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1	Dietary antioxidants, food deprivation and growth affect differently oxidative
2	status of blood and brain in juvenile European seabass (Dicentrarchus labrax)
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22	Running title: Early life and oxidative stress

### 23 ABSTRACT

24 Compensatory growth may increase molecular oxidative damage, which may be 25 mitigated through the intake of dietary antioxidants. However, dietary antioxidants may 26 also reduce concentration of antioxidant enzymes, which have a key role in regulating 27 the oxidative status. Here we investigated whether feeding on a diet rich in antioxidants 28 (vitamin E) enables juvenile European seabass (Dicentrarchus labrax) to catch up after 29 a period of food deprivation with negligible effects on the oxidative stress to blood and 30 brain as compared to fish feeding on a normal diet (i.e., not enriched in antioxidants). 31 The results show that a higher intake in antioxidants favoured compensatory growth, but 32 this came at a cost in terms of increased oxidative damage. Increased intake of 33 antioxidants also resulted in changes in the activity concentration of enzymatic 34 antioxidant defences and increased protein oxidative damage in both brain and blood. In 35 addition, food deprivation caused increased protein oxidative damage in brain. Our 36 findings show that the beneficial effects of dietary antioxidants on growth may be offset by hidden detrimental effects and that different early life events affect different 37 38 components of oxidative status of a given tissue.

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*Keywords:* Antioxidants, Aquaculture, Early life, Fish, Food restriction, Oxidative
stress

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## 44 **1. Introduction**

45 Positive selection of individuals capable of reaching an adult size quickly should be 46 expected to sift out slowly growing individuals. Achieving an adult size as soon as 47 possible should carry potential benefits to the individual, such as earlier time to sexual 48 maturity or reduced predation risk (Dmitriew, 2011). Conversely to this prediction, 49 animals do not grow at the maximum rate and much variation in growth strategies 50 persists in both wild and laboratory animals, even when trophic resources are abundant 51 (Blanckenhorn, 2000). The reason for this might lie with the costs of growing fast 52 (Metcalfe and Monaghan, 2001). For example, rapid growth can increase metabolic 53 demands (Criscuolo et al., 2008; Careau et al., 2013) and daily energy expenditure 54 (Careau et al., 2013), or reduce body performance and lifespan (Lee et al., 2016). An 55 additional mechanism through which growth may relate to self-maintenance is that 56 increased cellular activity due to fast growth increases the production of reactive 57 species, such as free radicals (Costantini, 2014; Smith et al., 2016). These unstable 58 molecules, if not counteracted by antioxidants, oxidise proteins, lipids and nucleic acids, 59 causing damage, and hence a state of oxidative stress (Halliwell and Gutteridge, 2015). 60 Recent correlative, experimental and meta-analytical work has suggested that oxidative 61 stress might play a key role as a constraint on, and cost of, growth (Alonso-Alvarez et 62 al., 2007; Nussey et al., 2009; Smith et al., 2016).

Oxidative costs of a rapid growth are particularly relevant when individuals accelerate growth to achieve the body size that is supposed to be normal for their life stage (known also as catch-up growth; Metcalfe and Monaghan, 2001). For example, abrupt reductions in food availability - common in wild populations - might lead young to slow down growth. When food availability becomes adequate again to meet the nutritional requirements of a growing individual, juveniles will compensate by accelerating their growth. Although there are benefits in doing so, an acceleration of growth may also carry costs, such as a reduction in motor skills, sexual ornamentation, cognitive abilities or longevity (Metcalfe and Monaghan, 2001; Lee et al., 2016), but also in increased molecular oxidative damage (Smith et al., 2016).

73 Organisms may mitigate oxidative stress through an increased intake of 74 antioxidants from diet. Animals normally ingest a cocktail of antioxidants and distribute 75 them differently across tissues, depending on the chemical properties (lipophilic vs. hydrophilic) of the antioxidant and the specific requirements of a certain tissue (Surai, 76 77 2002; Halliwell and Gutteridge, 2015). Dietary antioxidants may be particularly 78 important for growing individuals as the endogenous antioxidant machinery takes time 79 to become fully mature (Surai, 2002; Costantini, 2014). In many vertebrate species, 80 there is a greater reliance upon non-enzymatic antioxidants at earlier stages of 81 development, such as those acquired from diet or from the mother (Surai, 2002; 82 Costantini, 2014). It would therefore be expected that if an individual catches up while 83 feeding on a diet rich in antioxidants, it might mitigate the oxidative costs of a fast 84 growth. This simple prediction, however, does not consider that increases in the intake 85 of a given antioxidant may also result in no net reduction in oxidative damage because 86 any potential beneficial effects might be offset by compensatory reductions in other 87 antioxidants. For example, long-term intake of a dietary antioxidant (vitamin C) in 88 laboratory mice reduced expression of several genes linked to free radical scavenging, 89 with no net effect on various oxidative damage metrics (Selman et al., 2006). A recent 90 meta-analysis also showed that the effects of antioxidant supplementation on growth are 91 highly variable (from positive to negative), implying that effects of dietary antioxidants 92 on oxidative status might be difficult to predict (Smith et al., 2016).

93

Another aspect that has received little attention is the impact that accelerated 94 growth has on brain oxidative status and protein oxidative damage (Smith et al., 2016). 95 This is surprising given that acceleration of growth may reduce cognitive abilities 96 (Metcalfe and Monaghan, 2001), brain is a tissue particularly sensitive to oxidative 97 stress (Barja, 2004) and protein turnover is altered during fast growth (Samuels and 98 Baracos, 1995).

99 Compensatory growth is particularly common in fish at almost every stage of 100 their life cycles because of their indeterminate growth patterns (Ali et al., 2003). 101 Despite the expansion of the field of ecological oxidative stress, relatively little is 102 known about the role of oxidative stress in mediating key life-history stages of fish 103 (Birnie-Gauvin et al., 2017), such as those fish go through during growth and 104 development.

105 In this study, we tested experimentally whether feeding on a diet rich in 106 antioxidants enables young European seabass (Dicentrarchus labrax) to catch up after a 107 period of food deprivation with negligible effects on the oxidative stress as compared to 108 fish feeding on a normal diet (i.e., not enriched in antioxidants). To this end, we have 109 measured metrics of oxidative damage and of antioxidant enzyme activity (i.e., proxy of 110 enzyme concentration) in both blood and brain in juvenile fish that underwent a food 111 deprivation period and then a refeeding protocol either with or without an enriched diet.

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113 2. Materials and methods

114 2.1. Ethical note All experiments complied with the Guidelines of the European Union Council and were
approved by the Ethical Committee of the Tuscia University for the use of live animals
(D.R. n. 677/16 and D.R. 644/17).

118 At the end of the food deprivation period, the loss of body mass in fish that were 119 food-deprived was 11.4 and 12.6 % for the two experimental groups (mean  $\pm$  se: prior 120 food deprivation,  $59.7 \pm 5.1$  g and  $59.1 \pm 5.1$  g; end of food deprivation,  $52.8 \pm 5.2$  g 121 and 51.6  $\pm$  5.2 g), respectively. Hence, it was below the threshold of 15 % 122 recommended by the UK Home Office and the Canadian Council for Animal Care. The 123 mortality rates calculated using sample sizes at the beginning of Phase 1 were similar 124 among groups at the end of the experiment: 12.5 % (number of fish dead were 3), 25.0 125 % (number of fish dead were 7), 14.3 % (number of fish dead were 4) and 14.3 % 126 (number of fish dead were 4) for no food deprivation/standard diet, no food 127 deprivation/enriched diet, food deprivation/standard diet, and food deprivation/enriched 128 diet, respectively.

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### 130 2.2. Animals and husbandry

131 Juvenile European seabass (n = 112, one year of age) were obtained from a commercial hatchery (Cooperativa Orbetello Pesca Lagunare, Santa Liberata, Orbetello, Italy) 132 133 located on the coast at about 40 km from the facility where the study took place. The 134 fish were transported to the facility and randomly allocated into four fiberglass tanks 135 (80 x 80 x 403 cm, all equipped with aeration system) in groups of similar size (26, 29, 136 29 and 28 individuals). Before being released into the tanks, each fish was massed and 137 measured (mean  $\pm$  SD, body mass: 64.5  $\pm$  22.7 g; body length 18.9  $\pm$  2.0 cm; body 138 width  $4.2 \pm 0.5$  cm, see below for method details). In each tank, fish were maintained in 139 a volume of 1,290 liters on an open circuit system with flow of 30 liters/minute 140 (replicating the conditions of the supplier hatchery) on natural light dark cycle and 141 illumination, as ample windows were present in the building. Being an open circuit 142 meant that each tank was subject to natural seasonal variations of chemical/physical 143 parameters: pH ranged between 8.0 and 8.2, temperature varied from 15°C in January to 144 23°C in August, while salinity ranged between 35 and 38 PSU; ammonia and nitrite 145 were below detectable levels, while nitrate was about 5 mg/L. The European seabass is 146 an euryhaline coastal species known to tolerate (at both juvenile and adult stage) a large 147 range of chemical-physical water parameters, including salinity and temperature (in the 148 order of 5-28°C) extremes (e.g. Dulger et al., 2012).

149 Tanks were cleaned daily to avoid accumulation of food remains. Fish were fed 150 by hand *ad libitum* once per day on the same commercial standard feed (company name 151 undisclosed) used by the supplier for 12 acclimation days. In order to make sure the 152 ration was ad libitum, we provided food until fish did not eat pellets anymore. The 153 residual pellets that deposited at the bottom of the tank were removed within around 60 154 minutes in order to minimise any potential confounding effects arising from very 155 bold/dominant individuals monopolising the access to the food. The feed composition was as follows: crude proteins 48.0 %; crude fats and oils 22.0 %; crude fibre 3.0 %; 156 157 crude ash 6.0 %; vitamin E 150 mg/Kg.

- 158
- 159 2.3. Experimental procedure and sampling

Phase 1, food deprivation: after the acclimation period, the fish of two tanks started a food deprivation protocol that lasted 81 days, while the control fish in the other two tanks continued to be fed daily on the same commercial feed as previously (Fig. 1). All fish were massed and measured on the day Phase 1 was started; the sample size was 108because four fish died before the start of Phase 1.

165 Phase 2, re-feeding and enriched diet: as the food deprivation protocol stopped, 166 the pool of animals was re-fed as follows: two groups were fed daily on the standard 167 feed, and the other two groups were fed by hand ad libitum once per day on a feed 168 enriched with vitamin E (feed composition: crude proteins 48.0 %; crude fats and oils 169 20.0 %; crude fibre 2.0 %; crude ash 6.0 %; vitamin E 600 mg/Kg). In this way any 170 effect of the enriched diet on growth and oxidative status could be tested on both the 171 food restricted and the control fish (Fig. 1). This phase lasted until the final sampling, 172 which occurred 180 days later.

173 In order to quantify the growth, fish were caught with hand nets by two 174 operators, massed (Sartorius TE 612) and measured with a caliper for length and body 175 width (excluding dorsal and ventral fins) by a third operator once at the end of phase 1 176 and three times during phase 2 (Fig. 1). At the final sampling, the animals were bled after a mild anesthesia and were straightaway euthanized with an overdose of 2-177 178 phenoxiethanol (8 ml/l). Soon after euthanisation, the brain was collected from each 179 individual. Tubes containing blood samples were spun to separate plasma from red 180 blood cells. All samples were stored at -80°C and later transported in dry ice to the 181 laboratories of the University of Antwerp for the biochemical analyses.

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183 2.4. Laboratory analyses

We randomly selected samples of blood and brain of 10 individuals per experimental group. Whole brain was homogenised in cold PBS (supplemented with 20% (v/v) of glycerol and with 0.2 mM of phenylmethylsulfonyl fluoride as an inhibitor of proteases)

187 using a pestle. Samples were then sonicated for 10 minutes and then centrifuged for 10 188 minutes at 15,000 rpm. The supernatant was taken and split into different tubes, which 189 were stored at -80°C for later analyses. Haemolysates were centrifuged to separate cell 190 membranes from the supernatant, which were used for the analyses straightaway. The d-191 ROMs assay (reactive oxygen metabolites; Diacron International, Grosseto, Italy) was 192 used to measure plasma oxidative damage metabolites (e.g., organic hydroperoxides) 193 that are generated early in the oxidative cascade. Values were expressed as  $mM H_2O_2$ 194 equivalents. Protein carbonyls (marker of oxidative damage to proteins) were measured 195 in plasma and brain homogenates using the Protein Carbonyl Colorimetric assay 196 (Cayman Chemical Company, Ann Arbor, MI, USA). The assay is based on the protocol of Levine et al. (1990). The concentration of protein carbonyls has been 197 198 expressed as nmoles/mg proteins. The Ransod assay (RANDOX Laboratories, Crumlin, 199 UK) was used to quantify the activity of the enzyme superoxide dismutase (SOD) in 200 both red blood cells and brain. The activity of SOD has been expressed as Units/mg 201 proteins. The Ransel assay (RANDOX Laboratories, Crumlin, UK) was used to 202 quantify the activity of the enzyme glutathione peroxidase (GPX) in both red blood cells 203 and brain. The activity of GPX has been expressed as Units/mg proteins. The assay is 204 based on the original method of Paglia and Valentine (1967). Protein concentration in 205 the haemolysate or homogenate was measured using the Bradford protein assay (Bio-206 Rad Laboratories, Hercules, CA, USA).

207

208 2.5. Statistical analyses

Statistical analyses were performed using SPSS Version 23. General Linear Modelswere used to test the effect of our experimental manipulation on growth. Three main

211 factors and their interactions were included in the model. The main factors were pre-212 refeeding group (fed or food deprived), refeeding group (food enriched or not enriched 213 with antioxidants) and sampling day (three levels). As a response variable (body size 214 index), we used the first principal component (PC1, 95.8 % of the total variance) 215 extracted from a principal component analysis (PCA) on body mass, body length and 216 body width. The loadings (expressed as Pearson correlation coefficients) of these three 217 body measurements on the PC1 were as follows: 0.984 for body mass, 0.984 for body 218 length and 0.969 for body width. Analyses were repeated including body length or body 219 size (PC1 of a PCA on body mass and body width) as response variables, respectively. 220 General Linear Models were also used to test the effect of our experimental 221 manipulation on each metric of oxidative status. Each model included two main factors 222 (pre-refeeding group and refeeding group) and their interaction. To meet normality of 223 residuals and homogeneity of variances, plasma reactive oxygen metabolites and brain 224 superoxide dismutase were both square-root transformed, while brain protein carbonyls 225 were log-transformed. Transformations were chosen according to the fitting of the 226 models using values of Akaike Information Criterion as a reference. Post-hoc tests were 227 used to explore further any significant interaction across logical sets of means. When 228 non-significant, interactions were removed from the models. Outcomes of full and 229 reduced models coincided when there was not backward elimination because the 230 interaction/s was/were significant.

231

## 232 **3. Results**

The difference in body size between control and treated fish depended on whether fish were re-fed on a normal or enriched diet (three-way interaction: p = 0.047). To

235 investigate this further, the effect of the experimental manipulation on body size was 236 tested separately for fish re-fed on a normal or enriched diet, using similar linear models 237 as explained above. Fish re-fed on a normal diet did not catch-up, resulting in a smaller 238 body size than those fish that were not food-deprived (Fig. 2, Table 1). Conversely, fish 239 re-fed on an enriched diet were capable to catch-up, resulting in a body size as that of 240 fish that were not food-deprived (Fig. 2, Table 1). Similar results were obtained if body 241 size was extracted from a PCA calculated on body mass and body width (three-way 242 interaction: p = 0.012). As for body length, fish that were food-deprived were shorter 243 than fish that were not food-deprived at the end of the food deprivation period, while 244 they were of similar body length at the end of the experiment (Fig. 3; Table 1). The 245 effect of food deprivation was not influenced by the quality of diet given after the food 246 deprivation period (Table 1).

The difference in plasma reactive oxygen metabolites between control and treated fish depended on whether fish were re-fed on a normal or enriched diet. Fish that were fed on an enriched diet after food deprivation had significantly higher plasma reactive oxygen metabolites than all other experimental groups (Fig. 4, Table 1).

Plasma protein carbonyls were higher in fish that were either fed continuously or
re-fed on an enriched diet than those either fed continuously or re-fed on a normal diet
(Fig. 5, Table 1). Irrespective of the quality of re-feeding diet, brain protein carbonyls
were higher in fish that were food-deprived than in fish that were not (Fig. 6, Table 1).

The activity of glutathione peroxidase in red blood cells was higher in fish that were either fed continuously or re-fed on an enriched diet than those either fed continuously or re-fed on a normal diet (Fig. 5, Table 1). The activity of brain glutathione peroxidase was unaffected by the experimental manipulation in both phases(Table 1).

The activity of superoxide dismutase in both red blood cells and brain was lower in fish that were either fed continuously or re-fed on an enriched diet than those either fed continuously or re-fed on a normal diet (Fig. 5, Table 1).

263

# 264 **4. Discussion**

265 The results of this study show that a diet rich in vitamin E may favour compensatory 266 growth after a prolonged period of fasting, but this comes at a cost in terms of increased 267 oxidative damage. As expected, our food deprivation regime slowed down growth of 268 fish. A re-feeding period of six months enabled fish to compensate through a faster 269 growth, but the compensatory growth was mainly evident for mass/condition, while 270 there was not compensation for body length (proxy of skeletal size). There are at least 271 two possible explanations. Previous studies on fish have shown that (i) growth 272 compensation may be limited in some species to restoring lost tissues/energy reserves, 273 with no acceleration of skeletal growth (Alvarez and Nicieza, 2005), or (ii) growth 274 compensation of body length may take longer than body mass/condition to be evident 275 (e.g., Johnsson and Bohlin, 2006).

The compensatory growth occurred only in those fish that were re-fed with a diet enriched with vitamin E, highlighting the importance of this nutrient for fish growth. Vitamin E may interact with many metabolic and cellular pathways important for growth, such as lipid transport by lipoproteins, vitamin C synthesis, function of erythrocytes, immunity or metabolism of other nutrients (Hamre, 2011). Previous work on juvenile gilthead seabreams (*Sparus aurata*) also showed that increased intake of

282 vitamin E can reduce skeletal anomalies, favouring growth (Izquierdo et al., 2013). 283 Vitamin E promotes membrane repair and such repair ability is particularly important 284 when cells are being exposed to an oxidant challenge (Howard et al., 2011; Labazi et 285 al., 2015). Moreover, vitamin E may reduce haemolysis of red blood cells through a 286 chain-breaking activity, i.e., able to scavenge free radical by hydrogen donation (Niki et 287 al., 1988; Niki, 2014). Although vitamin E has antioxidant properties, those fish that 288 made a compensatory growth had higher levels of plasma reactive oxygen metabolites 289 than those fish that did not make a compensatory growth. High metabolic activity 290 typical of compensatory growth might have caused a high production of oxygen 291 metabolites. Conversely, protein oxidative damage was not increased in fish that made a 292 compensatory growth. A reason for this might lie with a higher protection of proteins 293 against oxidation. Reactive oxygen metabolites as measured in our work come from 294 early oxidation of biomolecules, and their accumulation is prevented by the activity of 295 the glutathione peroxidase (Halliwell and Gutteridge, 2015). On the other hand, protein 296 carbonylation is mostly irreversible damage, thus avoidance of carbonylation might 297 have been prioritised and traded-off against other kinds of damage.

298 The higher production of reactive oxygen metabolites in those fish that made a 299 compensatory growth might have also been favoured by the impact that vitamin E 300 intake had on antioxidant enzymes. Previous work on fish showed that vitamin E may 301 reduce the expression of superoxide dismutase gene and increase the expression of the 302 glutathione peroxidase gene, respectively (Izquierdo et al., 2013). These results on 303 expression of enzymatic antioxidant genes are in agreement with our results on the 304 activity of superoxide dismutase and glutathione peroxidase. Superoxide dismutase 305 catalyses the dismutation of the superoxide anion generated by mitochondria into

306 oxygen and hydrogen peroxide; glutathione peroxidase reduces hydrogen peroxide and 307 organic hydroperoxides into water and alcohols, respectively (Halliwell and Gutteridge, 308 2015). Hence, lower activity of superoxide dismutase might have resulted in reduced 309 protection against the peroxidative action of superoxide anion, which may increase 310 generation of organic hydroperoxides (Halliwell and Gutteridge, 2015). Upregulation of 311 glutathione peroxidase might have been needed to detoxify cells from organic 312 hydroperoxides, which may propagate the oxidative cascade (Halliwell and Gutteridge, 313 2015). Fish feeding on the diet enriched with vitamin E had actually higher protein 314 oxidative damage than those feeding on a normal diet.

315 Although vitamin E has antioxidant properties, any beneficial effects of having 316 large amounts of vitamin E available do not necessarily come through its antioxidant 317 properties. Selman et al. (2008) showed that supplementation of vitamin E significantly 318 increased median lifespan of mice by 15 % relative to non-supplemented controls and 319 also increased maximum lifespan. Lymphocyte and hepatocyte oxidative DNA damage 320 and hepatic lipid peroxidation were, however, unaffected by supplementation. Using a 321 cDNA macroarray, Selman et al. (2008) showed that genes associated with xenobiotic 322 metabolism were significantly upregulated in the livers of mice, suggesting that vitamin 323 E is initially metabolized by cytochrome P450s in a manner similar to xenobiotics. The 324 absence of any significant effect on oxidative damage suggests that the lifespan 325 extension observed in mice was not due to the antioxidant properties of vitamin E.

In various invertebrates (e.g., molluscs, crustaceans) and fish, data on development of antioxidant defences suggested that dietary and other low molecular weight antioxidants are predominant in the earliest stages of development, while antioxidant enzymes become predominant later in life (Rudneva, 1999; Surai, 2002;

Fontagné et al., 2008). We do not know if any beneficial effects of dietary antioxidants could have been stronger at earlier stages of development. This is an open question that deserves further work to understand at which developmental stage intake of dietary antioxidants might be more beneficial. It will also be important to assess the amount of tocopherols that can be retained in tissues and which forms of tocopherols and of their metabolites have a stronger impact on the antioxidant machinery during development (Brigelius-Flohé and Traber, 1999; Surai, 2002).

337 Our results also show that experiencing a period of food deprivation may have 338 long-term effects on the protein oxidative damage in brain. Irrespective of the quality of 339 re-feeding diet, all the fish that underwent food restriction had higher oxidative damage 340 in brain later in life than those that did not. Restriction of intake of specific nutrients in 341 early life may have pervasive, long-lasting effects on oxidative status. For example, rats 342 subjected to prenatal and postnatal protein malnutrition had increased oxidative damage 343 to lipids and proteins in brain (Feoli et al., 2006). Zebra finches (*Taeniopygia guttata*) 344 that experienced a short period of low quality nutrition during the nestling period had a 345 twofold reduction in plasma levels of carotenoids, vitamins A and E in adulthood 346 (Blount et al., 2003). In fish, starvation may elicit profound changes in brain energy 347 metabolism in a way similar to that demonstrated in other vertebrates like mammals. 348 For example, when the fasting period is prolonged and glucose becomes less available 349 with time, the brain obtains a portion of its energy from ketone bodies (Hasselbalch et 350 al., 1994; Campbell and Farrel, 2006), which are important generators of protein 351 carbonylation (Halliwell and Gutteridge, 2015). Moreover, during prolonged fasting, 352 animals face increased protein catabolism and body mass loss (McCue, 2010, 2012), 353 which might favour accumulation of protein carbonyls.

354 In conclusion, our work provides evidence that any beneficial effects of dietary 355 antioxidants on growth might be offset by an impaired capability of regulating the 356 oxidative status. If investing more into compensatory growth is more advantageous than 357 into self-maintenance and mechanisms favouring lifespan likely depends on the 358 environmental conditions young develop in, such as predation risk or food availability. 359 Work on oxidative stress resistance under this life-history framework would help to 360 clarify its role as a mediator of some early life tradeoffs across different environments. 361 Our results also point for a role of oxidative stress as a mechanism linking the long-term 362 negative effects of early life poor nutrition on brain function.

363

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369

- **370 Competing interests**
- 371 No competing interests.
- 372

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Table 1. Outcomes of full and reduced statistical models. FD = Food Deprivation;
POSTFD = diet after the food deprivation. The reduced models were obtained after
backward elimination of non-significant interactions from the full models, starting from
the three-way interaction. Outcomes of full and reduced models coincided when there
was not backward elimination because the interaction/s was/were significant. Markers
of oxidative status were analysed in a subsample of 40 individuals (10 for each group).

		Full model			Reduced		
Variable	Effect	df	F	P	df	F	n
Body size PC1 – Postdeprivation standard diet	Treatment group	1,147	28.6	<0.001	1,147	28.6	<0.001
	Sampling day	2,147	160.8	< 0.001	2,147	160.8	< 0.001
	Treatment group × Sampling day	2,147	8.3	<0.001	2,147	8.3	<0.001
Body size PC1 – Postdeprivation enriched diet	Treatment group	1,152	14.1	<0.001	1,152	13.5	<0.001
	Sampling day	2,152	209.2	<0.001	2,152	205.2	<0.001
	Treatment group × Sampling day	2,152	2.9	0.06			
Body length	Treatment group ED	1 280	15.7	<0.001	1 20/	15.6	<0.001
body length	Treatment group POSTED	1,209	0.05	0.82	1,294	0.06	0.80
	Sampling day	2 289	356.9	<0.02	2 294	357.2	<0.00
	Treatment group FD × sampling day	2,289	4.5	0.012	2,294	4.6	0.011
	Treatment group POSTFD × sampling day	2,289	0.8	0.46			
	Treatment group FD × Treatment group POSTFD	1,289	1.4	0.23			
	Treatment group FD × Treatment group POSTFD × sampling day	2,289	0.9	0.40			
Disama DOMa	Treater ant array of ED	4.00	0.4	0.007	4.00	0.4	0.007
Plasma ROIvis	Treatment group PO	1,39	0.1	0.007	1,39	0.1	0.007
	Treatment group FD × Treatment group POSTFD	1,39	16.4	<0.001	1,39	16.4	<0.001
Plasma protein carbonyls	Treatment group FD	1,39	3.7	0.063	1,39	3.7	0.063
•	Treatment group POSTFD	1,39	4.5	0.042	1,39	4.4	0.043
	Treatment group FD × Treatment group POSTFD	1,39	1.3	0.26			
Brain protein carbonyls	Treatment group FD	1,39	7.2	0.011	1,39	7.4	0.01
	Treatment group POSTFD	1,39	1.8	0.19	1,39	1.9	0.18
	Treatment group FD × Treatment group POSTFD	1,39	0.01	0.92			
	Treatment group ED	1 20	2.2	0.15	1 20	2.2	0.15
	Treatment group POSTED	1,39	2.2	0.10	1.39	2.2	0.15
	Treatment group FOSTFD Treatment group POSTFD	1,39	1.4	0.24	1,00	3.1	0.004
		1.20	0.04	0.05	1.20	0.04	0.95
Diain GPX	Treatment group FD	1,39	0.04	0.85	1,39	0.04	0.85
	Treatment group POSTED Treatment group FD × Treatment group POSTED	1,39	2.5	0.36	1,39	0.0	0.37
		1					
Red blood cell SOD	Treatment group FD	1,39	1.0	0.32	1,39	1.0	0.32

	Treatment group POSTFD	1,39	5.7	0.02	1,39	5.8	0.022
	Treatment group FD × Treatment group POSTFD	1,39	0.9	0.35			
Brain SOD	Treatment group FD	1,39	0.2	0.68	1,39	0.2	0.68
	Treatment group POSTFD	1,39	5.8	0.02	1,39	5.7	0.022
	Treatment group FD × Treatment group POSTFD	1,39	1.6	0.21			

### 487 **Figure captions**

488 Figure 1. Experimental setting and timeline.

489

Figure 2. Food deprivation caused a slowdown of growth as suggested by the smaller
body size compared to fish that were not food-deprived. Refeeding caused accelerated
growth only in those fish that were fed on a diet richer in vitamin E.

493

Figure 3. Food deprivation caused a slowdown of skeletal growth as suggested by the smaller body length of food-deprived fish compared to fish that were not food-deprived soon after the food deprivation period (Tukey test, p<0.01) and at the end of the refeeding experiment (Tukey test, p<0.01).

498

Figure 4. Fish that made a compensatory growth (black square) had higher plasmaoxidative damage (expressed as reactive oxygen metabolites).

501

Figure 5. Fish fed on the diet richer in vitamin E had higher plasma protein oxidative
damage (protein carbonyls) and activity of glutathione peroxidase in red blood cells
(RBS) and lower activity of superoxide dismutase in both red blood cells and brain.

505

Figure 6. Fish that experienced food deprivation had higher brain protein oxidativedamage (protein carbonyls) than those fish that were fed ad libitum.











