other hand some essential trace elements (e.g. Cu, Zn) are necessary in small amount to insure vital physiological 128 129 functions (Wood et al., 2012) but have detrimental effect in high concentrations (Minghetti et al., 2008; Oliva et al., 2007; Shuhaimi-Othman et al., 2015). Such relationships urge the use 130 131 of several markers of individual health and insist on the necessity to consider both positive 132 and negative relationship between contaminant and stress response.

133

Besides these direct effects, extrinsic and intrinsic factors also influence individual stress response through indirect pathways. In particular, environment-driven changes (e.g. temperature, light, salinity, pH) affecting lower trophic levels of the food web, can also affect higher trophic levels indirectly by modifying prey availability (Chassot et al., 2010). Assessing variation in individual diets may therefore present a convenient tool for tracking indirect effects on individual energy stored, growth and stress response. Organic matter inputs

are central to aquatic ecosystem functioning, affecting primary production (phytoplankton vs. 140 141 aquatic angiosperms in eutrophic vs. oligotrophic conditions), as well as higher trophic levels (i.e. affecting the nature of reserves fish assimilate as well as their growth capacities; Escalas 142 et al., 2015; Isnard et al., 2015; Le Fur et al., 2019). For instance, whereas fish grow faster at 143 intermediate salinities, salinity also has complex interactions with temperature and oxygen 144 concentrations, and the consequences on food intake and food conversion are species-145 dependant (Bœuf and Payan, 2001). Increasing salinity also limits metal contaminant 146 accessibility in animal (Ifremer, 2009; Lee et al., 1998) and bioaccumulation most probably 147 due to osmo-regulation modulation (Lee et al., 1998; Wright, 1995) but biodisponibility in the 148 149 water and sediments tend to improve (Zhao et al., 2013).

150

151 Study aim and choice of a relevant fish model

As a case study, we focused on the ecophysiology of gilthead seabream (Sparus 152 aurata) in the Gulf of Lions (Mediterranean Sea), which migrate yearly between coastal 153 154 lagoons (for growth and development) and the open sea (for reproduction) during its juvenile and adult stages (Mercier et al., 2012). The physiology of gilthead seabreams has been well 155 studied and physiological markers of chronic stress, including high cortisol and oxidative 156 157 stress and decreased immune function, have been identified (Salati et al., 2016; Tort et al., 1998; Vera et al., 2014). As a ectothermic species, Seabream growth and movement are 158 affected by environmental changes in temperature (Heather et al., 2018; Madeira et al., 2016; 159 Mignucci et al., 2021). In addition, the environmental conditions of coastal lagoons are known 160 to vary over short spatial scales, including in dissolved oxygen concentrations, eutrophication, 161 162 and contaminants known to cause oxidative stress and depress fish immune system (Benhamed et al., 2016; Cerezuela et al., 2016). 163

Here, we focused on young seabreams (0-3 years old) captured in ten coastal lagoons, 164 165 at an age where lagoon conditions, including diet, have marked long-term consequences on individual growth (Escalas et al., 2015; Isnard et al., 2015) and stress response (Médale et al., 166 2005; Mourente et al., 2000). The early life period of fish is a critical period of growth and 167 development, associated with high mortality rates (Cushing and Horwood, 1994). To assess 168 the direct and indirect effects of environmental factors on individual physiology, stress 169 170 response and health, including synergic, agonistic or additive effects (e.g. cocktail effects between contaminants; Celander 2011), we used a Partial Least Squares-Path Modelling 171 (PLS-PM) approach. PLS-PM evaluates interactions through a combination of multiple linear 172 regressions and principal component Analyses (Wold, 1980; Esposito Vinzi et al., 2010; 173 Sanchez, 2013). 174

We expected contaminant loads to directly impair individual health. We further 175 expected abiotic and biotic environmental features (eutrophication, organic matter, salinity 176 and depth), as well as individual age to affect seabream growth and body condition (i.e. lipids 177 178 and protein content), through changes in their diet. Indeed eutrophication and organic matter inputs are known to boost the entire trophic web as well as seabreams growth rate (Brehmer et 179 al., 2013; Escalas et al., 2015; Isnard et al., 2015). Salinity and organic matter contents play a 180 role in contaminant biodisponibility (Richards et al., 2001; Schwartz et al., 2004; Zhao et al., 181 2013) and since Gilt-head seabreams is a necto-benthic species it feeds and uses bottom 182 habitats. Moreover the depth of the lagoon may play a role in hidden from predator, especially 183 early in life. Finally organisms allow energy to growth to reach maturity and reproduce as 184 early as possible but also to avoid predation. Ultimately this structural investment (hereafter 185 186 estimated by growth rate, protein content and structural lipids known as phospholipids) will necessarily require energy for maintenance (structural cost, Stearns, 1989). When resources 187 exceed these cost, energy can be stored under the form of reserve lipid (Triacylglycerol, 188

Anedda et al., 2013; Queiros et al., 2019). Since fish condition is known to interact with contaminant bio-accumulation, and therefore indirectly with stress response, the use of a PLS-PM approach (see Fig. 1 for the general construct) and comparison of seven alternative models, allowed for a comprehensive study into how environmental conditions, individual condition and environmental contaminant affected fish health in coastal lagoons.

194

195 METHODS

196 Environmental descriptors

We monitored ten coastal lagoons (Arnel, Ayrolle, Gruissan, Ingril, Leucate, Or, Prévost, 197 Bages-Sigean, Thau, and Vic; latitude: from 48° 51' 18" N to 43° 34' 50" N; longitude from 3° 198 0' 21" E to 4° 2' 16" E; see Fig. 2) in the Gulf of Lions (France, NW Mediterranean Sea, Fig. 199 2). Lagoon morphological features included surface area (km²), maximum depth (m), and 200 hydrobiological characteristics of the water column (collected 1 m below the surface), 201 including salinity, dissolved inorganic nitrogen (DIN, µM), total nitrogen (µM), total 202 phosphorus (μ M), phosphate (μ M), chlorophyll *a* (μ g.L⁻¹), dissolved oxygen concentration 203 $(O_2, mg.L^{-1})$, and Turbidity (NTU), as integrative descriptors of the environment at a lagoon 204 205 scale (ESM1). Those data were acquired via the OBSLAG network and the Water Framework Directive (WFD) (Bouchoucha et al., 2019; Derolez et al., 2017). Data were collected 206 monthly from 2014-17 (time period corresponding to the age in years of the sampled 207 seabreams, see below) during summer (June, July, August), from a single station in smaller 208 lagoons (Arnel, Ayrolle, Gruissan, Ingril, Prévost, and Vic), two stations in intermediate sized 209 lagoons (Bages-Sigean, Leucate and Or), and six stations in the largest lagoon (Thau). 210

Linking lagoon morphometric and hydrobiological features to relevant biological
functions or mechanisms is not trivial and previous studies have addressed this issue (see

Derolez et al., 2020 and references therein). Following recommendations of Derolez and 213 214 collaborators, we partitioned lagoons over a multidimensional gradient running a principal component analysis over all environmental variables. The three first principal components of 215 this analysis explained 52.7%, 17.8% and 12.6% of the total variance (83.1% in total; ESM1), 216 highlighting three fundamental descriptors. Principal component 1 (PC1) loaded strongly and 217 positively on total nitrogen, total phosphorus, chlorophyll a, phosphates and turbidity, 218 representing increasing organic matter inputs and primary productivity with increasing values 219 of PC1 and highlighting eutrophication (Derolez et al., 2020). PC2 loaded strongly and 220 positively on lagoon morphology (increasing values of PC2 reflecting increasing lagoon 221 222 surface and depth), whereas increasing values of PC3 reflected increasing concentrations of dissolved inorganic nitrogen and decreasing salinity in the water column (ESM1). Each 223 lagoon was then characterized by its centroid value on each of these three axes that have been 224 225 considered in the following analyses to describe the environment.

226

227 Fish sampling

Seabream were sampled from the same ten coastal lagoons in autumn 2017. Fish were 228 sampled just before their annual winter migration from the lagoons. Individuals were fished 229 using traditional Mediterranean fishing gear ("capetchades") from October 10th to 30th 2017. 230 This fishing method uses a thin mesh to capture and sample live fish. The device was 231 232 deployed in the evening; fish entered during the night and were collected early the next morning. We captured 100 individuals, focusing on the smallest fish, to insure all were young 233 individuals and incidentally males (as seabreams are proteandrous hermaphrodites). Fish were 234 kept in oxygenated lagoon-water tanks until sacrificed by immersion in ice-cooled lagoon-235 water tanks and a blood sample collected within 1 hour of capture. We collected 0.5 mL of 236 blood from the caudal vein using a 25G heparinised syringe. After centrifugation (3500g for 237

10 min) plasma was separated and frozen in a dry shipper containing liquid nitrogen (~-238 239 190°C, ©CX100 Worthington Industries) and then kept frozen at -80°C in the lab until analysed. Individuals were weighed to the nearest $\pm 1g$ and their total length measured to the 240 nearest \pm 1mm using a metal ruler (Table 1). Fish were kept on crushed ice for transport to the 241 lab, then frozen at -20°C until analyses. Four weeks later, all individuals were unfrozen and 242 we collected the scales over the entire right flank of the fish flank for stress hormone 243 244 (cortisol) content measures, and 20 to 30 scales below the pectoral fin on the opposite flank to confirm individual age from scale growth annuli. Three pieces of dorsal epaxial muscle (white 245 muscle) were saved and kept frozen at -80°C for contaminant, lipid and protein content 246 247 analyses. Muscle samples were collected using ceramic tools and rinsed with ethanol 99% and 248 double-deionized water (MilliQ) between each individual to avoid any contamination. Muscle weight was measured before and after lyophilisation to account for individual hydration level. 249 250 Measurements on muscle (contaminant, lipid, FA) are reported relative to muscle dry weight.

251

252 Individual age and growth rate

Individual age (in years) was determined by counting annual increments on five to seven scales immersed in aqueous solution using transmitted light, as routinely done in seabream (for details see Mahé et al., 2009). We estimated individual growth rates by dividing individual total length by age. As growth rate drastically decreases after the first year and to account for variation in growth rate between ages, we standardized growth rate by age class (1, 2, 3). Descriptive statistics of total length, mass, age estimated by scale reading, and growth rate centred by age are summarized for each lagoon Table 1.

260

261 **Body composition**

262 Lipid content

Lipids were extracted from fish muscle following Folch et al. (1957) and lipid classes 263 separated and quantified using an Iatroscan MK-VI (Iatron Laboratories, Inc.) following 264 Parrish (1999) (for details; see Sardenne et al., 2019). Lipids were broadly classified into 265 reserve (e.g. fat stores) or structural (e.g. membrane phospholipids) lipids. Reserve lipids are 266 constituted of triacylglycerols (TAG), diacylglycerols (DAG as precursors of TAG) and free 267 fatty acids (FFA). In contrast, structural lipids include phospholipids (PL) and sterols (ST). 268 Finally, acetone-mobile polar lipids (AMPL), and alcohols (ALC) are eluted at the same time 269 but present in relative low proportions (Tocher, 2003). PL represented the predominant lipid 270 (ca. 87%; 45 to 98% of total lipids, TL) in our samples, whereas TAG concentration showed 271 important inter-individual variation (1 to 51 % of TL). FFA, ALC, DAG and AMPL were 272 very low (<0.05%) or undetected, while ST were always detected at low level (< 2% of TL, 273 see ESM2). Low FFA proportion indicated that lipid integrity was conserved during sample 274 275 storage (Parrish, 1999).

276

277 Muscle Protein content and choice of body composition indices

Protein content was assessed using a piece of epaxial muscle after lyophilization and grinding 278 (ball mill, MM400, Retsch GmbH, Germany). Proteins were extracted from 10 mg of dry 279 muscle powder by immersion and incubation in a 1.5-mL solution of a 10% SDS (Sigma 280 Aldrich), 1.5% Protease inhibitor Cocktail (cOmplete, Sigma Aldrich, France), miliO water 281 solution (lysis solution adapted from Campus et al. 2010). Cycles of successive 15-min 282 ultrasonic baths (300 Ultrasonik, Ney Company, USA) and 3-min vortex were repeated four 283 times to allow total dissolution of the sample. Extracts were clarified by a 10-min 284 centrifugation at 3000g at 4°C (Ericsson and Nistér, 2011). Muscle protein content in 25 µL 285 of the collected fraction was quantified using a Bicinchoninic Acid method (BCA, Pierce, 286

Thermo Fisher Scientific, France). Intra- and inter-plate protein variations (based on the same
sample repeated over plates) were 7.62% and 16.7%, respectively.

As body composition indices, we used (1) TAG and TAG/PL as indicators of individual endogenous energetic reserves (fish reserve); and (2) PL, protein content, and individual growth rate as indicators of individual investment into structure components (fish structure) (Anedda et al., 2013; Queiros et al., 2019).

293

294 Trophic markers and Fatty Acid ratio indicators

295 Fatty acid identification

296 The polar fraction of lipids, mainly PL, plays an important role in cell structure and shows a high degree of plasticity in response to changes in abiotic environmental features. The neutral 297 fraction plays a role as a resource compartment, destined either for oxidation to provide 298 energy (ATP) or for incorporation into PL. Predominantly made out of TAG and wax esters, 299 its composition is less influenced by changes in abiotic features and therefore better reflects 300 individual diet (reviewed in Dalsgaard et al., 2003). Therefore, only the neutral fraction was 301 considered in this study. Neutral and polar lipids were separated by column chromatography 302 303 on silica gel and methylated using 12% BF₃-MeOH (see ESM3). Fatty acid methyl esters (FAMEs) were analysed in an Agilent 6890 gas chromatograph equipped with an on-column 304 injector and a flame-ionization detector, with hydrogen as a carrier gas. They were identified 305 by their retention times with reference to those of a standard 37-component fatty acid methyl 306 esters (FAME) mix and designated following the formula C:XovY where C is the number of 307 carbon atoms, X is the number of double bonds and Y is the position of the first double bond 308 counted from the CH terminal. 309

310

In aquatic food webs, FAs are conservatively transferred along trophic levels (Lovern, 1935). 312 Determining the FA composition of fish tissues has been widely used to track individual diet 313 314 composition (Dalsgaard et al., 2003; Gao et al., 2006), as well as its quality in terms of essential FAs (Izquierdo 2005, Benedito-Palos et al. 2008, ESM3). As the present study aimed 315 at identifying broad-scale diet profiles, we selected nine generic aquatic trophic indicators 316 (either sum or ratio of specific FA) relevant to aquatic systems (Blanchet-Aurigny et al., 317 2015), as well as Arachidonic acid (ARA), Docosahexaenoic acid (DHA) and 318 eicosapentaenoic acid (EPA) contributions (%) to the neutral lipid fraction (Koussoroplis et 319 320 al., 2011) (Table 2).

321

322 Metal contaminant load in fish muscle

Muscle samples were treated with trace mineral grade nitric acid (69% Suprapure, Merck) and 323 33% H₂O₂ (Suprapure, Merck) in Teflon reaction tubes, heated in a microwave digestion 324 system (UltraClave-Microwave Milestone®) for 20 min at 220 °C, and then diluted to 10 mL 325 with double-deionized water (MilliQ). Total mercury (Hg) content was measured using an 326 atomic absorption spectrometer AMA254 Advanced Mercury Analyzer (Leco) (Bouchoucha 327 et al., 2018), without pre-treating or pre-concentrating the samples (wavelength = 253.65 nm, 328 detection limit (DL) = 0.003 ug.g^{-1}). Samples were further analysed using inductively coupled 329 plasma optical emission spectrometry (ICP-OES, ICAP 6500 Duo, Thermo) to quantify levels 330 of aluminium (Al), arsenic (As), boron (B), cadmium (Cd), chrome (Cr), copper (Cu), nickel 331 (Ni), lead (Pb), rubidium (Rb), Selenium (Se) and zinc (Zn) (García-Navarro et al., 2017). 332 The DL for the analysed elements was 0.001 $\mu g.g^{-1}$. Each sample was read in duplicate and 333 averaged. Uncertainty remained between 4.12 and 6.43% (Al: 5.38%, As: 4.23%, B: 4.82%, 334 Cd: 4.56, Cr: 4.30%, Cu: 4.12%, Hg: 5.26%, Ni: 4.83%, Pb: 6.14%, Rb: 5.36%, Se: 6.43% 335

and Zn: 5.02%). Based on UNE-EN ISO reference 11885, multi-element calibration standards
(SCP Science, in 4% nitric acid) were assembled with different concentrations of inorganic
elements.

339

340 Individual stress response

341 Stress hormone: scale cortisol content

342 We assessed chronic individual stress response levels by measuring scale cortisol levels (adapted from Carbajal et al. 2018), the principal glucocorticoid in teleost. Measuring scale 343 content, after removing external cortisol from mucus, allows assessing glucocorticoid 344 secretion retrospectively over an extended period of time (Aerts et al., 2015; Sadoul and 345 346 Geffroy, 2019). The scales of an entire flank of each individual were collected, washed and vortexed three times (2.5 min; 96% isopropanol) to remove external sources of cortisol. 347 Residual solvent traces were evaporated under nitrogen flux, samples frozen at -80°C and 348 349 lyophilised for 12 hours before grinding to a powder (ball mill, MM400, Retsch GmbH, Germany). Cortisol was extracted from ~20 mg of dry scale powder by incubation in 1.5mL 350 of methanol (MeOH) on a 30°C rocking shaker for 18 hr. After centrifugation (9500g, 10 351 352 min), the supernatant was evaporated under nitrogen flux and reconstituted with 0.2 mL of EIA buffer provided by the cortisol assay kit (Neogen® Corporation Europe, Ayr, UK). 353 Cortisol concentrations were determined in 50 µL of extracted cortisol samples using 354 competitive EIA kits (Neogen® Corporation Europe, Ayr, UK). Samples were run in 355 duplicates and averaged. Intra-plate variation (each sample run in duplicates) was 3.36% and 356 357 inter-plate variation (2 control samples were run in duplicate over all plates) was 5.67%.

358

359 *Fish oxidative status*

We assessed global measures of oxidative damages and antioxidant defences in plasma using 360 361 the d-ROMs test and OXY Adsorbent tests (Diacron International©, Grosseto, Italy). Oxidative damage was assessed in 8 μ L of plasma by measuring hydroperoxides, the main 362 compounds contributing to plasma oxidant activity, expressed as mg H₂O₂ equivalent/dL 363 (Beauvieux et al., 2022). Oxidative defences were evaluated in 5 μ L of 1:100 diluted plasma 364 by measuring the ability of plasma to buffer massive oxidation through hydroperoxide acid, 365 expressed in µmol HCl/mL (Beauvieux et al., 2022). All sample measurements were run in 366 duplicate and averaged. Intra-plate variations (each sample run in duplicates) were 2.40% and 367 7.51% for ROMs and OXY, respectively. Inter-plate variations (2 control samples were run in 368 369 duplicate over all plates) were 10.02% and 7.92% for ROMs and OXY, respectively.

370

371 Immune functions: Natural haemolytic complement and lysozyme activities

Fish immune systems are precursors and quite primitive relative to that of most vertebrates, 372 mostly relying on non-specific/innate/natural immunity. The complement and lysozyme 373 systems are important mechanisms of the immune system for recognizing and eliminating 374 non-self-substances in fish (Sitjà-Bobadilla et al., 2003; Tort et al., 1998). Plasma activity of 375 the complement pathway was assayed against rabbit red blood cells collected in Alsever's 376 solution (Sigma Aldrich, Lyon, France). SRBC were washed in phenol red-free Hank's buffer 377 (HBSS) containing Mg²⁺ and EGTA and re-suspended at 3% (v/v) in HBSS. 100 μ L test 378 379 serum was serially diluted in HBSS solution to obtain four final serum concentrations ranging from 10% to 0.078% in a round-bottomed 96-well plate. An equal volume (100 µL) of SRBC 380 was added to each well. Following incubation for 90 min at room temperature, samples were 381 centrifuged at 400 g for 5 min at 4°C to remove un-lysed erythrocytes. The relative 382 haemoglobin content of supernatants was assessed by measuring its optical density at 550 nm 383 (©Tecan Infinite 200 microplate reader). The values of maximum (100%) haemolysis were 384

obtained by adding 100 mL of distilled water to 100 mL samples of SRBC and values of minimum (spontaneous) haemolysis obtained from SRBC without serum. The power of haemolysis (%) was estimated as the slope of its lysis curve (Díaz-Rosales et al., 2006). Intra-(each sample run in duplicates) and inter-plate (2 control samples were run in duplicate over all plates) variations were 9.06% and 14.71%, respectively.

Lysozyme activity assays were performed by a turbidimetric method using the lysis of 390 *Micrococcus luteus* by plasma samples to determine their enzymatic activity. Serum lysozyme 391 activity was measured according to methods described by Ellis (1990) and previously applied 392 to gilthead seabream (Tort et al., 1998). Briefly, 25 µL of individual serum was mixed with a 393 75 µL Micrococcus lysodiekticus (Sigma Aldrich, Lyon, France) suspension at 75 µg.mL⁻¹ in 394 0.1 M phosphate citrate buffer, pH 5.8. After rapid mixing, turbidity changes were measured 395 every 60s over 30 min at 415 nm, 28°C (©Tecan Infinite 200 microplate reader). The 396 dilutions of chicken egg white lysozyme (Sigma Aldrich) served as a standard. Lysozyme 397 activity was estimated by comparing individual dynamic turbidity decrease over time 398 399 compared to that of the standard's and is expressed as Unit/mL serum. Samples were run in duplicate and averaged. Intra-plate (each sample run in duplicates) and inter-plate (2 control 400 samples were run in duplicate over all plates) variations were 15.7% and 5.68%, respectively. 401

402

403 Statistics

All statistical analyses were run in the R v3.5.1 (R Development Core Team 2008). We gathered all information required for the present study for 96 out of the 100 sampled fish (age 0: N=42, age 1: N=27 & age 2: N=48). For four individuals, one or two assays were lacking due to limited blood material and were removed from the analysis. Altogether we thus considered 96 individuals * 39 variables = 3744 data points (see ESM1 & 6 for more details).

We investigated the multi-scale causal pathways by which extrinsic (environmental 410 and anthropogenic) and intrinsic factors affected seabream ecophysiology using Partial Least 411 412 Squares Path Modelling (PLS-PM) (R package plspm, Sanchez 2013). PLS-PM allows studying complex multivariate relationships among groups of manifest (observed) and latent 413 (construct) variables (MVs and LVs respectively), testing for direct and indirect effects. These 414 415 effects are necessarily unidirectional (Sanchez, 2013; Vinzi et al., 2010) and can be used to draw causal inferences in the path diagram (more detailed framework presented in ESM4). 416 LVs are hypothetical constructs that are not measured directly but can be assumed to reflect 417 418 their underlying MVs. The general construct of this study is detailed in Fig. 1.

Using a PLS-PM approach, we defined extrinsic and intrinsic constructs (LVs), built 419 in a formative way (for detail see ESM4), characterizing relevant environmental and 420 physiological pressures affecting seabreams in coastal lagoons. We then explored how 421 422 individual stress response was related to those morphological and hydrobiological, 423 anthropogenic (contaminant), dietary and intrinsic constructs using an iterative algorithm (Trinchera and Russolillo, 2010), and comparing seven alternative models (Table 3). The 7 424 constructs (LVs) always included: (1) lagoon features (morphological and hydrobiological); 425 and fish (2) diet, (3) contaminant loads, (4) endogenous reserves, (5) structure components, 426 (6) age, and (7) stress response. Each of these latent variables was the combination of one or 427 more MVs (such as lagoon eutrophication, lagoon depth, fish cortisol levels, etc.). Expected 428 directional relationships were based on empirical and theoretical research to draw a 429 comprehensive picture of the complex ecosystem interactions likely affecting seabream/fish 430 431 stress response in coastal lagoons, and alternatives were assessed for seven different causal models (see ESM5). We tested for potential effects of lagoon features on fish age (Model 2), 432 and tested if the LV fish age was best predicted by structural size, age in years or both (Model 433

434 1, 3 & 4). Models 1, 5 & 6, differed in terms of directional relationships (e.g. fish reserve
435 affected fish structure, the opposite, or no relation was specified between those LVs; Table 3).
436 We computed a final model including all of the interactions retained in previous models
437 (Model 7). For each model, we computed and compared the overall variance explained by the
438 path model (goodness of fit, GoF) (Table 3, ESM5). The model displaying the highest GoF
439 was retained and discussed in the manuscript.

440 First, we consider the model from an integrative perspective. We consider which resources individuals might extract from the environment and how they might distribute those 441 to different functions, ultimately asking how individuals deal with environmental variations in 442 443 terms of stress management. We therefore describe causal relationships likely linking a given LV to other independent LVs in our model. Second, we discuss the indirect effects affecting 444 individual stress response. The overall path model (PLS-PM; Fig. 3) represents: (1) the 445 manifest models including all correlations between MVs (e.g. lagoon eutrophication, depth, 446 area, dissolved inorganic nitrogen and salinity) and their respective LVs (e.g. lagoon 447 448 features); and (2) the structural model in which all causal relationships between our LVs (lagoon features, fish age, fish diet, fish reserve, fish structure, fish contaminant load, fish 449 stress) are determined. 450

451

452 **RESULTS**

453 Model Comparisons

The seven models displayed similar global explanatory performances (GoF) ranging from 27.3% to 32.6% (Table 3). Regardless of the model considered the relationships between connected latent variables (LVs) that were significant remained identical in terms of sign (positive or negative; ESM5) and only differed slightly in terms of strengths. Model 7
displayed the highest GoF (32.6 %) and therefore was selected for further investigation.

459

460 Global model assessment: overall variance explained by the path model 7

461 *Manifest model variance*

462 The variance of the manifest variables (MVs) reflected by the latent variables (LVs) is given by the block communality coefficients (Table 5). For instance, the LV "lagoon features" 463 reflected 44% of the variance in its three MVs, *i.e.* the three principal components of the PCA 464 representing eutrophication, lagoon morphology (depth and area), and dissolved inorganic 465 nitrogen (DIN) and salinity variables (Table 5). The LV "fish age" included the age and size 466 of fish and accounted for 94% of the variation in age and size. Other LVs shared a lesser 467 proportion of variance with their underlying MVs, ranging from 12 to 36%. Table 5 468 summarizes all explained variance from LVs and Fig. 3 describes how they related to the 469 underlying (measured) MVs. First, the LV "lagoon features" was positively related to 470 eutrophication (r = 0.91), lagoon depth and area (r = 0.60) and the third PCA axis 471 encompassing increasing DIN and decreasing salinity (r = 0.36). Second, the LV "fish diet" 472 473 assessed by measurement and composition of free fatty acid, was negatively related to dinoflagellates 2 (r = -0.51), diatoms (r = -0.49), terrigenous inputs (r = -0.33), bacteria (r = -474 and arachidonic acid (ARA) (r = -0.34) and 0.18), detrital sources (r = -0.65), 475 eicosapentaenoic acid (EPA) (r = -0.69), while positively related to carnivorous (r = 0.78) and 476 copepods (r = 0.15) inputs. DHA (r = 0.10), dinoflagellates 1 (r = -0.06), and non-methylene 477 interrupted (NMI) (r = -0.06) contributions to the LV "fish diet" remained small (r \leq 0.10). 478 Third, the LV "fish reserve" corresponded to an increase in TAG (r = 0.35) and TAG/PL ratio 479 (r = 0.56), both being proxies of fish body condition. Fourth, the LV "fish structure" 480

positively correlated to growth rate (r = 0.52) and negatively correlated (r = -0.90) with the 481 amount of phospholipid present in the neutral FA stored. Muscle protein concentration only 482 marginally contributed to the LV "Fish structure" (r = 0.07). Fifth, the LV "fish contaminant" 483 load positively correlated to metal concentrations of Al (r = 0.40), As (r = 0.14), B (r = 0.50), 484 Cu (r = 0.57), Rb (r = 0.35), Zn (r = 0.46), Hg (r = 0.20) and negatively correlated to 485 concentrations of Se (r = -0.51) and Ni (r = -0.11). Cd, Cr, and Pb contributions were 486 marginal ($r \le 0.10$). Finally, the LV "Fish stress" highly and positively correlated to cortisol 487 (r = 0.71), oxidative damage (ROM, r = 0.67), complement (r = 0.29), and antioxidant 488 defences (OXY r = 0.17) levels. In contrast, it was negatively related to lysozyme activity (r =489 - 0.24) (see Fig. 3 and ESM6). 490

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492 Structural model variance

Within the structural model, the amount of variance of a given LV (*e.g.* fish contaminant load) that is explained by all other LVs (lagoon features, etc.) in the path model is presented Table 5 (\mathbb{R}^2 values). The amount of explained variance in LVs "fish structure" (55%), "fish contaminant" (63%) and "fish stress" (54%) was high. In contrast, the amount of explained variation for LV "fish reserve" was relatively low (16%) and the amount of variation in LV "fish diet" explained by other LVs (fish age and lagoon features) was intermediate (28%).

Direct effects: Starting from the top of our constructs, individual diet composition was affected by individual age and environmental lagoon features encountered during the summer (Fig. 3; Table 6). The greater lagoon eutrophication, depth, area and dissolved inorganic nitrogen content, the more fish diet shifted from planktivorous, terrigenous, detrital, and EPA sources to carnivorous ones. The same relationship was found as fish age increased. Consequently, when fish shifted to a more carnivorous diet, structural lipid concentrations decreased but growth increased (LV "fish structure"). Such diets had a positive effect on fish energy stores (reserve variable), reflected by increased TAG and TAG/PL ratio. Age showed opposite trends on fish reserve and fish structure: increasing age was associated with increasing structural lipids (although limited growth) but decreasing energy stores. We found no substantial effect of the LV "lagoon feature" on fish reserves or fish structure. Notably, fish structure was strongly affected by fish reserve with higher reserves leading to higher growth.

Fish age was also negatively associated with muscle contaminant concentrations, 512 whereas fish reserve, structure and diet had non-significant effects on contaminant 513 514 concentrations. Most importantly, fish contaminant concentrations had the highest direct positive (+0.38) impact on fish stress response. Increasing lagoon eutrophication, depth, area, 515 dissolved inorganic nitrogen and decreasing salinity (LV "lagoon feature") were associated 516 with increased fish stress response (+0.25), whereas stress response largely decreased with 517 increasing fish age (-0.41). Finally, the direct effects of individual diet and energy stores (LV 518 519 "Fish reserve") on individual stress response was non-significant (Fig. 3, Table 6).

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Indirect effects: The PLS-PM framework further allowed assessing indirect effects by 521 multiplying coefficients along the path. Focusing on total effects affecting fish stress 522 response, positive direct effects of lagoon features (+0.25) on individual stress response was 523 524 counteracted by negative indirect effects through muscle contaminant loads (total effect of lagoon features on contaminant load * total effect of contaminant on stress response = -0.77 * 525 0.38 = -0.29). The later was slightly balanced out by indirect effect through fish diet (0.57 * 526 0.20 = 0.11), fish reserve (0.22 * 0.10 = 0.02) and fish structure (0.28 * 0.17 = 0.05) (Figure 4 527 and ESM7). As a result, the overall effect of lagoon features on fish stress response was low 528 (+0.11). In contrast, direct and indirect age effects on fish stress response were cumulative 529

530 leading to the highest negative coefficient (-0.61): increasing age was associated with 531 decreasing stress loads through direct and indirect effects. The indirect effects of fish diet, 532 reserve and structure on fish stress response were marginal (<0.06). A summary of direct and 533 indirect effects is given Figure 4 (for a full description of the other LVs relationships see 534 ESM7).

535

536 **DISCUSSION**

Using a multifactorial PLS-PM analysis, we identified key direct and indirect contributions of 537 538 extrinsic (lagoon features, fish diet, fish contaminant loads) and intrinsic (fish reserves, fish structure components, fish age) factors affecting fish stress response in coastal lagoons. 539 Understanding how individuals deal with multiple ecological pressures using stress response 540 as a proxy is a first step to better understand the constraints that apply to population dynamics 541 542 (Brosset et al., 2021; McKenzie et al., 2016; Saraux et al., 2019). Although a larger sample size might have improved the overall statistical power, the PLS path model explained 33% of 543 the total variance of the 46 variables accounted in this study, reflecting ten Mediterranean 544 lagoon systems, a substantial amount considering that the work was performed at the scale of 545 multiple ecosystems. More importantly, 54% of the variance of the LV "fish stress" was 546 explained by the dependent LVs "lagoon features", "fish age", "fish diet", "fish reserve", 547 "fish structure", and "fish contaminant" included in the model. 548

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551 Extrinsic and intrinsic factors affecting fish metal contaminant loads

Lagoon features had an important negative effect on fish contaminant loads. When the LV "lagoon features" increased, eutrophication (inorganic matter content and primary

production) increased, while salinity decreased. This is consistent with the fact that both the 554 555 toxicity of metal contaminant to fish and their bioavailability in water are known to vary with water physicochemical properties, including pH, alkalinity, suspended solids, organic 556 compound content, and hardness (Di Giulio and Meyer, 2008). Increasing organic matter 557 concentration reduces metal accumulation to aquatic organisms by reducing the amount of 558 free cationic metal by chelation (Richards et al., 2001; Schwartz et al., 2004). In rainbow 559 560 trout, increased dissolved organic matter concentration has been associated to increased survival when faced with lethal Pb and Cu metal concentrations allowing reduced binding to 561 gills (Richards et al., 2001; Schwartz et al., 2004). In contrast, while increasing salinity has 562 563 been shown to reduce metal contaminant accessibility in bivalves and some teleost fish (Somero et al. 1977; Lee et al. 1998; Ifremer 2009), its biodisponibility in the water and 564 sediments tend to increase (Zhao et al., 2013). Our results suggest that for seabream the 565 566 opposite effect of eutrophication and salinity on contaminant concentration in the fish could be due either to differences in gill osmoregulation processes in seabream compared to other 567 species tested, or non-exclusively, to confounding effects of decreasing salinity and increasing 568 organic matter concentration affecting contaminant loads. Indeed, the strong negative effect (-569 0.65) of the LV "lagoon feature" on fish contaminant load could be due to a stronger effect of 570 increasing organic matter than decreasing salinity (higher loading and weight of 571 Eutrophication of Lagoon features compared to Salinity; ESM6). 572

We also found that Gilthead seabream displayed a general pattern of metal contaminant bio-diminution with age. The PLS-PM showed that contaminant load decreased markedly with age (direct effect of fish age on contaminant load; -0.55, Fig. 3, Table 6). When focusing on individual metal concentrations and their respective relationships with the total length of individual (see ESM8), we found that aluminium (Al), boron (B), copper (Cu), nickel (Ni), rubidium (Rb), selenium (Se) and zinc (Zn) concentrations were all negatively

related to total length. This suggests that dilution of those metals occurred with age-related 579 580 changes in diet shift and size (Lee et al., 1998) and is consistent with similar relationships that have been described in trophic webs (cephalopods, coelenterates, crustaceans, echinoderm, 581 fish, and gastropods) for cadmium (Cd) and lead (Pb) in freshwater and marine ecosystems 582 (Asante et al., 2008). In contrast, arsenic (As) and mercury (Hg) were the only two elements 583 to bioaccumulate and/or biomagnify in seabream (reflected by positive correlations of their 584 585 concentration with size). Cd, chromium (Cr) and Pb showed no relationship with total length (ESM8). This absence of relationships suggests that these metals were neither biomagnified 586 nor biodiminished during ontogenetic development or through the food web. Although these 587 588 relationships deserve to be considered with cautious, indeed the relatively low overlap of age classes between each lagoon do not allow to exhaustively disentangle age and environmental 589 effects, similar bio-accumulation and dilution are common in marine organisms and food 590 591 webs, including seabreams (Chouvelon et al., 2014; Cresson et al., 2015; Harmelin-Vivien et al., 2012). 592

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594 Synergetic effects and consequences for individual body composition and growth

Lagoon features had an overall (direct + indirect) positive effect on fish reserve (+0.22) and 595 596 fish structure (+0.28). Increasing relative eutrophication (without decreasing dissolved O2 concentration ESM1B) was positively linked to fish growth rates and the quantity of 597 endogenous reserves individuals accumulated during the summer. This concurs with previous 598 findings that seabream growth and body condition between and within lagoons generally 599 increase with increasing organic matter inputs (Brehmer et al., 2013; Escalas et al., 2015; 600 Isnard et al., 2015). These two effects were largely captured indirectly in our model (direct 601 effects non-significant; coefficients < 0.05; Table 6) by dietary shifts from planktivorous, 602 terrigenous, detritic, and EPA-based diets to carnivorous diets when eutrophication increases. 603

Further, lagoon features and fish age appear to act synergistically on fish diet: as individuals 604 605 age and grow, they progressively switch to bigger prey, which at juvenile stage lead to the developing of a more carnivorous diet toward molluscs and fish larvae (Escalas et al., 2021; 606 Rosecchi, 1987). 607

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Integrating multiple factors in our understanding of fish stress response and health

611 We found that fish age (age in years and size) displayed the strongest direct effect on stress response (-0.41). Older individual are thought to have greater experience since they 612 have logically faced and survived several challenges as they age. As they age, individuals that 613 are exposed to stressors might also develop mitigating strategies allowing them to deal with a 614 variety of challenges and acclimate resulting in decreasing stress responses (Moreira et al., 615 616 2019). Moreover, fish that survive longer may be of better intrinsic quality (biased selection), leading to individuals displaying lower maintenance costs in the sampled population (Birnie-617 618 Gauvin et al., 2017; Wilson and Nussey, 2010). These non-exclusive hypotheses may explain 619 the lower stress response of older (max. 3 years old) individuals.

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Interestingly, the effect of several lagoon features on stress response was independent of the 621 622 other studied LVs, *i.e.* not captured as indirect effects but as direct effects (+0.25). In other words, increasing eutrophication, lagoon morphology (depth and surface), inorganic nitrogen, 623 and decreasing salinity directly increased individual stress response. This may be due to direct 624 effects on individual physiology, such as individual osmoregulation processes, that can be 625 energetically stressful when salinity decreases and eutrophication increases (Bodinier et al., 626 2010; Cuesta et al., 2005). An alternative may be that these effects are captured indirectly, 627 through other unmeasured lagoon ecosystem features known to increase stress and decrease 628

629 immune functions, for instance predation risk (Werner et al., 1983) or competition (Goldan et630 al., 2008; Salati et al., 2016), which deserve further investigation.

631

Fish structure (decreasing PL and increasing growth rate) had a non-significant effect (+0.18, 632 p=0.089) on fish stress response (increase in cortisol level, oxidative damages, anti-oxidant 633 defences and complement activity, and a decrease in lysozyme activity). Nonetheless, from a 634 635 life-history perspective at the organism level, it is assumed that the greater the investment in growth, the lower the investment in prevention or repair of molecular damage (Cichon, 1997; 636 Stearns, 1989). Pushing physiological machinery to grow faster may lead to metabolically 637 induced damage (Metcalfe and Alonso-Alvarez, 2010; Monaghan et al., 2009). Studies in 638 several taxa indicate that the trade-off underlying accelerated growth early in life comes at an 639 oxidative cost (Merry, 1995). Long-term consequences may be offset by subsequent impaired 640 641 performance, including decreased immunological competence, energy reserves and longevity (Inness and Metcalfe, 2008; Rollo, 2002), or reduced investment in protein maintenance 642 (Morgan et al., 2000). This is notably the case for animals having experienced a period of 643 slow growth followed by rapid compensation (compensatory growth), relative to those having 644 grown steadily throughout (Metcalfe and Monaghan, 2001). However, there are many 645 646 ecological advantages to attaining a large body size as fast as possible. Reaching maturation rapidly may be a successful strategy in terms of reproductive success or competition for food 647 resources (Lee et al., 2012; Scharf et al., 2000). Larger sizes may also be associated with 648 649 reduced risks to predation (Sogard, 1997). Such long-term benefits may outweigh the shortterm stress costs of investing in structural tissues and growth. 650

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Most importantly, our analysis shows that when considering several component of the costal lagoon ecosystem and physiological trait of the gilthead seabream, its stress response

was mostly affected by muscle inorganic contaminant content. Over the past decade, similar 654 655 causal relationships have been shown by studies on contaminant loads under controlled laboratory conditions. Exposure to heavy metals is usually associated with increased cortisol 656 and oxidative stress levels (Sevcikova et al., 2011) and lower immune function (Cerezuela et 657 al., 2016; Guardiola et al., 2016, 2015). In small amounts, metal elements such as As, Cu, Cr, 658 Ni, and Zn are essential for immune and oxidative defence functions, whereas they impair 659 these same functions in excess (Sevcikova et al., 2011). For instance, Zn or Cu deficiencies 660 lead to deficiencies in immune and antioxidant systems and to the development of life 661 threatening infections (Rink and Kirchner, 2000). In contrast, at high doses, those same 662 663 elements have inhibiting effects on the immune system (Rink and Kirchner, 2000) or increase oxidative stress (Noh and Koh, 2000; Valko et al., 2005). Because of their deleterious impacts 664 at low concentration on organism physiology, As, Cd, Hg and Pb are the most commonly 665 666 (often individually) studied metal contaminants. Here, we considered a wider array of contaminant, considering cocktail effects in interaction with other environmental features 667 rather than single contaminant effects. It is remarkable that individual contaminant loadings 668 on the LV "contaminant load" were relatively low for these widely studied metal contaminant 669 (As +0.14, Cd +0.02, Hg +0.20 and Pb +0.10). Rather, it appeared that interactions between B 670 (+0.50), Cu (+0.57), Rb (+0.35), Zn (+0.46), Al (+0.40) and a decrease in Se (-0.56) were 671 critical in explaining individual stress response in seabream. It is important to understand that 672 the physiological action of these two different groups of contaminant in causing oxidative 673 stress are fundamentally different (Valko et al., 2005). Metals without redox potential, such as 674 As, Cd, and Hg, cause oxidative stress by impairing antioxidant defences, especially those 675 involving thiol-containing antioxidants and enzymes (Stohs and Bagchi, 1995). In contrast, 676 redox active metals such as Fe, Cu and Al participate in the formation of reactive oxygen 677 species (e.g. Fe and Cu catalyse the formation of reactive hydroxyl radicals (•OH) by Fenton 678

reaction) (Gaetke and Chow, 2003; Liaquat et al., 2019). Elements such as dietary Se have
been shown to protect against lipid peroxidation and Cu (Kadiiska and Mason, 2002) and Cd
toxicity (Ulusu et al., 2003). The above highlights the complexity of interactions between
contaminant, and the importance of integrated approaches to evaluate cocktail effects.

Measuring cortisol in the scales of the fish, enabled us to assess a proxy of the total 683 amount of corticoid hormone secreted through the life of the individual (Goikoetxea et al., 684 2021; Harris, 2020; Sadoul and Geffroy, 2019). Cortisol level accumulated in the scales was 685 highly loaded on the LV "stress response", therefore its regulation tended to positively relate 686 to an increasing long term stress response. Circulating glucocorticoid hormone is commonly 687 688 used as a marker of individual stress response as they mediate a cascade of physiological (increased metabolism) and behavioural response to stress (Munck et al. 1984, Johnson et al. 689 1992) and has been widely employed to evaluate several stressor effects in fish, from disease 690 691 to chemical contaminants (Zimmer et al., 2011). Although under acute exposure its concentration tend to increase and lead to rapid mitigating response toward a stressor and 692 protection against toxic effects (Bury et al., 1998; Laflamme et al., 2000), long term intense 693 stress event may induce a decrease of glucocorticoid secretion and a decline in the 694 hypothalamic-pituitary-adrenal response to stress with subsequent consequences on 695 696 immunity and reproduction (Hontela et al., 1997; for a meta-analysis see Rohonczy et al., 2021). In the present study, the stress response of the fish was highly related to increasing 697 contaminant load. It suggests that long term exposure led to constant increase in cortisol 698 secretion without impairing their ability to mount an efficient HPA regulation. Although 699 critical stress responses do not seem to be reached, contrary to some acute responses observed 700 in experimental approaches (Hontela et al., 1997; Hontela and Fortin, 1995), it still rises 701 questions on the long term consequences of such maintained high stress response on future 702 reproduction and metabolism trade-offs. 703

Over the last decade, contaminant monitoring programs for coastal ecosystems were 704 705 developed for the Mediterranean Sea to provide scientific knowledge to assess ecosystem health and sustainability (Bonito et al., 2016; Naccari et al., 2015; Tomasello et al., 2012). 706 707 However, mainly due to European legislation, most studies on fish species have focused on As, Cd, Hg and Pb contamination or have compared contaminant loads for different species at 708 different locations (Cresson et al., 2016; Mille et al., 2018; see Chouvelon et al., 2017 for a 709 710 multi-population approach for tuna), although few studies have focused on the consequences on individual health in nature. Yet, as monitoring of organic contaminants such as POPs 711 (HAPs, PFOS, PFAS, DDT, Lindane...) become more common in marine environments, it 712 urges further research that would certainly bring new insights on the impact of changing 713 lagoon environment on fish health (Munaron et al., 2023). 714

715

716 *Conclusion*

Using the Gilthead seabream and Mediterranean lagoons as a model, our study emphasized 717 the importance of integrated approaches to identify direct and indirect contributions of 718 multiple environmental factors affecting individual stress response in coastal ecosystems. 719 Abiotic and biotic features of the environment have complex direct and indirect, synergetic or 720 721 antagonistic, interactions with individual intrinsic features. A proper evaluation of these 722 effects on stress response and health requires multi-factorial frameworks and may benefit from further inclusion into energetic models (such as the allostasis or reactive scope models) 723 724 aiming to understand the tipping-point at which physiological systems are pushed beyond their capacity to cope with stressors in the environment leading to pathology and disease 725 (McEwen & Wingfield 2003; Romero et al. 2009). The challenge ahead lies in extending such 726 727 approaches from the individual to population scale.

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737 ACKNOWLEDGEMENTS

This research was founded by Ifremer and Fisheries and Ocean Canada through the Galway 738 Statement on Atlantic Ocean Cooperation. The authors are grateful to local fishermen for fish 739 collection. We kindly thank the Ifremer LER-LR of Sète (France) and especially Martine 740 Fortuné, Gregory Messiaen, Annie Fiandrino, Vincent Ouisse as well as Béatrice Bec and 741 742 Elise Hatey from the University of Montpellier for their participation in the environmental 743 data collection over the years. We thank the Agence de l'Eau Rhône Méditerranée Corse for financing the OBSLAG and WFD monitoring networks of the lagoons between 2014 and 744 2017. We thank Marion Richard for her help and expertise on lagoon functioning. We thank 745 746 Bastien Sadoul for his help and suggestions in adapting the Cortisol assay for fish scales. We thank Blandine Brisset, Jean-Hervé Bourdeix, Eric Gasset and Gilbert Dutto for their 747 unwavering support and their incredible ability to find practical solutions for any technical 748 issue. We thank the sclerochronology center of Brest, Ifremer for the annuli-scale reading and 749 ageing of all fish. 750

751

752 AUTHORS CONTRIBUTION

QS, CS, VAV and JB designed the experiment and collected the data. QS, LM, AB and LL 753 754 developped the methods and did most of the laboratory analyses. DR measured metal contaminant concentrations in all seabream muscles. VD and DM managed the long term 755 environmental data collection (OBSLAG-WFD), provided environmental datasets, and 756 contributed to the interpretation of lagoons features. CQ and FP identified and quantified fatty 757 acid. CWM provided logistical and financial support in the realisation of the study. QS did the 758 statistical analyses and wrote the manuscript. CS, VAV, CWM, AB and JB commented on the 759 paper. All authors agreed on the final version. 760

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762 COMPLIANCE WITH ETHICAL STANDARDS

763 All experiments were approved by the national ethic comitee (APAFIS#8945-201

764 7021612469374 v4 & APAFIS#10759-2017072512353814 v3) and realized in compliance

765 with French and European Law.

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- 1228 FIGURE CAPTIONS
- 1229 Figure 1. Path diagram of the structural model.

Figure 2. Map of the sampling area. Each sampling site is identified by its local Frenchname.

Figure 3. Partial Least Square (PLS) path model showing the strength and direction of the relationships among Latent variables. Red represents negative direct effects while

1234 positive relationships are represented in green. The thickness of the arrows is 1235 proportional to the strength of the effect and dashed arrows represent non-significant

relationships ($p_{value} > 0.05$). Direct effects are shown as standardized path coefficients. Loadings of each Measured Variable (MV) in relation to its Latent Variable (LV) are

- 1238 shown for each block in its respective frame.
- 1239 Figure 4. Summary of direct and indirect effects of independent latent variables (LVs)
- 1240 on fish stress. The total effect of an independent LV on its dependent LV is the sum of
- 1241 all indirect effects and the direct effect.

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1243 TABLE CAPTIONS

Table 1. Fish sampling summary table. Descriptive statistics are given for each lagoon in terms of total length, mass, age estimated by scale reading, and growth rate centred by age. We gathered all information required for the present study for 96 of the 100 sampled fish. For four individuals, one or two assays were lacking due to limited blood material and they were removed from the analysis.

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Table 2. Summary of the different trophic and diet quality markers selected for the analysis and associated references. Based on Blanchet-Aurigny et al. 2015. For an exhaustive review, see Pernet (2016).

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Table 3. Models comparisons. GOF Goodness-of-fit statistics, calculated as the 1254 geometric mean of the average block communality and the average R^2 value, thus 1255 accounting for the quality of both the measurement and the structural models. The 1256 1257 quality of the manifest model is evaluated for each latent variable (LV) by block communality, which is calculated as the average of all squared correlations between the 1258 given LV and its underlying manifest variables (MVs). Block communality thus 1259 measures how much of the variance is common between a LV and its MVs (Vinzi et al., 1260 1261 2010). Finally, the fit of the structural model is assessed for each endogenous LV (\mathbb{R}^2) , which highlights the amount of variance in the endogenous LV explained by its 1262 independent LVs (Vinzi et al., 2010). 1263

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1265 Table 4. Summary of the different parameters measured in fish, the tissue considered,

- 1266 their unit, their role as biomarkers and the latent variable they refer to.
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Table 5. Summary of the model 7; Exogenous refers to Latent Variable in the model for which no incoming effect are tested (no arrows pointed at them; lagoon features and age).

1271 Table 6. Summary of the latent model. Effect size of the relationships between each LVs

1272 within the model 7 are given (estimates) as well as their error (standard deviation) and t-

1273 values. *Pvalues are considered when $\alpha < 0.05$.

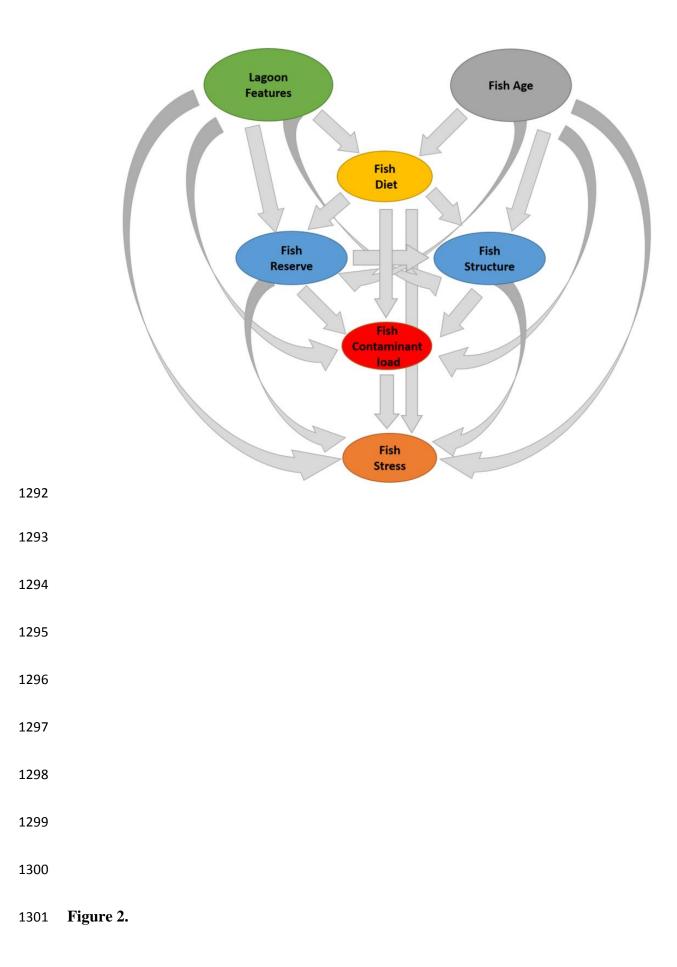
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1290	FIGURES
1291	Figure 1.



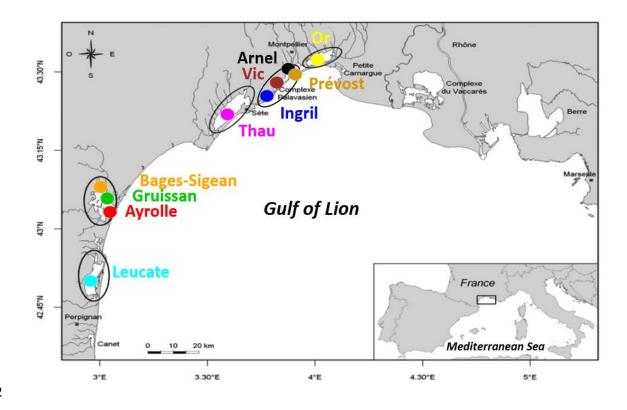
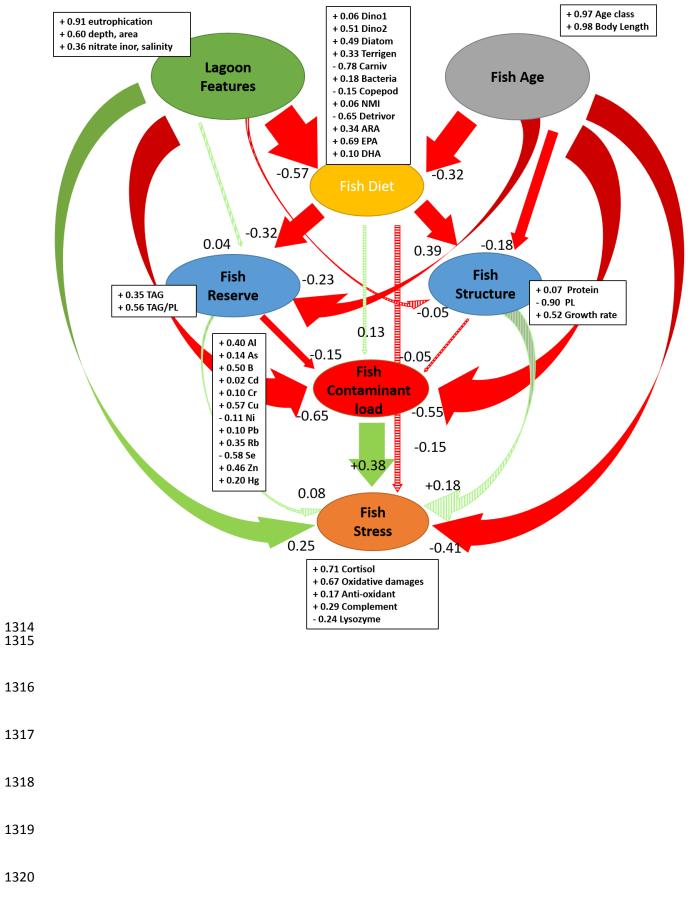
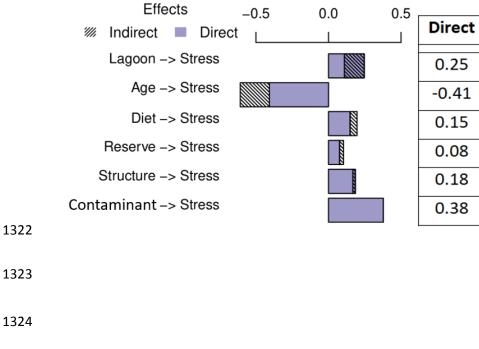


Figure 3.



1321 Figure 4.



Indirect

-0.14

-0.20

0.05

0.03

-0.02

0.00

Total

0.11

-0.61

0.20

0.10

0.17

0.38

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- 1335 TABLES
- 1336 **Table 1.**

		Total Le	ength	Ma	ISS			Grow	th rate
		(mm	1)	(g	;)	Ag	e	(centred	by age)
Lagoon	n	Mean \pm sd	min-max	Mean \pm sd	min-max	Mean \pm sd	min-max	Mean \pm sd	min - max
Arnel	10	190.6 ± 4.7	185 - 200	97 ± 6	89 - 108	0.0 ± 0.0	0-0	0.17 ± 0.18	-0.07 - 0.5
Ayrolle	10	281.3 ± 17.9	254 - 300	331 ± 65	238 - 430	1.6 ± 0.5	1-2	-0.01 ± 0.47	-0.77 - 0.6
Gruissan	9	299.1 ± 57.2	158 - 324	429 ± 153	52 - 516	1.8 ± 0.7	0-2	1.09 ± 1.00	-1.32 - 1.69
Ingril	10	274.5 ± 28.2	232 - 313	305 ± 104	160 - 436	1.7 ± 0.5	1-2	-0.27 ± 1.11	-1.8 - 1.15
Leucate	9	277.0 ± 9.4	265 - 295	293 ± 32	248 - 354	1.8 ± 0.4	1-2	-0.37 ± 0.43	-1.05 - 0.32
Or	9	201.6 ± 8.3	191 - 214	139 ± 22	116 - 188	0.0 ± 0.0	0-0	1.01 ± 0.38	0.45 - 1.75
Prévost	10	261.4 ± 48.9	132 - 305	282 ± 90	174 - 408	1.5 ± 0.5	1-2	-0.35 ± 0.98	-2.01 - 0.85
Bages- Sigean	9	280.1 ± 9.6	265 - 295	324 ± 32	274 - 361	1.0 ± 0.0	1-1	1.04 ± 0.55	0.19 - 1.83
Thau	10	264.7 ± 11.1	249 - 286	267 ± 37	218 - 333	2.0 ± 0.0	2-2	-1.09 ± 0.58	-1.89 - 0.02
Vic	10	190.8 ± 30.5	106 - 209	108 ± 9	97 - 125	0.0 ± 0.0	0-0	0.56 ± 0.28	0.09 - 0.92
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1346	Tab	le 2.							

Marker	Source	Reference			
Bacteria: iso15:0 + ant15:0 + 15:0 + iso16:0 + iso 17:0 + 17:0	Bacteria	(Volkman et al., 1980)			
Dino1: 16:100 7/16:0	Dinoflagellates	(St John and Lund, 1996)			
Dino2: 20:5\u03c03/22:6\u03c03	Diatoms vs dinoflagellates	(Budge and Parrish, 1998)			
Diatom : 16:1\u07/20:5\u03	Diatoms	(Pernet, 2016)			
Detrital: SFA/PUFA	Phytoplankton vs detrital	(Biandolino et al., 2008; Maazouzi et al., 2007; Pommier et al., 2010; Prato et al., 2012)			
Copepods: $20:1\omega9 + 22:1\omega11$	Copepods	(Budge et al., 2002; Kattner et al., 2012)			
Terrigenous: $18.2\omega6 + 18.3\omega3$	Terrestrial vascular plants, Green microalgae / Terrigenous	(Budge and Parrish, 1998; Dalsgaard et al., 2003; Kelly and Scheibling, 2012)			
non-methylene interrupted (NMI) 20:2i + 20:2j + 22:2i + 22:2j	Bivalves and gastropods	(Budge et al., 2007; Joseph, 1982)			
Carnivorous: 18:1@9/18:1@7	Carnivorous	(Auel et al., 2002; El-Sabaawi et al., 2009; Graeve et al., 1997)			
Essential FA	Role	Reference			
Docosahexaenoic acid (DHA):	Essential FFA: neuronal	(Arts and Kohler, 2009; Bell			
22:6ω3	development and maintenance	and Sargent, 2003, 1996; Bell et al., 1995; Koussoroplis et al.,			
Arachidonic acid (ARA): 20:4ω6	Essential FFA: precursor of eicosanoid hormone	2011; Koven et al., 2003;			
Eicosapentaenoic acid (EPA): 20:5ω3	Essential FFA: precursor of eicosanoid hormone	Schmitz and Ecker, 2008; Van Anholt, 2004)			

Table 3.

						Lagoon	D	iet	Res	erve	Stru	cture	Conta	minant	Str	ess	Ag	ae
Model	Model specificity	Lvs	Mvs	Direct Paths	GOF		Block		Block		Block		Block		Block		Block	
1	Basic	7	38	19	0.2908									0.638				
7	Basic + effect of Lagoon on Age	7	38	21	0.2733									0.587				
2	Basic with Size instead Age	7	38	20	0.283									0.613				
3	Basic with Size + Age	7	39	19	0.203									0.631				
4	Basic + effect of Structure on Reserve	7	38	20	0.2972									0.634				
5	Basic + efffect of Reserve on Structure	7	38	20	0.3074												1.000	
6	Basic + efffect of Reserve on Structure with Size + Age	1	39	20	0.3262	0.439	0.193	0.283	0.212	0.160	0.359	0.550	0.124	0.626	0.225	0.538	0.943	farmer i
	1356 1357																	
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13/1 Table 4.	1371	Table 4.
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Parameters	Tissue	Unit	Role	Latent
				Variable
Cortisol	Scales	μg.g ⁻¹	Systemic stress hormone	Fish Stress
Oxidative damages	Plasma	$\begin{array}{c} mg H_2O_2 \\ equivalent.dL^{-1} \end{array}$	Oxidative damages accumulated	Fish Stress
Antioxidant	Plasma	µmol HCl.mL ⁻¹	Oxidative defences available	Fish Stress
Complement	Plasma	%lysis	Efficiency of the innate immune function (complement pathway)	Fish Stress
Lysozyme	Plasma	U.mL ⁻¹	Efficiency of the innate immune function (Lysozyme enzymatic efficiency)	Fish Stress
Inorganic contaminants	Muscle	μg.g ⁻¹ dry muscle	Contamination load by each inorganic contaminants	Fish contaminant
Protein	Muscle	mg.g ⁻¹ dry muscle	Body condition index – quantity of proteins allocated in structure	Fish Structure
Lipids	Muscle	mg.g ⁻¹ dry muscle	Body condition index - Types and quantity of lipids stored in the muscle	Fish Reserve & Fish Structure
Fatty acids	Muscle	% total - ratios	Trophic markers - Types and ratios of fatty acids present in the muscle	Fish Diet
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1383 Table 5.

LATENT VARIABLE	Туре	Manifest variables	R ²	Communality
Lagoon features	Exogenous	↑ Eutrophication, depth & area and dissolved inorganic nitrogen ↓Salinity		0.439
Fish Age	Exogenous	↑ Age ↑ Fish total length		0.943
Fish Diet	Endogenous	 ↑ Carnivorous, Copepods ↓ Dinoflagellates, Diatoms 1 & 2, Terrigenous, Bacteria NMI, Detrital, ARA, EPA, DHA 	0.283	0.193
Fish Reserve	Endogenous	↑ TAG and TAG/PL	0.160	0.212
Fish Structure	Endogenous	\uparrow Growth rate \downarrow PL	0.550	0.359
Fish contaminant load	Endogenous	 ↑ Al, As, B, Cr, Cu, Rb, Zn, Hg ↓ Se, Ni r≤0.10 Cd, Cr, Pb 	0.626	0.124
Fish Stress	Endogenous	↑Cortisol, oxidative stress damages, anti-oxidant, complement activity ↓Lysozyme activity	0.538	0.225
Goodness of fit		0.3262		
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Table 6.

	Estimate	Std. error	t value	P value
\$ Fish Diet				
Intercept	0.000	0.088	0.000	1
Lagoon features	0.571	0.096	5.960	4.48e-8*
Fish Age	0.321	0.096	3.650	1.15e-03*
\$ Fish Reserve				
Intercept	0.000	0.096	0.000	1
Lagoon features	0.037	0.123	0.302	0.763
Fish Age	-0.231	0.110	-2.100	0.039*
Fish Diet	0.318	0.113	2.820	0.006*
\$ Fish Structure				
Intercept	0.000	0.070	0.000	1
Lagoon features	-0.048	0.090	-0.329	0.596
Fish Age	-0.183	0.083	-4.370	0.030 *
Fish Diet	0.391	0.087	3.680	1.84e-05 *
Reserve	0.499	0.077	6.500	4.28e-09
\$ Fish contaminant load				
Intercept	0.000	0.064	0.000	1
Lagoon features	0.650	0.083	-7.840	8.56e-12*
Fish Age	-0.555	0.078	-7.090	2.94e-10*
Fish Diet	-0.132	0.088	-1.510	0.135
Fish Reserve	-0.149	0.085	-1.750	0.084
Fish Structure	-0.047	0.096	-0.485	0.629
\$ Fish Stress				
Intercept	0.000	0.072	0.000	1
Lagoon features	0.246	0.120	2.050	0.043*
Fish Age	-0.408	0.109	-3.730	3.3e-4*
Fish Diet	0.149	0.099	1.500	0.138
Fish Reserve	0.076	0.097	0.790	0.432
Fish Structure	0.185	0.107	1.720	0.089*

Fish contaminant load	0.378	0.118	3.210	0.002*
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