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**Dietary amino acids impact sperm performance traits for a catadromous fish, *Anguilla anguilla* reared in captivity**

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24 **Abstract**

25

26 Little is known about the role of dietary amino acids on male reproductive performance and gamete  
27 quality in fishes. Thus, the objective of this study was to investigate how “enhanced” feeds (EH-  
28 4, EH-5, EH-6), with modified amino acid composition, and the standard on-growing diet (DAN-  
29 EX) impact body composition, milt biochemistry, and sperm performance in male European eel,  
30 *Anguilla anguilla*. The fatty acid composition of EH-4, EH-5, and EH-6 was similar but differed  
31 to that in DAN-EX, while amino acid composition varied between all four diets. Diet did not  
32 influence organ-somatic indices (e.g. HSI, GSI), while males fed EH-4 were heavier than other  
33 groups. Arginine, alanine, and lysine were the most abundant amino acids in milt (>11%), followed  
34 by glycine, aspartic acid, valine, glutamic acid, and leucine (>5.66%). Diet impacted milt arginine,  
35 serine, proline, methionine, and histidine levels. Specifically, males fed DAN-EX, EH-4, and EH-5  
36 had the highest percentages of arginine, while males fed EH-4 to EH-6 had higher percentages of  
37 serine. Proline was most abundant in males fed DAN-EX, EH-5, and EH-6. Both methionine and  
38 histidine were detected at low percentages (<2%), and were impacted by diet, where males fed  
39 EH-4 and EH-5 had higher percentages of methionine, and males fed DAN-EX, EH-4, and EH-6  
40 had the highest percentage of histidine. Milt production increased over time, where eels fed EH-4  
41 and EH-6 showed the highest probability of producing suited milt volumes (>0.5 mL) for  
42 fertilization procedures. Spermatocrit ( $43.1 \pm 1.80\%$ ) did not differ between the diets (ranged from  
43 37.57 to 47.21%). Dietary regime had an impact on sperm motility, such that eels fed EH-5 and  
44 EH-6 had the greatest percentage of motile cells. In addition, fish fed EH-5 and EH-6 (or DAN-  
45 EX) had the fastest swimming sperm. Spermatogenic maturity index of hormonally treated eels  
46 varied within groups but did not differ between dietary treatment groups after 9 weeks of injections

47 (ranged from 0.54 to 0.80). The most interesting amino acids to scrutinize from PCA plots were  
48 proline, histidine, and valine as well as lysine and arginine. Here, eels with highly motile sperm  
49 had milt with high relative proportions of proline, histidine, and valine, but were particularly low  
50 in lysine and arginine. Together, our findings add evidence that certain amino acids regulate milt  
51 biochemistry, and that male ejaculate traits may be promoted by amino acid intake. Further studies  
52 to evaluate effects of supplemented amino acid diets on fertilization ability and inter-linked early  
53 developmental stages are required.

54

55 Keywords: Aquaculture; Broodstock diet; Assisted Reproduction; Gamete quality; European eel

56

## 57 **1. Introduction**

58

59 When reared in captivity, fishes may exhibit different degrees of reproductive dysfunction,  
60 such as spermiation failure, spontaneous ovulation, or decreases in gamete quality (Mylonas et al.,  
61 2017). Poor gamete quality may jeopardize the survival of offspring, especially during the  
62 “critical” early life history stages, which can ultimately impact aquaculture production and limit  
63 possibilities for selective breeding (Bromage et al., 1992; Kjørsvik et al., 2003). Therefore, a stable  
64 supply of “high-quality” gametes is required for establishment of sustainable culture for any given  
65 species. In this scenario, controlling sperm quality can be a major issue for the aquaculture  
66 industry, since it is affected by a variety of factors including broodstock nutrition, epigenetics,  
67 and/or sperm handling (Cabrita et al., 2014).

68 It is widely accepted that broodstock nutrition or enriched diets with certain compounds  
69 greatly modulate sperm physiology and functionality (Labbé et al., 1995; Lahnsteiner et al., 2009;



70 Beirão et al., 2015) as well as male reproductive success (Asturiano et al., 2001). In this regard,  
71 most studies have supplemented diets with modified lipid and fatty acid profiles (Asturiano et al.,  
72 2001; Vassallo-Agius et al., 2001; Nandi et al., 2007; Cabrita et al., 2014; Beirão et al., 2015).  
73 While manipulation of lipids in broodstock diets offers excellent opportunities to improve gamete  
74 quality, the protein and amino acids should similarly receive consideration. The biological  
75 functionality of amino acids is diverse, as they are involved in feed intake, nutrient utilization, and  
76 reproduction (Izquierdo et al., 2001; Li et al., 2009; Wu, 2009). Additionally, they are precursors  
77 for a wide variety of macromolecules, such as nucleotides, lipids, glycogen, steroids (Finn and  
78 Fyhn, 2010), and serve as oxidizable substrates for sperm (Mann and Lutwak-Mann, 1981).

79 Evidence has accumulated that amino acids are highly beneficial for improving reproduction  
80 and directly affect fertilization success and the number of viable embryos (Amirkhanov, 1980;  
81 Dabrowski et al., 1985; Kwasek et al., 2014a). For example, certain amino acids (e.g. arginine) are  
82 abundant in physiological fluids (Wu et al., 2009) and play a crucial role in regulating reproductive  
83 functions (Li et al., 2009), survival of juvenile fish (Buentello and Gatlin, 2001), as well as  
84 neuronal development and neurotransmission (Jobgen et al., 2006; Yao et al., 2008). Additionally,  
85 in fish, amino acid-rich proteins called protamines within sperm nuclei are involved in sperm cell  
86 growth and differentiation (Martinage et al., 1985), thus underlining a critical role for amino acids  
87 in spermatogenesis (Wu et al., 2009). Previous studies reported that taurine (derived from cysteine)  
88 at levels of 10 g/kg improved spawning in some marine fish species (Matsunari et al., 2006). In  
89 Japanese eel, Higuchi et al. (2012) determined that taurine is essential for spermatogenesis,  
90 although it can be synthesized from cysteine in the testis under the action of dihydroxyprogesterone  
91 (DHP). The consequences of dietary supplementation of amino acids on male gamete quality and  
92 performance are fascinating, due to the fact that amino acids are a key component of seminal

93 plasma and sperm (Lahnsteiner 2009, 2010; Kwasek et al., 2014b), impacting sperm metabolism  
94 and sperm motility and/or fertility (Patel et al., 1998; He and Woods, 2003). In addition, a complex  
95 antioxidant defense system exists in milt (e.g. superoxide dismutase, catalase, and the glutathione  
96 peroxidase-glutathione reductase system), which is highly influenced by nutrition (Mansour et al.,  
97 2006). Thus, dietary supplementation of specific amino acids may provide a novel approach to  
98 improve fertility in males.

99 Sperm performance and composition in vertebrates are clearly affected by dietary  
100 supplementation of amino acids (Wu et al., 2009; Kwasek et al., 2014b; Pourkhazaei et al., 2017).  
101 For example, in mammals, either supplementation of diets with amino acids or direct injection  
102 improved sperm traits (e.g. motility, velocity, morphologically normal sperm, and acrosome  
103 integrity) and subsequent fertilization success (Dong et al., 2016; Abd-Elrazek and Ahmed-Farid,  
104 2018). In teleosts, only a few studies have investigated incorporation of dietary amino acids on  
105 male gonadal development and gamete quality (Akiyama et al., 1996; Kwasek et al., 2014b;  
106 Pourkhazaei et al., 2017), and without directly linking the diet to the kinetic characteristics of  
107 sperm. As such, knowledge regarding the role of dietary amino acids on gamete quality and  
108 reproductive performance of male fish remains incomplete.

109 Here, we use European eel, *Anguilla anguilla* as our model organism. Despite increasing  
110 efforts (Asturiano et al., 2005; Gallego et al., 2012; Tomkiewicz, 2012; da Silva et al., 2018; Politis  
111 et al., 2018a; Benini et al., 2018), variability in gamete quality is still an issue affecting larval  
112 production for this species. During the last decade, different strategies have been employed to  
113 improve European eel gamete quality, especially by conducting studies on reproductive  
114 performance and broodstock diets at specific developmental windows (Støttrup et al., 2013; Baeza  
115 et al., 2014, 2015a,b; Butts et al., 2015; da Silva et al., 2016). For example, supplementation of

116 broodstock diets with lipids [mainly polyunsaturated fatty acid (PUFA)], improved spermiation  
117 and milt quality in males (Butts et al., 2015), and oocyte growth and ovarian development in  
118 females (Støttrup et al., 2016; da Silva et al., 2016). Therefore, further information on the effects  
119 of broodstock diets on European eel gamete performance could enhance the process of  
120 domestication for mass production of larvae. Nevertheless, the effect of supplementation of the  
121 broodstock diet with amino acids on gamete quality has not yet been investigated for this species.  
122 As such, we hypothesize that dietary supplementation of specific amino acids will be beneficial  
123 for male gamete performance.

124 The objective of this study was to investigate how “enhanced” feeds (EH-4, EH-5, EH-6),  
125 with modified amino acid composition, and the standard on-growing diet (DAN-EX) impact body  
126 composition, milt biochemistry, and sperm performance in male European eel. Together, these  
127 data may be used to improve broodstock diets for this catadromous fish and increase our  
128 understanding on reproductive physiology in fishes.

129

## 130 **2. Materials and methods**

131

132 All fish were handled in accordance with the European Union regulations concerning the  
133 protection of experimental animals (Dir 86/609/EEC). Eel experimental protocols were approved  
134 by the Animal Experiments Inspectorate (AEI), Danish Ministry of Food, Agriculture and  
135 Fisheries (permit number: 2015-15-0201-00696). All efforts were made to minimize animal  
136 handling and stress.

137

138

139 *2.1. Eel broodstock husbandry*

140

141 Eels were raised from wild-caught glass eels in a commercial eel farm, Stensgården Eel Farm  
142 A/S in Jutland, Denmark. Eels were grown on commercial feed (DAN-EX; BioMar A/S, Brande,  
143 Denmark) in concrete tanks (18-30 m<sup>3</sup>) at stocking density of ~50 kg/m<sup>2</sup>. Rearing tanks were  
144 equipped with recirculating aquaculture system (RAS) technology, which consisted of a rotating  
145 drum filter, biofilter, trickle-filters, UV system, and oxygen cones. Water was salted to 1-2 PSU  
146 and heated to 23 ± 2°C. Upon start of experimental trials, the eels were sorted based on size and  
147 then transferred to 4 fiberglass tanks (2 × 2 m, volume of 2.5 m<sup>3</sup>, flowrate 1.5 to 2 m<sup>3</sup>/h) at an  
148 initial stocking density of 50 kg/m<sup>2</sup>. Males were fed with automatic feeders six times per day at  
149 0.5 to 0.8% body weight from 1 August 2015 to 2 February 2016 on three “enhanced” feeds (EH-4,  
150 EH-5, EH-6; Table 1) and the standard on-growing diet (DAN-EX) varying in fatty acid (Table 2)  
151 and amino acid composition (Table 3). Final stocking density was ~80 kg/m<sup>2</sup>.

152 Upon completion of the commercial feeding trail, ~35 males from each dietary regime were  
153 randomly selected and transported to a Technical University of Denmark (DTU Aqua) research  
154 facility in Hirtshals, Denmark (57.585971 N, 9.985036 E). The eels were housed in 4 × 500 L  
155 tanks equipped with RAS technology at a flowrate of 600 L/h. The RAS system consisted of a 350  
156 L gravel filter, 0.3 m<sup>3</sup> trickle filter, and a 300 L temperature regulated sump. Temperature ranged  
157 between 19-21°C, salinity ranged from 36-37 PSU, and photoperiod was kept at 12h light/12 h  
158 dark at ~20 lux. Natural seawater salinity of ~32.5 PSU from the North Sea was adjusted using  
159 Tropic Marin Sea Salt (Dr. Biener GmbH, Wartenberg, Germany) and verified using a  
160 conductivity meter (WTW multi3410, Wissenschaftlich-Technische Werkstätten GmbH,  
161 Weilheim, Germany). Acclimatization to sea water took place over one week before entering the

162 facility. No feed was provided during experimentation as eels cease feeding during the silvering  
163 process and are not anticipated to feed during spawning migration (Tesch, 2003).

164 After acclimatization, 32 randomly selected males (8 per diet, >100 g) were euthanized by  
165 submergence in an aqueous solution of benzocaine and their body weight and length were recorded  
166 to get an indication of their morphological status after receiving the experimental diets. These 32  
167 eels were also dissected and liver weight was obtained. The remaining eels (n = 97) were  
168 anaesthetized by short-term submergence in benzocaine and tagged with a passive integrated  
169 transponder (PIT tag). The PIT tag was placed in the dorsal muscle. Male eels from each diet  
170 (DAN-EX = 27, EH-4 = 25, EH-5 = 25, EH-6 = 20) received weekly injections of human chorionic  
171 gonadotropin (hCG, Sigma Aldrich Denmark A/S) at 1.5 IU/g fish in order to induce milt  
172 production (Pérez et al., 2000). Body weight was recorded at the time of first injection and then at  
173 injection Week 4, Week 7, and Week 9. In addition, ~24 h after the Week 9 injection, 8-9 males  
174 were randomly selected per diet. Among these, the spermiating males ranged between 94-126 g.  
175 Again, after eels were euthanized, body weight, testes weight, and liver weight were recorded.  
176 Gonadosomatic index (GSI;  $100 \times \text{testes weight/body weight}$ ) and hepatosomatic index (HSI;  $100$   
177  $\times \text{liver weight/body weight}$ ) were later calculated.

178

## 179 *2.2. Broodstock feed*

180

181 Previous studies have investigated dietary impacts on egg quality (Heinsbroek et al., 2013;  
182 Støttrup et al., 2013, 2016). Based on these findings, fatty acid composition of the best performing  
183 diet for females was used as basis for the fatty acid composition of these tested diets (hereafter  
184 EH-4, EH-5, and EH-6). In terms of fatty acid composition, EH-4, EH-5, and EH-6 differed

185 significantly from the commercial DAN-EX feed (Table 2). Amino acid composition varied  
186 between all four diets, where the enhanced feeds were originally modified with respect to arginine  
187 content (Table 1, Table 3). Here, arginine was lowest in EH-5 and highest in EH-6, while EH-4  
188 showed intermediate levels.

189

### 190 *2.3. Amino acid analysis of milt*

191

192 Amino acid composition was analyzed using a EZ:fast™ amino acid analysis kit  
193 (Phenomenex Inc., Torrance, CA, USA). Between ~300 to 2000 mg of sperm samples were used  
194 for the analysis depending on how much sample was available. Separation and detection of amino  
195 acids occurred by liquid chromatography using Agilent G6120BA single-quadrupol LC-MS in the  
196 ESI ionization mode (Agilent Technologies, Hørsholm, Denmark). The LC was equipped with an  
197 EZ:fast™ Liquid chromatography-Mass spectroscopy (LC-MS) column (250 mm × 3.0 mm,  
198 Phenomenex). Amino acids were analyzed in duplicate or triplicate depending on milt volume.  
199 Further details for amino acid analyses are described in Safafar et al. (2016).

200

### 201 *2.4. Milt production*

202

203 Males (n = 97 eels at start; n = 27 for DAN-EX; n = 25 for EH-4; n = 25 for EH-5; n = 20  
204 for EH-6) were assessed weekly for spermiation one day after injection from Week 5 until Week  
205 9. Fish ejaculate sperm spontaneously during natural spawning, but in captivity, it is typically  
206 expressed from the sperm ducts by “stripping”, where gentle abdominal pressure is applied. Thus,  
207 slight pressure was applied to the abdomen of each male to assess their degree of maturity. Milt

208 was then classified between 0-5, where 0 = no milt, 1 = few drops with gentle pressure, 2 = low  
209 volume of milt with gentle pressure (<0.5 mL), 3 = medium volume of milt with gentle pressure  
210 (<1.0 mL), 4 = flowing milt with gentle pressure (>1.0 mL), and 5 = freely flowing. Approximately  
211 24 h after injection on Week 9, the total weight of milt per male was recorded and standardized  
212 (g/100 g eel).

213

#### 214 *2.5. Spermatozoa kinetic traits*

215

216 Milt samples from 83 males (n = 22 for DAN-EX; n = 22 for EH-4; n = 20 for EH-5; n = 19  
217 for EH-6; one male was excluded from further analyses due to possible milt contamination) were  
218 collected 24 h after the Week 9 injection, as the highest sperm quality is obtained at this time  
219 (Pérez et al., 2000). Males were anesthetized with benzocaine (60 mg/L) to minimize any adverse  
220 effects during stripping. The urogenital pore was cleaned with ionized water and wiped dry to  
221 avoid contamination with feces, urine, or blood. To avoid any bias associated with time during  
222 sampling, the milt was collected into weight boats from two males per diet, and this was repeated  
223 until all males were sampled.

224 Immediately after stripping, a milt sample (100  $\mu$ L) from each male was diluted in 900  $\mu$ L  
225 of P1-extender medium (Peñaranda et al., 2010) and then kept in a cooler with frozen icepacks to  
226 maintain viability. Before activation, the samples were inverted for ~5 s for homogenization. The  
227 immobilized sperm suspension (0.2  $\mu$ L) was then pipetted into a chamber of an 80  $\mu$ m 2X-CEL  
228 glass slide (Hamilton Thorne, MA, USA) and covered with a 22  $\times$  22 mm coverslip. The cells  
229 were then activated with 15  $\mu$ L of modified seawater (36 ppt) with 1% bovine serum albumin,  
230 BSA (w/v). The BSA was added to the activation media as it prevents the sperm from sticking to

231 the glass slide.

232 Sperm activity was captured at 10, 20, and 30 s ( $\pm 1$ ) post-activation using a compound  
233 microscope (Nikon Eclipse 55i microscope, Nikon Corporation, Tokyo, Japan) equipped with a  
234 Nikon DS-Fi1 camera head and negative phase objective (PL 40 x /0.66,  $\infty$ /0.17). The digital video  
235 camera was attached to a personal computer and images were captured using a frame grabber at  
236 50 frames/s (Procadi, PROiSER 1.4, Paterna, Spain). Three replicate activations were performed  
237 for each male. Two observers continuously did all activations (752 video recordings) to avoid any  
238 subjective deviation. Total motility (MOT, total number of motile spermatozoa/total number of  
239 cells  $\times 100$ ) and curvilinear velocity (VCL, defined as velocity of sperm along its actual curvilinear  
240 path) were then assessed using a computer-assisted sperm analysis (CASA; ISAS v1; PROiSER  
241 R+D, S.L., Paterna, Spain).

242

## 243 *2.6. Spermatoxrit*

244

245 Spermatoxrit, defined as the ratio of packed sperm cells to the total volume of milt  $\times 100$ ,  
246 was used to estimate sperm concentration on Week 9 (Sørensen et al., 2013). For each male ( $n =$   
247 84 eels), samples of milt were drawn into three microhematocrit capillary tubes (75 mm length  
248 and 1.1-1.2 mm opening) and sealed at the end with Vitrex Sigillum wax. The tube was then  
249 centrifuged for 10 min at  $6000 \times g$  (Haematokrit 210, Andreas Hettich GmbH & Co. KG,  
250 Tuttlingen, Germany) and the average of three replicate tubes was used for statistical analysis.

251

252

253



254 2.7. *Histological analyses*

255

256 To assess testes development, testicular lobes were sampled after 9 weeks of hormonal  
257 treatment from the middle of testes and preserved (n = 8 per diet) in a 4% solution of formalin  
258 buffered by NaH<sub>2</sub>PO<sub>4</sub>-H<sub>2</sub>O and Na<sub>2</sub>HPO<sub>4</sub>-2H<sub>2</sub>O. Subsequently, the tissue samples were  
259 dehydrated, embedded in paraffin, and sectioned at 5 µm. The sections were stained with  
260 haematoxylin and eosin (H & E, VWR-Bie & Berntsen A/S, Denmark). The histological sections  
261 were photographed (Olympus BX53 digital camera) at 200× magnification for identification of  
262 gamete development stages and tissue, i.e. spermatogonia (Sg), spermatocytes (Sc), spermatids  
263 (St), and spermatozoa (Sz). Testes tissues of five micrographs per male were categorized according  
264 to gamete stages (i.e. Sg, Sc, St, and Sz) and their relative area fraction (*F*) and progression of  
265 spermatogenesis was assessed using a spermatogenic maturity index (SMI) (see Tomkiewicz et  
266 al., 2011). SMI was estimated for each of the testes images in order to compare the morphological  
267 development of the testes tissue in males receiving different amino acid diets (n = 8-9 fish per  
268 diet).

269

270 2.8. *Statistical analysis*

271

272 Data were analysed using SAS statistical analysis software (v.9.1; SAS Institute Inc., Cary,  
273 NC, USA) and R programming language (Venables and Ripley, 2002). Residuals were tested for  
274 normality (Shapiro-Wilk test) and homogenous of variances (Levene's test). The significance level  
275 was set at  $\alpha = 0.05$ . Treatment means were determined using the honest significant difference  
276 Tukey's test. Prior to or at the end of the experiment (Week 9), length, weight, liver weight, testes

277 weight, GSI, and HSI were compared across the diets using a series of one-way ANOVAs. During  
278 injections, eel body weights were compared using a repeated measures ANOVA model that  
279 contained the factors: Time, Diet, and Time  $\times$  Diet. One-way ANOVAs were used to compare  
280 amino acids between the dietary groups. For statistical comparisons the amino acid composition  
281 was compared as a percentage.

282 Logistic regression analysis was applied to estimate the parameters of a logistic model  
283 consisting of a dependent variable (*milt production*) with two possible values, “unsuited”  
284 (representing categories 0-2) or “suited” (categories 3-5) for further fertilization procedures (see  
285 Section 2.4). In the logistic model, the log-odds for the value labeled "suited" is a linear  
286 combination of two independent variables; a binary variable (*diet*) and a continuous variable  
287 (*time*). The corresponding probability varied between 0 ("unsuited") and 1 ("suited"). At the end  
288 of the experiment, the total weight of milt per male was recorded and standardized across the diets  
289 using one-way ANOVA model.

290 Spermatoctrit was compared across the diets with a one-way ANOVA model. A one-way  
291 ANOVA model was also used to compare SMI between the diets on Week 9. Total motility and  
292 curvilinear velocity were compared using a repeated measures ANOVA model that contained the  
293 factors: Time (10, 20 and 30 s), Diet, and Time  $\times$  Diet.

294 Additionally, two principal components analyses (PCA) were performed to study the 1)  
295 correlation between amino acid composition (in % of total amino acids) in the diet and in the milt  
296 samples, and 2) the correlation between amino acid composition (in % of total amino acids) in the  
297 milt samples and sperm motility and velocity. In both PCAs, the diets (DAN-EX, EH-4, EH-5, and  
298 EH-6) were used as category variables. All parameters were weighted by 1/SD and full cross  
299 validation was used. For the first PCA, the mean value for each amino acid in each diet was used

300 for all milt samples receiving the same diet.

301

### 302 **3. Results**

303

#### 304 *3.1. Body morphometric measures*

305

306 Prior to hormonal treatment, total body weight, length, and liver weight of eels ranged from  
307 72 to 163 g, 34 to 44 cm, and 0.72 to 1.52 g respectively. There were no significant differences  
308 detected between diets for these three morphometric traits ( $P > 0.05$ ).

309 For the hormonally treated males, total body weight ranged from 104 to 167 g for males fed  
310 DAN-EX, 101 to 179 g for males fed EH-4, 100 to 150 g for males fed EH-5, and 98 to 130 g for  
311 males fed EH-6. For the repeated measures ANOVA, the Time  $\times$  Diet interaction and Time effect  
312 were both non-significant ( $P > 0.05$ ). On the contrary, dietary regime impacted total weight of the  
313 males ( $P < 0.0001$ ; Fig. 1A), such that males fed EH-4 were the heaviest, while males fed EH-5  
314 and EH-6 were the lightest (Fig. 1B).

315 At the end of the experiment (i.e. after Week 9 injection) liver weight, testes weight, GSI,  
316 and HSI were compared across the diets. Here, liver weight and testes weight ranged from 1.04 to  
317 1.09 and 9.42 to 11.41 g, respectively, and there were no significant differences between the diets  
318 ( $P > 0.05$ ). Additionally, GSI (ranged from 7.05 to 9.44) and HSI (ranged from 0.94 to 1.03) were  
319 not significantly influenced by dietary regime ( $P > 0.05$ ).

320

321

322

323 3.2. *Amino acid analysis of milt*

324

325 The amino acids detected in eel milt are displayed in Table 4. Arginine, alanine, and lysine  
326 were the most abundant amino acids in eel milt (all >11%), followed by glycine, aspartic acid,  
327 valine, glutamic acid, and leucine (all > 5.66%). Dietary regime significantly impacted arginine (P  
328 = 0.01), serine (P < 0.0001), proline (P = 0.02), methionine (P = 0.01), and histidine (P < 0.0001)  
329 levels. Specifically, males fed DAN-EX, EH-4, and EH-5 had the highest percentages of arginine,  
330 while males fed EH-4 to EH-6 had highest percentages of serine. Proline was most abundant in  
331 males fed DAN-EX, EH-5, and EH-6. Both methionine and histidine were detected at low levels  
332 (<2%), but were still impacted by dietary regime, where males fed EH-4 and EH-5 had the highest  
333 percentages of methionine, and males fed DAN-EX, EH-4, and EH-6 had highest percentages of  
334 histidine.

335

336 3.3. *Milt production and Spermatocrit*

337

338 Milt production was initially graded from 0 (no milt release) to 5 (flowing milt) and then  
339 grouped as “unsuited” (categories 0-2) or “suited” (categories 3-5; Fig. 2A-F). Logistic regression  
340 showed that milt production was significantly influenced by the time of hormonal treatment (p <  
341 0.001) and dietary regime (p = 0.020). Generally, milt production increased over time from almost  
342 no milt on Week 5 to reach highest values on Week 9, with 65% and 68% probability of milt  
343 “suited” for fertilization procedures when males were fed EH-4 or EH-6 respectively, compared  
344 to 18% or 28% when fed DAN-EX or EH-5, respectively (Fig A-F). Mean ± SEM spermatocrit  
345 for the males was 43.1 ± 1.80% and it did not differ between the diets (P > 0.05, ranged from

346 37.57% for males fed EH-6 to 47.21% for males fed EH-4; Fig. 2G). The one-way ANOVA model  
347 showed no impact of dietary regime on total milt weight, where it ranged from 1.91 g/100 g eel  
348 for males fed EH-6 to 2.32 g/100 g eel for males fed EH-4 ( $P > 0.05$ ; Fig. 2H).

349

#### 350 *3.4. Spermatozoa kinetic traits*

351

352 For sperm motility, the Time  $\times$  Diet interaction ( $P > 0.05$ ; Fig. 3A) and Time effect were  
353 both non-significant ( $P > 0.05$ ). Dietary regime had an impact on sperm motility ( $P = 0.007$ ; Fig.  
354 3B), such that eels fed EH-5 or EH-6 had the greatest percentage of motile cells. The Time  $\times$  Diet  
355 interaction ( $P > 0.05$ ; Fig. 3C) and Time effect were also not significant for sperm velocity ( $P >$   
356  $0.05$ ), while the dietary regime had an impact ( $P = 0.003$ ; Fig. 3D). Here, fish fed the DAN-EX  
357 diet or EH-5 and EH-6 had the fastest swimming sperm.

358

#### 359 *3.5. Histological analyses*

360

361 The SMI of hormonally treated eels ranged from 0.54 to 0.80 after 9 weeks of hormonal  
362 injections and did not differ between dietary treatment groups ( $P > 0.05$ ). All hormonally treated  
363 males responded to treatment with testes showing progressed development, including  
364 spermatocytes (Sc), spermatids (St) and spermatozoa (Sz) (Fig. 4). Sc and St dominated the least  
365 developed males, but also tubules with attached developing Sz were observed (Fig. 4). In contrast,  
366 free Sz in enlarged tubules dominated the most developed males, but still with prevalent Sc and  
367 St. The continuous presence of Sg and Sc showed ongoing spermatogenesis in all males  
368 independently of the diet they received (Fig. 4).

369

370 *3.6. Principal components analysis of amino acid composition in diet and milt samples*

371

372 PCA was performed to study the correlation between the amino acid composition (% of total  
373 amino acids) in the diet and milt, where diets were used as category variables. From this PCA, it  
374 was evident that three samples (two receiving EH-6 and one receiving EH-4) behaved as outliers.  
375 A new PCA without these samples was therefore performed (Fig. 5A and B). PC1 and PC2  
376 explained 43 % and 15 % of the variation in the data, respectively (Supplementary Table 1). Both  
377 the scores and the loadings plot showed that PC1 explained the difference between the DAN-EX  
378 diet/milt and the other samples, whereas PC2 mainly explained differences between EH-5 and EH-  
379 6 diet/milt. When taking these findings into consideration, the most interesting amino acids to  
380 scrutinize further in the second PCA were proline, histidine and valine as well as lysine and  
381 arginine (Fig. 5A and 5B). In Fig. 5B, proline, histidine and valine in the milt samples were located  
382 close to EH-6 (in the first quadrant). These three amino acids in the diets (D-PRO, D-HIS, and D-  
383 VAL) were all located to the far right indicating that the DAN-EX diet had a low content of these  
384 three amino acids and that there was a positive correlation between the content of these amino  
385 acids in the diet and in the milt samples. The same was also the case for glutamic acid and D-GLU,  
386 which were both located to the left. Interestingly, lysine and arginine in the milt were located in  
387 the 3<sup>rd</sup> quadrant, whereas the variables for these two amino acids in the diet (D-LYS and D-ARG)  
388 were located directly opposite in the 1<sup>st</sup> quadrant, suggesting a negative correlation between the  
389 presence of these amino acids in the diet and in the milt samples. The same was also the case for  
390 other amino acids such as hydroxyproline (HYP to the left) and D-HYP (to the right) and for  
391 phenylalanine (PHE to the right) and D-PHE (to the left).

392 3.7. *Principal components analysis of sperm motility and amino acid composition in milt*

393

394 In order to study the correlation between sperm motility and amino acid composition (in %  
395 of total amino acids) a PCA including all samples and using diets as category variables was  
396 performed. All samples excluding the three outliers mentioned above were used for the analysis  
397 (Fig. 6A and B). PC1 and PC2 together explained 43% of the variation in the data (Supplementary  
398 Table 2). The scores plot (Fig. 6A) showed that PC1 mainly explained differences between DAN-  
399 EX samples to the left and the samples obtained from eel receiving the other diets (EH-4 to EH-  
400 6), which were mainly located to the right. PC2 mainly explained differences between eels  
401 receiving diet EH-4 in the top of the plot and EH-6 in the bottom of the plot. This interpretation of  
402 the scores plot was also confirmed by the location of the diet category variables in the correlation  
403 loadings plot (Fig. 6B). The correlation loadings plot showed a clear positive correlation between  
404 motility parameters in the lower right corner and eel receiving diet EH-6. However, this  
405 interpretation of the model was only partly confirmed by the original data in Figure 3, which did  
406 not show a significant difference between EH-5 and EH-6. Further inspection of the scores plot in  
407 Fig. 6A showed some overlap between the locations of the samples receiving different diets.  
408 Particularly EH-4 and EH-5 samples were scattered in the plot. It was, however, clear that all  
409 DAN-EX samples were located to the left in the scores plot and all EH-6 samples were located in  
410 the lower part of the plot and mainly to the right. These findings thus suggest that eels fed EH-6  
411 to a higher degree than eels fed DAN-EX had high motility sperm. Eels with high motility sperm  
412 had milt with high relative proportions of proline, histidine and valine, but were particularly low  
413 in lysine and arginine. They were also to some extent high in the amino acids located in the lower  
414 part of the 1<sup>st</sup> quadrant (threonine, isoleucine, serine, glycine).

415 **4. Discussion**

416

417 In teleosts, studies have focused on amino acid requirements for growth and metabolism (Li  
418 et al., 2009), however, their physiological significance in relation to reproduction and/or gamete  
419 performance has not been well elucidated. Proteins and amino acids are the most abundant  
420 constituents in fish gametes [e.g. free amino acids (FAAs) constitute up to 50% of the total amino  
421 acid pool in marine pelagic fish eggs]. Here, they serve as an energy source during embryonic  
422 development (Rønnestad et al., 1992; Syama Dayal et al., 2003), are important osmotic effectors  
423 during oocyte hydration (Cerdà et al., 2007), and can even impact fertilization success (Kwasek et  
424 al., 2014a). Moreover, dietary protein/amino acids modulate the time of puberty and rate of  
425 maturation indirectly by impacting growth (Gunasekera et al., 1995). Therefore, dietary  
426 supplementation of amino acids can provide quantitative evidence on whether the reproductive  
427 performance or gamete quality in fish can be modified, particularly under captive conditions.

428 Amino acids or short peptides are produced as hydrolysates of intercellular matrix proteins,  
429 to act as signals for maturation and timing of spermiation (Kawabata et al., 1992). In the present  
430 study, diet did not influence organ-somatic indices (e.g. HSI, GSI) and testes histology, while it  
431 impacted total weight of the males, where males fed EH-4 were the heaviest in comparison to other  
432 groups. Moreover, induction of spermiation was significantly impacted by diet and time of  
433 hormonal treatment, such that eels fed EH-4 and EH-6, on each week, showed the highest  
434 probability of producing “suited” milt for further fertilization procedures, reaching values of 65%  
435 and 68% on week 9, respectively. This observation is in line with several other studies in which  
436 certain amino acids positively impacted spermiation and male or female reproductive success. For  
437 example, in ayu, *Plecoglossus altivelis*, additional tryptophan in the broodstock diet advanced the



438 peak of serum testosterone levels and spermiation time in males and final maturation in females  
439 (Akiyama et al., 1996). In the male rose bitterling *Rhodeus ocellatus*, Kawabata et al. (1992)  
440 indicated that spermiation and sexual behavior was induced by several amino acids, such as  
441 cysteine, serine, alanine, glycine, and lysine. In addition, supplementation of a diet with higher  
442 levels of lysine significantly increased milt volume in silver catfish, *Rhamdia voulezi* (Diemer et  
443 al., 2014), indicating relationships between levels of certain amino acids in the diet and  
444 spermiation. Therefore, in the present study, it is quite likely that well-balanced amino acids or  
445 specific amino acids in diets EH-4 and EH-6 were more favorable for European eel  
446 spermatogenesis in comparison to the diets EH-5 and DAN-EX. Moreover, certain amino acids  
447 (e.g. tyrosine, phenylalanine, glutamine, and leucine) are precursors for the synthesis and secretion  
448 of hormones such as thyroid hormones, insulin hormones, growth hormones, prolactin, and  
449 progesterone (Wu, 2009). Presumably, in the present study, elevated levels of these hormones in  
450 males fed EH-4 and EH-6 may have partly mediated induction of spermatogenesis, which warrants  
451 further investigation also in relation to transmission of effects to offspring as these hormones play  
452 a key role during early life development (Politis et al., 2017, 2018b, 2018c).

453 Sperm motility and velocity are regarded as primary determinants of reproductive success  
454 and are commonly used to assess male gamete quality and fertilization potential (Gage et al., 2004;  
455 Rurangwa et al., 2004; Gallego and Asturiano, 2018a,b; Zadmajid et al., 2019). Typically, sperm  
456 with higher velocity and motility have the advantage of reaching the micropyle within a shorter  
457 window of time during a fertilization event, while correlations have been found between sperm  
458 motility parameters and fertilization rates in several fish species (reviewed by Gallego and  
459 Asturiano, 2018a). Sperm motility parameters have been positively impacted by broodstock  
460 nutrition for various fish species, such as barbel, *Barbus barbus* (Alavi et al., 2009), Senegalese

461 sole, *Solea senegalensis* (Beirão et al., 2015), and European eel (Butts et al., 2015). Contrary to  
462 fish, in mammals, the role of amino acids on sperm quality/function has received great attention.  
463 For instance, incubation of goat sperm with specific amino acids (e.g. arginine), not only enhances  
464 the pH and metabolic activity of sperm, but also the synthesis of ATP, which is essential for sperm  
465 motility (Patel et al., 1998). In addition, supplementation of diets with amino acids improved sperm  
466 motility and subsequent fertilization success in mice (Bahadorani et al., 2019), while for humans,  
467 amino acid-deficient diets resulted in a ~10-fold increase in the percentage of non-motile sperm  
468 (Wu et al., 2000). This striking observation underlines a critical role of amino acids for male  
469 gamete performance.

470 We observed that dietary amino acids impacted sperm quality, where eels fed EH-5 or EH-  
471 6 presented an improvement in sperm motility parameters, which most likely increases chances  
472 for sperm to achieve fertilization. The underlying mechanism(s) may be related to enhanced  
473 synthesis of polyamines and amino acid-rich basic proteins in the sperm cells (Wu et al., 2009) or  
474 activation of signaling molecules such as nitric oxide, which acts as a stimulator of sperm  
475 motility/velocity (Creech et al., 1998; Barman et al., 2013). On the other hand, protein  
476 phosphorylation processes, which trigger further cell signaling processes such as cyclic adenosine  
477 monophosphate (cAMP) and hydrolysis of ATP catalyzed by dynein ATPase, are important  
478 regulatory components for sperm swimming trajectories (Dzyuba and Cosson, 2014; Zilli et al.,  
479 2008,2017). Therefore, balanced amino acids in the diets EH-5 or EH-6 may have increased  
480 intracellular ATP stores (Perchee Poupard et al., 1998), or changed the protein phosphorylation  
481 state via production of proteins, which are involved in sperm motility activation, such as motor  
482 proteins [e.g. A-kinase anchor protein (AKAP), axonemal dynein; Zilli et al., 2017], signaling  
483 proteins [e.g. protein kinase A (PKA), Caspase 3, cleavage of PARP; Silva et al., 2015], and

484 proteins involved in cell metabolism, including metabolism of reactive oxygen species (ROS) [e.g.  
485 Acetyl-CoA synthetase, Cu/Zn superoxide dismutase (Cu/Zn SOD); Zilli et al., 2017].  
486 Interestingly, in our study, PCA analysis showed that eels with high motility sperm had milt with  
487 high relative proportions of proline, histidine, and valine, but were particularly low in lysine and  
488 arginine. However, when comparing to other studies, the impact of amino acids on teleosts sperm  
489 traits shows high species-specific variability. For example, *in vitro* incubation of rainbow trout,  
490 *Oncorhynchus mykiss* spermatozoa with proline, isoleucine, and methionine had a positive effect  
491 on sperm traits (e.g. motility, velocity and viability), while, proline, glutamine, cysteine,  
492 asparagine, isoleucine, phenylalanine, serine, and histidine had a negative impact on common carp,  
493 *Cyprinus carpio* sperm viability (Lahnsteiner, 2009). In both perch, *Perca fluviatilis* and gilthead  
494 sea bream, *Sparus aurata*, glycine, lysine, methionine, and serine had a positive effect on sperm  
495 motility *in vitro* (Lahnsteiner, 2010). In male yellow perch, *Perca flavescens* sperm motility and  
496 fertilization rate were significantly decreased in the lysine deficient group (Kwasek et al., 2014a,b).  
497 In addition, there is some evidence that *in vitro* supplementation of arginine with sperm cells  
498 positively impacts sperm swimming behaviors in fish (Lahnsteiner, 2010), human (Keller and  
499 Polakoski, 1975), rabbit (Radany and Atherton, 1981), and rat (Abd-Elrazek and Ahmed-Farid,  
500 2018). The impact of amino acids on sperm motility and velocity has also been highlighted during  
501 the freeze-thawing processes by protecting sperm cells against free-radical-induced damage  
502 (Cabrita et al., 2011; Sangeeta et al., 2015). In ram semen, supplementation of freezing media with  
503 proline led to a significant improvement in sperm motility, velocity, and structural and functional  
504 integrity of biological membranes during the freezing and post-thawing process (Sangeeta et al.,  
505 2015). From the above reports, it clearly emerges that protein and amino acids are highly involved  
506 in sperm motility initiation by different mechanisms, but with specific amino acid preference

507 among species. Generally, it is not surprising that amino acids have species-specific impacts on  
508 teleosts sperm performance, as amino acid composition and metabolism in general differs greatly  
509 between fish.

510 In teleosts, it is well documented that dietary amino acid profiles are influencing post-  
511 feeding levels of amino acids in the body, tissues, liver, and muscles (Kaushik et al., 1988; Mai et  
512 al., 2006; Mozanzadeh et al., 2018). In addition, several authors have suggested that profiles of  
513 amino acids present in the plasma are directly related to dietary composition. For example, dietary  
514 supplementation of amino acids, enhanced the concentration of several amino acids in either blood  
515 plasma or seminal plasma in different fish or mammal species (Tantikitti and March, 1995; Wu et  
516 al., 2007; Dong et al., 2016). Nevertheless, relatively little information is available about the amino  
517 acid composition of male germ cells and whether their accumulation impacts sperm functionality.  
518 Available evidence shows that amino acid profiles in fish gametes are directly related to sperm  
519 quality and fertilization success (He and Woods, 2003; Kwasek et al., 2014b). In support of this  
520 notion, quantifying the free amino acid composition of sperm from several fish species revealed  
521 that amino acids play a significant role in stimulation of sperm metabolic activity and viability,  
522 and participates in various detoxifying functions (Lahnsteiner, 2009, 2010). Interestingly, in our  
523 study, even on a quantitative basis, there was high accumulation (>35% of total) of arginine,  
524 alanine, and lysine in milt, however, their incorporation at higher concentrations could not impact  
525 sperm function in eel compared to other amino acids such as proline, histidine, and valine as  
526 validated by PCA analysis. Similarly, previous studies on male European eel (Baeza et al., 2014,  
527 2015a) reported that the specific use of every type of fatty acid depends on the tissue and phase of  
528 spermatogenesis. Overall, the physiological roles of specific fatty acids and amino acids must be  
529 clarified. In addition, the concentration of amino acids in male gonads or germ cells varies among

530 and even within species, which makes it difficult to discern a particular pattern in amino acid  
531 profiles. For example, methionine, arginine, and cysteine were found to be the main amino acids  
532 in milt of rainbow trout (Lahnsteiner, 2009); leucine, arginine, glutamic acid, histidine, and lysine  
533 in common carp, *Cyprinus carpio* (Lahnsteiner, 2009); arginine, alanine, isoleucine, tyrosine,  
534 asparagine, methionine, tryptophan, glutamic acid, and lysine in perch (Lahnsteiner, 2010); and  
535 leucine, arginine, methionine, glycine, hydroxyproline, cysteine, isoleucine, serine, glutamic acid,  
536 lysine, phenylalanine, and asparagine in gilthead sea bream (Lahnsteiner, 2010). This variation is  
537 largely due to differences in dietary protein sources (Forster and Ogata, 1998), diet formulation,  
538 size, age of species, genetic differences, rearing conditions, and feeding practices (Ruchimat et al.,  
539 1997). Furthermore, a considerable amount of amino acids are produced from the spermatic duct  
540 epithelium (Lahnsteiner et al., 1993, 1994) or by proteolysis of seminal plasma (Ciereszko et al.,  
541 1998), which could both change amino acid profiles in male gametes.

542 Overall, these findings not only add evidence that certain amino acids are essential for  
543 regulating milt biochemistry, but also show that some ejaculate traits may be promoted by amino  
544 acid intake (e.g. proline, histidine, and valine in the present study). Thus, further studies to evaluate  
545 the effect of these supplemented diets on sperm fertilization ability and interlinked early  
546 developmental stages (i.e., egg/embryo to early juveniles) are required. Moreover, new approaches  
547 with high-throughput functional genomics, metabolomics, and proteomics may help to uncover  
548 regulatory roles of these amino acids for gene and protein function. Thus, it would be interesting  
549 to highlight these innovative methods in future attempts in order to expand our knowledge of  
550 amino acid function for fish reproduction, especially for a critically endangered and economically  
551 important catadromous fish species such as European eel.

552

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554

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559

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816 **Figure captions:**

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818

819 **Fig. 1.** Total weight of male European eel, *Anguilla anguilla* fed with four diets (DAN-EX, EH-4,  
820 EH-5, EH-6). Box-and-whisker plot represent the diets for total weight of eels from Week 1 to  
821 Week 9 (A) and the bar chart represents recorded total weight of fish between the diets (B). Values  
822 with different subscripts differ ( $P < 0.05$ ). Results are expressed as mean values  $\pm$  SEM.

823

824 **Fig. 2.** European eel males, *Anguilla anguilla* fed four diets (DAN-EX, EH-4, EH-5, EH-6) were  
825 assessed for milt production from injection Week 5 until Week 9. Milt production was initially  
826 graded from 0 (no milt release) to 5 (flowing milt) and then grouped as “unsuited” (categories 0-  
827 2) or “suited” (categories 3-5; A-E). Regression analysis was then applied to estimate parameters  
828 of a logistic model, where corresponding probabilities varied between 0 for "unsuited" and 1 for  
829 "suited" milt for fertilization procedures (F). Spermatocrit (G) and standardized milt production [g  
830 /100 g eel, (H)] for males fed the four diets was also determined. Values with different subscripts  
831 are significantly different ( $P < 0.05$ ). Results are expressed as mean values  $\pm$  SEM.

832

833 **Fig. 3.** Spermatozoa kinetic traits (A-D) of male European eel, *Anguilla anguilla* fed four diets  
834 (DAN-EX, EH-4, EH-5, EH-6). For sperm motility and velocity, the Time  $\times$  Diet interaction was  
835 non-significant (A, C), therefore the Diet main effect was interpreted for each trait (B, D). Values  
836 with different subscripts are significantly different ( $P < 0.05$ ). Results are expressed as mean values  
837  $\pm$  SEM.

838

839 **Fig. 4.** Photomicrographs of histological sections of testes from selected male European eel,  
840 *Anguilla anguilla*, in different developmental stages. Testis tissues were categorized according to

841 prevalence of different cell types using the spermatogenic maturity index (SMI). Examples include  
842 (A) SMI = 0.54, *male ID 39CD*; (B) SMI = 0.56, *male ID 2CBC*; (C) SMI = 0.71, *male ID 7DE6*;  
843 (D) SMI = 0.80, *male ID DCBA*. Symbols indicate germ cells: Sg = spermatogonia; Sc =  
844 spermatocytes; St = spermatids; Sz = spermatozoa; as well as Ad = adipocytes.

845

846 **Fig. 5.** Score plot (A) and correlation loadings (B) from principal component analysis to study the  
847 correlation between amino acid composition (in % of total amino acids) in the diet and milt of  
848 European eel, *Anguilla anguilla*. The diets (DAN-EX, EH-4, EH-5 and EH-6) were used as  
849 category variables and the locations of the category variables are shown in the loadings plot. All  
850 parameters were weighted by 1/SD and full cross validation was used. In the top panel (A), the  
851 blue squares correspond to milt samples from fish receiving the DAN-EX diet, red circles to milt  
852 from fish receiving the EH-4 diet, green triangles to milt from fish receiving the EH-5 diet, pink  
853 diamonds to milt from fish receiving the EH-6 diet. The mean value for each amino acid in each  
854 diet was used for all milt samples receiving the same diet. These amino acids are marked with “D”  
855 before the amino acid to show that these are the values from the diets.

856

857 **Fig. 6.** Score plot (A) and correlation loadings (B) from principal component analysis to study the  
858 correlation between amino acid composition (in % of total amino acids) in milt samples, sperm  
859 motility [MOT(10), MOT(20), and MOT(30)], and velocity [VLC(10) and VCL(20)] of European  
860 eel, *Anguilla anguilla*. In the top panel (A), the blue squares correspond to milt samples from fish  
861 receiving the DAN-EX diet, red circles to milt from fish receiving the EH-4 diet, green triangles  
862 to milt from fish receiving the EH-5 diet, pink diamonds to milt from fish receiving the EH-6 diet.  
863 The diets (DAN-EX, EH-4, EH-5 and EH-6) were used as category variables and the locations of

864 the category variables are shown in the loadings plot. All parameters were weighted by 1/SD and  
865 full cross validation was used.

866

867 **Table captions:**

868

869

870 **Table 1.** Dietary formulation for the “enhanced” feeds (EH-4, EH-5, EH-6) that were fed to male  
871 European eel, *Anguilla anguilla*.

872

873 **Table 2.** Composition of fatty acids in the diets (EH-4, EH-5, EH-6, n = 2 samples analyzed per  
874 diet) that were fed to male European eel, *Anguilla anguilla*. The commercial feed was DAN-EX  
875 2848 (n = 2 samples analyzed). Diets are presented as average (% of total fatty acids in the feed)  
876 ± standard deviation (SD).

877

878 **Table 3.** Composition of amino acids in the diets (EH-4, EH-5, EH-6; n = 3 samples analyzed per  
879 diet) that were fed to male European eel, *Anguilla anguilla*. The commercial feed was DAN-EX  
880 2848 (n = 2 samples analyzed). Diets are presented as average (% of total amino acids in the feed)  
881 ± standard deviation (SD).

882

883 **Table 4.** Composition of amino acids (percentage of total amino acids) in milt of male European  
884 eel, *Anguilla anguilla* fed different diets (EH-4, EH-5, EH-6, DAN-EX). Commercial feed used  
885 was DAN-EX 2848. Results are presented as average (% of total amino acids in the feed) ±  
886 standard deviation (SEM). Small letters show significant differences ( $P < 0.05$ ) in each amino acid  
887 over the dietary regimes.

**Fig. 1**

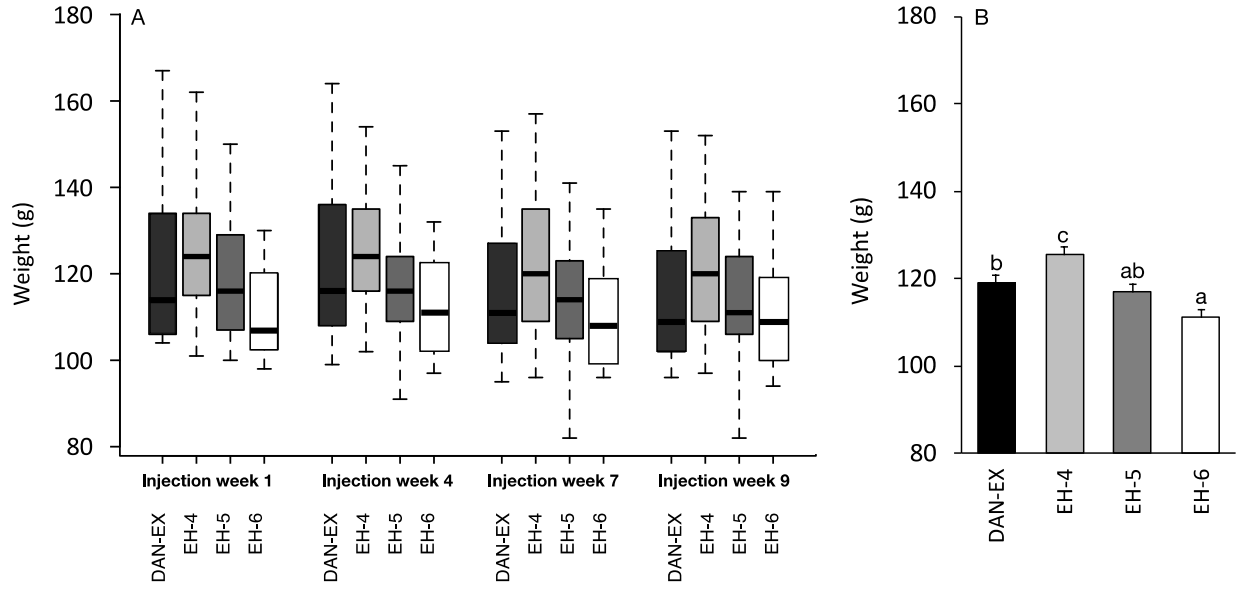
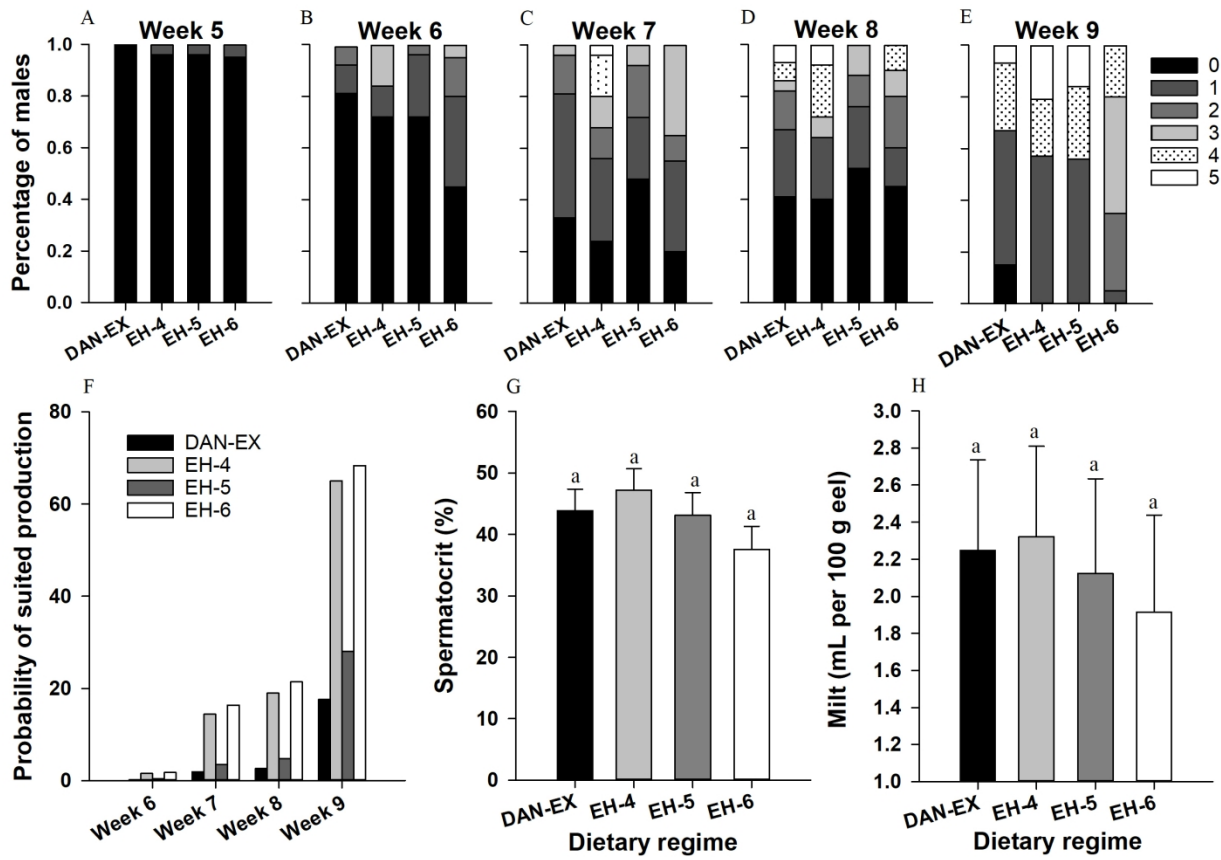
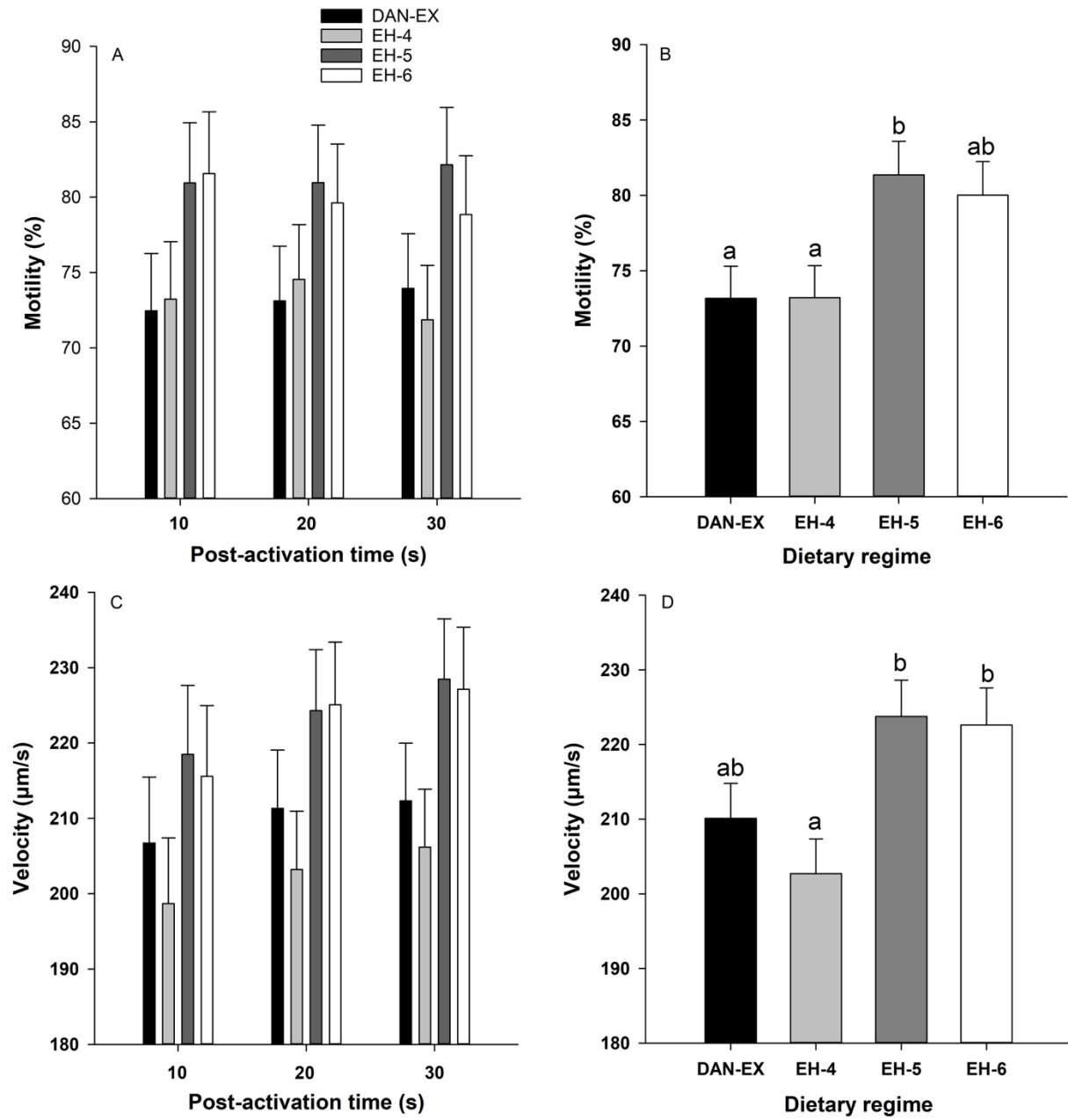


Fig. 2



**Fig. 3**





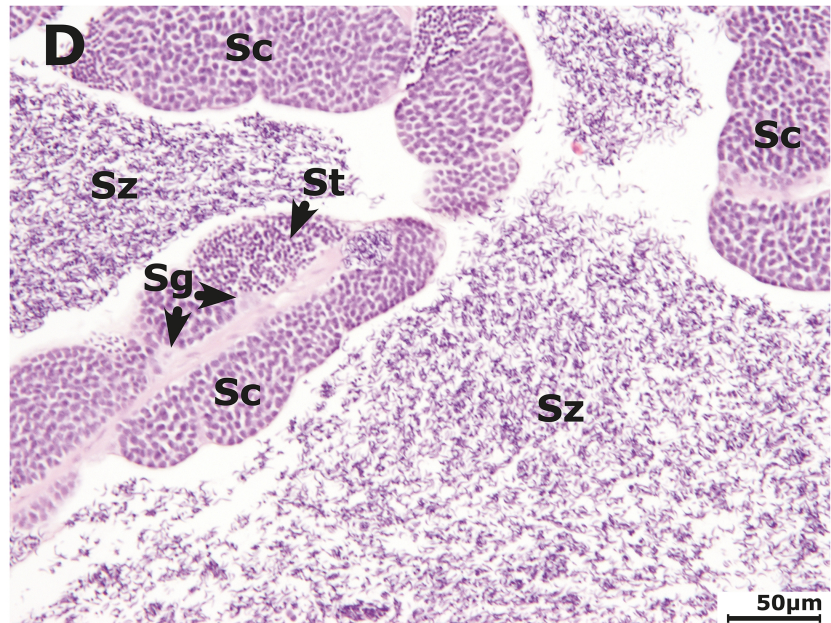
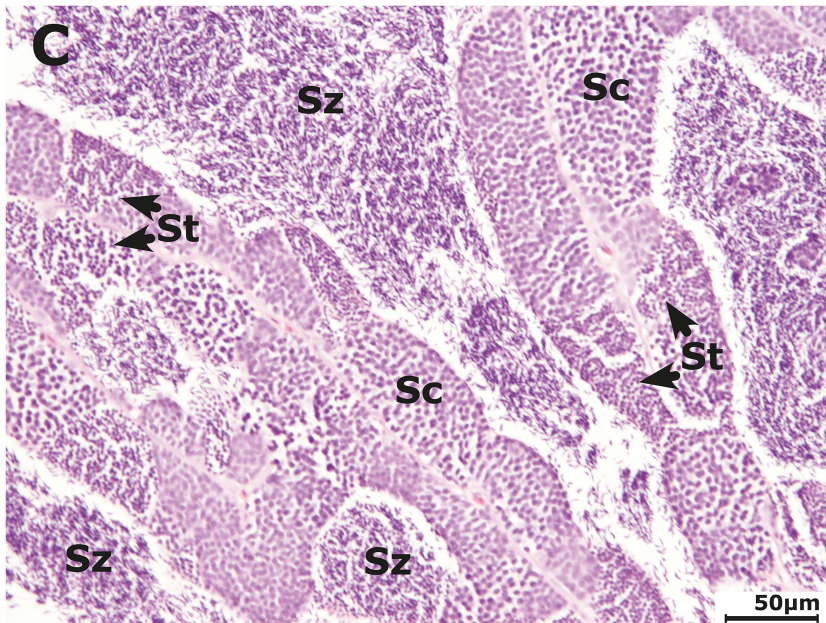
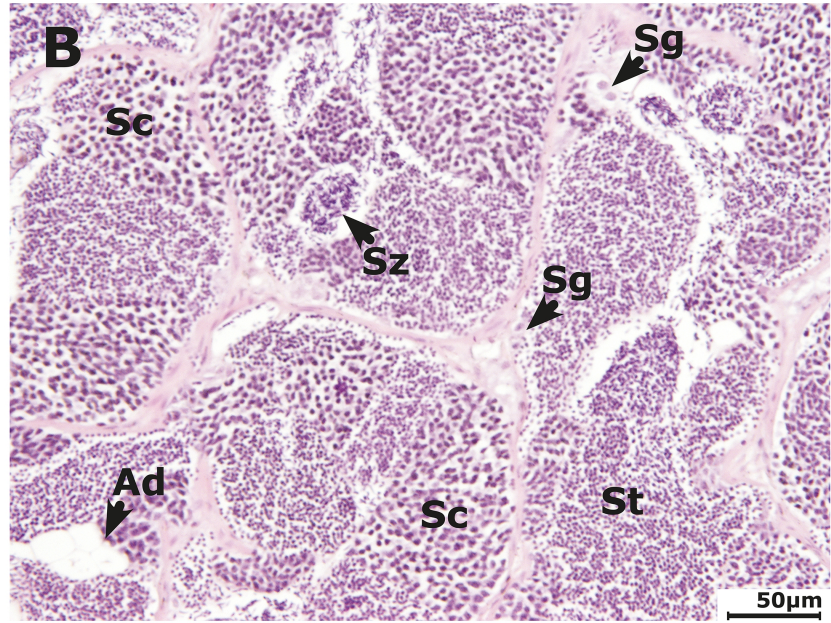
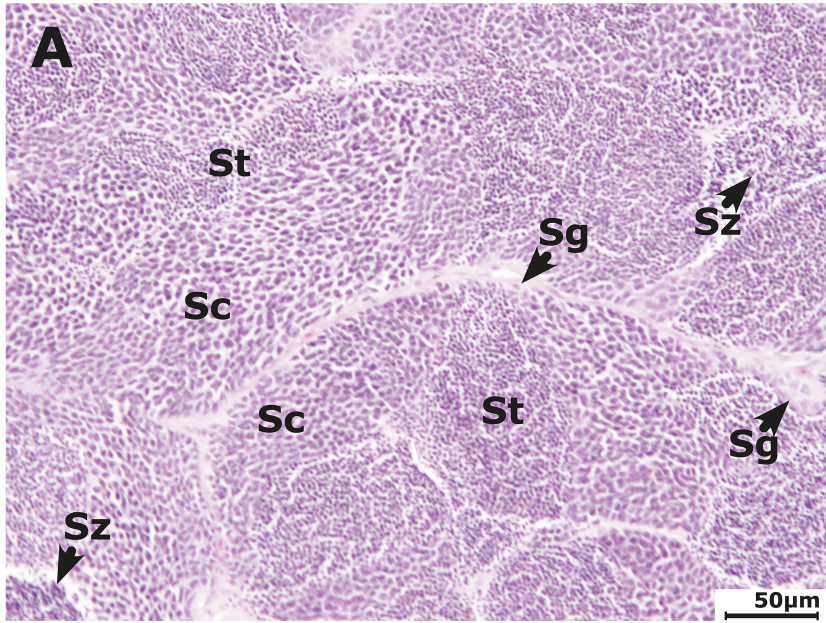
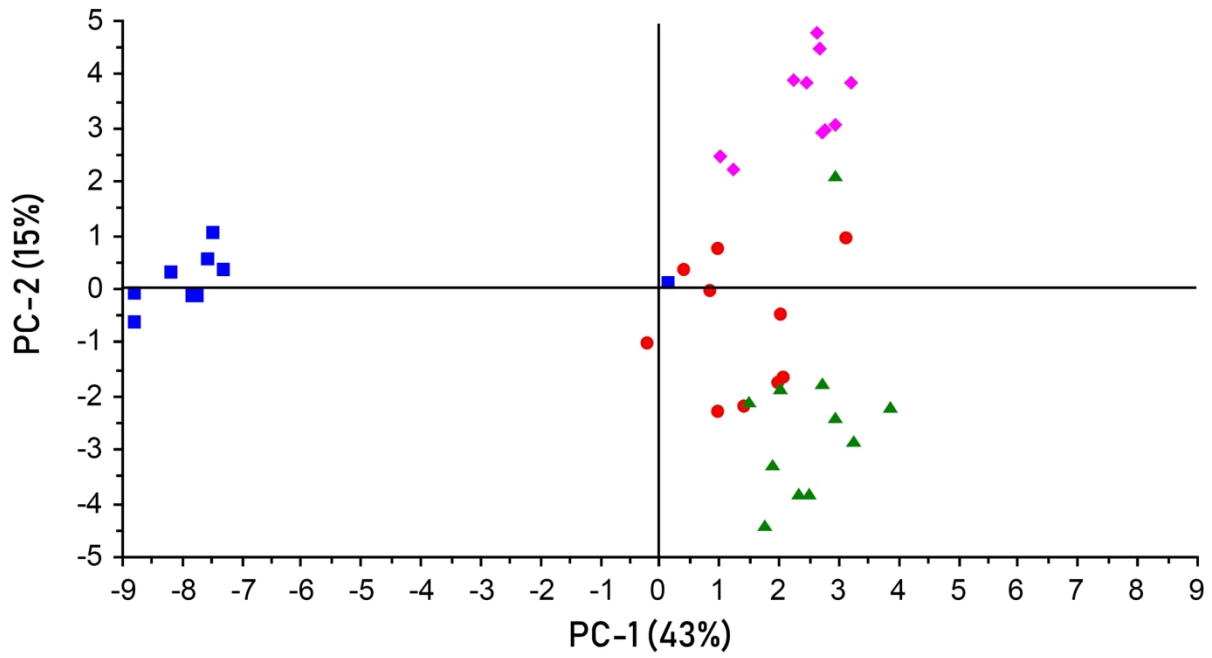




Fig. 5

A Scores



B Correlation Loadings (X)

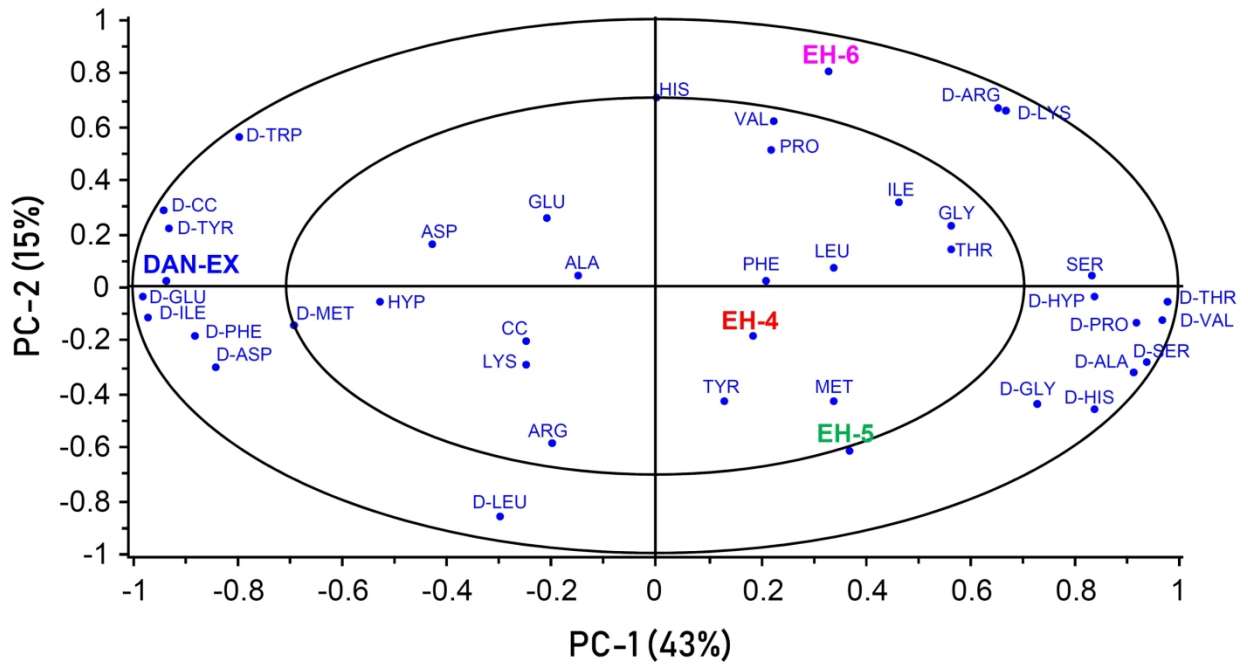
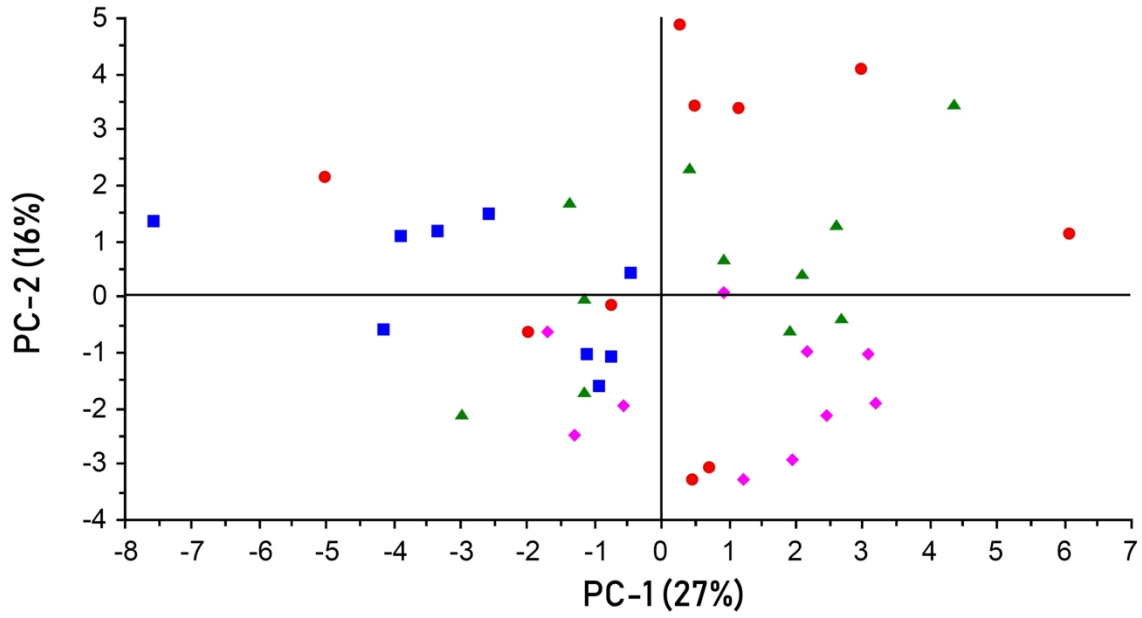


Fig. 6

A Scores



B Correlation Loadings (X)

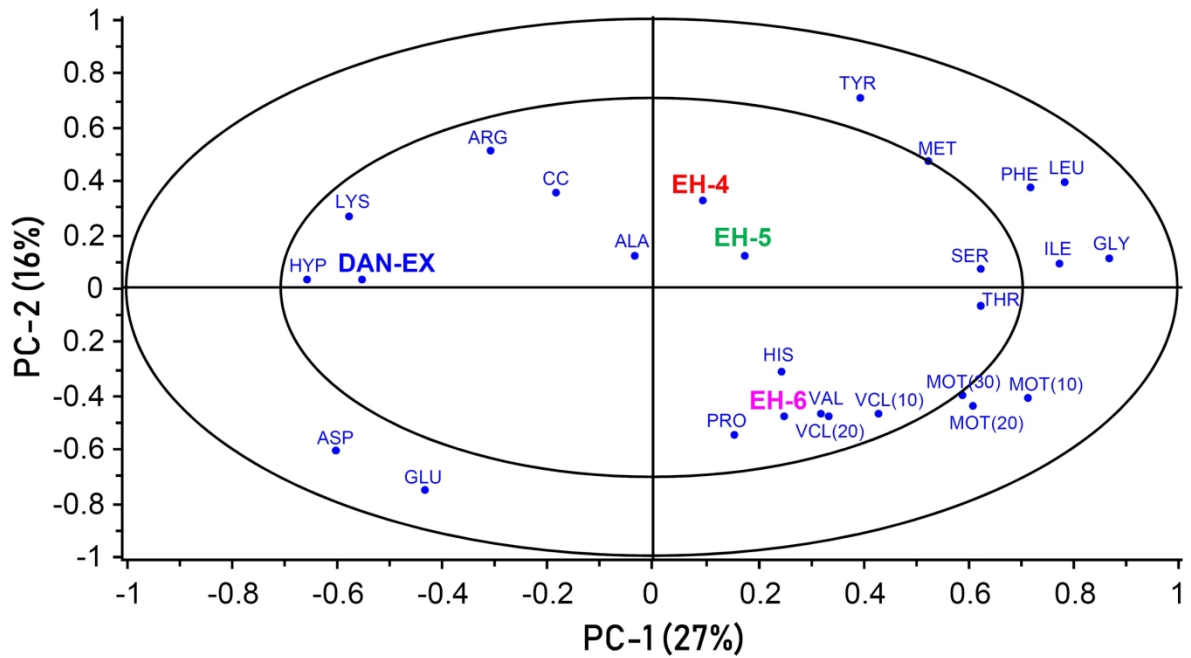


Table 1

| <b>Dietary formulation</b> | <b>EH-4</b> | <b>EH-5</b> | <b>EH-6</b> |
|----------------------------|-------------|-------------|-------------|
| Fish Meal LT               | 53.275      | 55.136      | 53.727      |
| Wheat Gluten               | 9           | 9           | 9           |
| Wheat                      | 22          | 20          | 20          |
| Fish Oil                   | 6.03        | 5.79        | 5.95        |
| Rapeseed Oil               | 5           | 5           | 5           |
| Vevodar                    | 2.35        | 2.39        | 2.37        |
| Premix                     | 2.03        | 1.82        | 1.93        |
| L-Lysine HCl (78%)         | .           | 0.17        | 0.28        |
| DL-Methionine (99%)        | 0.22        | 0.19        | 0.22        |
| L-Threonine (98%)          | .           | 0.19        | 0.23        |
| L-Histidine (74%)          | .           | .           | 0.03        |
| Water change               | -3.423      | -3.204      | -3.083      |
| Lecithin, liquid           | 0.5         | 0.5         | 0.5         |
| DHA Liquid                 | 3           | 3           | 3           |
| L-arginin                  | .           | .           | 0.828       |
| <i>Total</i>               | <i>100</i>  | <i>100</i>  | <i>100</i>  |
| Moisture (%)               | 5.47        | 5.528       | 5.529       |
| Protein - crude (%)        | 48.29       | 49.783      | 49.802      |
| Fat - crude (%)            | 23.032      | 22.979      | 22.976      |
| Ash (%)                    | 8.061       | 8.065       | 7.992       |
| Cellulose - crude (%)      | 0.579       | 0.535       | 0.535       |
| Crude fiber (%)            | 0.579       | 0.535       | 0.535       |

**Table 2.**

| <b>Fatty<br/>Acids</b> | <b>DAN-EX<br/>(n=2)</b> |           | <b>EH-4<br/>(n=2)</b> |           | <b>EH-5<br/>(n=2)</b> |           | <b>EH-6<br/>(n=2)</b> |           |
|------------------------|-------------------------|-----------|-----------------------|-----------|-----------------------|-----------|-----------------------|-----------|
|                        | <b>Mean</b>             | <b>SD</b> | <b>Mean</b>           | <b>SD</b> | <b>Mean</b>           | <b>SD</b> | <b>MEAN</b>           | <b>SD</b> |
| C14:00                 | 5.19                    | 0.01      | 3.44                  | 0.04      | 3.39                  | 0.02      | 3.48                  | 0.01      |
| C16:00                 | 10.79                   | 0.02      | 11.77                 | 0.04      | 11.72                 | 0.03      | 11.85                 | 0.04      |
| C16:1(n – 7)           | 7.35                    | 0.01      | 4.31                  | 0.02      | 4.16                  | 0.03      | 4.22                  | 0.02      |
| C18:00                 | 1.32                    | 0.03      | 3.04                  | 0.03      | 3.22                  | 0.00      | 3.15                  | 0.06      |
| C18:1(n – 9)           | 16.07                   | 0.02      | 19.28                 | 0.19      | 19.02                 | 0.12      | 19.02                 | 0.03      |
| C18:1(n – 7)           | 2.71                    | 0.01      | 2.00                  | 0.02      | 2.14                  | 0.00      | 2.13                  | 0.04      |
| C18:2(n – 6)           | 6.02                    | 0.04      | 12.90                 | 0.13      | 12.91                 | 0.05      | 13.00                 | 0.04      |
| C18:3(n – 3)           | 1.62                    | 0.01      | 1.05                  | 0.02      | 1.11                  | 0.05      | 1.06                  | 0.00      |
| C20:1(n – 7)           | 0.74                    | 0.00      | 0.37                  | 0.00      | 0.38                  | 0.01      | 0.38                  | 0.00      |
| C20:4(n – 6)           | 0.28                    | 0.00      | 4.33                  | 0.04      | 4.41                  | 0.04      | 4.39                  | 0.02      |
| C20:5(n – 3)           | 5.49                    | 0.02      | 3.90                  | 0.05      | 3.96                  | 0.02      | 4.01±                 | 0.02      |
| C22:1(n – 11)          | 13.36                   | 0.03      | 8.29                  | 0.08      | 8.13                  | 0.04      | 8.30                  | 0.05      |
| C22:5(n – 3)           | 0.49                    | 0.00      | 0.56                  | 0.00      | 0.56                  | 0.00      | 0.57                  | 0.00      |
| C22:6(n – 3)           | 5.32                    | 0.04      | 6.61                  | 0.09      | 6.79                  | 0.02      | 6.91                  | 0.02      |
| total n – 3            | 14.07                   | 0.12      | 13.01                 | 0.10      | 13.33                 | 0.12      | 13.59                 | 0.22      |
| total n – 6            | 6.74                    | 0.05      | 17.96                 | 0.17      | 18.12                 | 0.17      | 18.13                 | 0.07      |
| n – 3/n – 6            | 2.09                    |           | 0.72                  |           | 0.74                  |           | 0.75                  |           |
| total SFA              | 17.98                   | 0.07      | 19.18                 | 0.23      | 19.18                 | 0.07      | 19.34                 | 0.13      |
| total MUFA             | 54.22                   | 0.16      | 42.64                 | 0.39      | 42.41                 | 0.33      | 42.66                 | 0.20      |
| total PUFA             | 23.74                   | 0.20      | 34.44                 | 0.53      | 35.24                 | 0.49      | 35.29                 | 0.30      |

**Table 3.**

| <b>Amino<br/>Acids</b> | <b>DAN-EX (n=2)</b> |           | <b>EH-4 (n=3)</b> |           | <b>EH-5 (n=3)</b> |           | <b>EH-6 (n=3)</b> |           |
|------------------------|---------------------|-----------|-------------------|-----------|-------------------|-----------|-------------------|-----------|
|                        | <b>Mean</b>         | <b>SD</b> | <b>Mean</b>       | <b>SD</b> | <b>Mean</b>       | <b>SD</b> | <b>MEAN</b>       | <b>SD</b> |
| Alanine                | 4.47                | 0.01      | 7.01              | 0.66      | 7.21              | 1.04      | 6.37              | 0.70      |
| Arginine               | 4.83                | 0.28      | 7.20              | 0.11      | 6.42              | 1.71      | 10.26             | 1.29      |
| Aspartic acid          | 11.27               | 0.52      | 10.02             | 1.07      | 8.91              | 0.37      | 8.02              | 1.15      |
| Cysteine               | 0.96                | 0.04      | 0.72              | 0.04      | 0.67              | 0.08      | 0.75              | 0.09      |
| Glutamic acid          | 30.22               | 0.12      | 20.38             | 2.40      | 19.71             | 2.10      | 18.91             | 1.28      |
| Glycine                | 4.79                | .         | 5.23              | 0.53      | 6.15              | 0.67      | 5.43              | 0.35      |
| Histidine              | 1.54                | 0.01      | 1.81              | 0.31      | 2.00              | 0.36      | 1.78              | 0.39      |
| Hydroxyproline         | 0.64                | 0.04      | 0.93              | 0.03      | 1.00              | 0.06      | 0.98              | 0.10      |
| Isoleucine             | 3.80                | 0.23      | 3.08              | 0.10      | 2.93              | 0.37      | 2.80              | 0.32      |
| Leucine                | 6.97                | 0.04      | 6.92              | 0.79      | 7.03              | 0.73      | 6.57              | 0.63      |
| Lysine                 | 7.13                | 0.18      | 8.08              | 0.59      | 7.81              | 1.88      | 9.30              | 1.45      |
| Methionine             | 3.56                | 0.21      | 3.03              | 0.43      | 3.35              | 0.25      | 3.20              | 0.40      |
| Phenylalanine          | 5.22                | 0.17      | 4.17              | 0.72      | 4.53              | 0.56      | 4.23              | 0.62      |
| Proline                | 0.34                | 0.03      | 5.06              | 0.07      | 5.27              | 0.48      | 4.79              | 0.17      |
| Serine                 | 3.64                | 0.10      | 4.47              | 0.21      | 4.77              | 0.26      | 4.45              | 0.24      |
| Threonine              | 2.99                | 0.13      | 3.35              | 0.05      | 3.96              | 0.35      | 3.85              | 0.13      |
| Tryptophan             | 0.32                | 0.01      | 0.27              | 0.02      | 0.25              | 0.05      | 0.29              | 0.07      |
| Tyrosine               | 3.43                | 0.22      | 2.65              | 0.11      | 2.69              | 0.23      | 2.83              | 0.22      |
| Valine                 | 3.87                | 0.01      | 5.61              | 0.29      | 5.33              | 1.14      | 5.20              | 0.67      |

**Table 4.**

| <b>Amino<br/>Acids</b> | <b>DAN-EX</b>      |            | <b>EH-4</b>         |            | <b>EH-5</b>        |            | <b>EH-6</b>        |            | <b>P-value</b> |
|------------------------|--------------------|------------|---------------------|------------|--------------------|------------|--------------------|------------|----------------|
|                        | <b>Mean</b>        | <b>SEM</b> | <b>Mean</b>         | <b>SEM</b> | <b>Mean</b>        | <b>SEM</b> | <b>MEAN</b>        | <b>SEM</b> |                |
| Alanine                | 11.54              | 1.20       | 13.30               | 1.09       | 10.65              | 1.09       | 13.02              | 1.04       | 0.28           |
| Arginine               | 12.83 <sup>b</sup> | 0.75       | 11.51 <sup>ab</sup> | 0.68       | 13.00 <sup>b</sup> | 0.68       | 10.12 <sup>a</sup> | 0.65       | 0.01           |
| Aspartic acid          | 6.91               | 0.45       | 5.76                | 0.40       | 6.09               | 0.40       | 6.59               | 0.39       | 0.23           |
| C-C                    | 0.81               | 0.06       | 0.79                | 0.06       | 0.77               | 0.06       | 0.85               | 0.05       | 0.75           |
| Glutamic acid          | 8.29               | 0.63       | 7.42                | 0.57       | 7.67               | 0.57       | 8.87               | 0.55       | 0.28           |
| Glycine                | 5.66               | 0.21       | 6.13                | 0.19       | 6.21               | 0.19       | 6.17               | 0.18       | 0.2            |
| Histidine              | 1.67 <sup>b</sup>  | 0.12       | 1.92 <sup>b</sup>   | 0.11       | 1.26 <sup>a</sup>  | 0.11       | 1.91 <sup>b</sup>  | 0.10       | <0.0001        |
| Hydroxyproline         | 0.09               | 0.02       | 0.04                | 0.02       | 0.01               | 0.02       | 0.01               | 0.02       | 0.05           |
| Isoleucine             | 3.83               | 0.19       | 4.03                | 0.18       | 4.18               | 0.18       | 4.10               | 0.17       | 0.6            |
| Leucine                | 7.06               | 0.29       | 7.49                | 0.26       | 7.29               | 0.26       | 7.04               | 0.25       | 0.6            |
| Lysine                 | 13.80              | 0.67       | 12.89               | 0.60       | 13.33              | 0.60       | 12.42              | 0.58       | 0.44           |
| Methionine             | 1.31 <sup>a</sup>  | 0.10       | 1.57 <sup>ab</sup>  | 0.09       | 1.70 <sup>b</sup>  | 0.09       | 1.36 <sup>a</sup>  | 0.08       | 0.01           |
| Phenylalanine          | 3.12               | 0.15       | 3.37                | 0.13       | 3.18               | 0.13       | 3.27               | 0.13       | 0.6            |
| Proline                | 3.87 <sup>ab</sup> | 0.11       | 3.79 <sup>a</sup>   | 0.10       | 3.96 <sup>ab</sup> | 0.10       | 4.21 <sup>b</sup>  | 0.09       | 0.02           |
| Serine                 | 3.28 <sup>a</sup>  | 0.11       | 3.95 <sup>b</sup>   | 0.10       | 4.16 <sup>b</sup>  | 0.10       | 4.05 <sup>b</sup>  | 0.09       | <.0001         |
| Threonine              | 4.87               | 0.15       | 4.95                | 0.13       | 5.27               | 0.13       | 4.99               | 0.13       | 0.18           |
| Tyrosine               | 4.47               | 0.21       | 4.70                | 0.19       | 4.77               | 0.19       | 4.35               | 0.18       | 0.35           |
| Valine                 | 6.59               | 0.27       | 6.37                | 0.25       | 6.51               | 0.25       | 6.68               | 0.24       | 0.84           |

1 **Supplementary Table 1.** PCA was performed to study the correlation between amino acid  
 2 composition (% of total amino acids) in the diet and milt of European eel, *Anguilla anguilla*,  
 3 where diets (DAN-EX, EH-4, EH-5, EH-6) were used as category variables. The cumulative  
 4 proportion of variance explained and factor loadings for two principle components are displayed.

|                       | PC1         |             | PC2         |             |
|-----------------------|-------------|-------------|-------------|-------------|
| Cumulative proportion | 0.430       |             | 0.580       |             |
| DAN-EX                | -0.225      |             | 0.013       |             |
| EH-4                  | 0.045       |             | -0.074      |             |
| EH-5                  | 0.090       |             | -0.251      |             |
| EH-6                  | 0.080       |             | 0.320       |             |
|                       | <i>Milt</i> | <i>Diet</i> | <i>Milt</i> | <i>Diet</i> |
| Alanine               | -0.035      | 0.219       | 0.017       | -0.127      |
| Arginine              | -0.048      | 0.157       | -0.236      | 0.273       |
| Aspartic acid         | -0.103      | -0.201      | 0.063       | -0.126      |
| C-C                   | -0.059      | -0.225      | -0.082      | 0.111       |
| Glutamic acid         | -0.050      | -0.235      | 0.105       | -0.019      |
| Glycine               | 0.137       | 0.175       | 0.088       | -0.175      |
| Histidine             | 0.001       | 0.201       | 0.283       | -0.182      |
| Hydroxyproline        | -0.127      | 0.235       | -0.023      | -0.020      |
| Isoleucine            | 0.112       | -0.232      | 0.124       | -0.051      |
| Leucine               | 0.083       | -0.071      | 0.027       | -0.352      |
| Lysine                | -0.059      | 0.160       | -0.116      | 0.268       |
| Methionine            | 0.082       | -0.165      | -0.176      | -0.064      |
| Phenylalanine         | 0.052       | -0.210      | 0.008       | -0.078      |
| Proline               | 0.054       | 0.232       | 0.205       | -0.046      |
| Serine                | 0.201       | 0.225       | 0.013       | -0.110      |
| Threonine             | 0.137       | 0.202       | 0.054       | -0.011      |
| Tyrosine              | 0.032       | -0.222      | -0.173      | 0.085       |
| Valine                | 0.054       | 0.220       | 0.249       | -0.053      |

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14 **Supplementary Table 2.** PCA was performed to study the correlation between sperm  
 15 motility/velocity and amino acid composition (% of total amino acids) in the milt of European  
 16 eel, *Anguilla anguilla*, where diets (DAN-EX, EH-4, EH-5, EH-6) were used as category  
 17 variables. The cumulative proportion of variance explained and factor loadings for two principle  
 18 components are displayed.

|                                   | PC1    | PC2    |
|-----------------------------------|--------|--------|
| Cumulative proportion             | 0.27   | 0.43   |
| DAN-EX                            | -0.205 | 0.014  |
| EH-4                              | 0.036  | 0.156  |
| EH-5                              | 0.066  | 0.058  |
| EH-6                              | 0.093  | -0.230 |
| Alanine                           | -0.012 | 0.058  |
| Arginine                          | -0.114 | 0.244  |
| Aspartic acid                     | -0.223 | -0.291 |
| C-C                               | -0.067 | 0.169  |
| Glutamic acid                     | -0.159 | -0.360 |
| Glycine                           | 0.323  | 0.054  |
| Histidine                         | 0.091  | -0.148 |
| Hydroxyproline                    | -0.243 | 0.015  |
| Isoleucine                        | 0.287  | 0.044  |
| Leucine                           | 0.291  | 0.186  |
| Lysine                            | -0.213 | 0.127  |
| Methionine                        | 0.196  | 0.224  |
| Phenylalanine                     | 0.267  | 0.176  |
| Proline                           | 0.059  | -0.260 |
| Serine                            | 0.232  | 0.030  |
| Threonine                         | 0.232  | -0.034 |
| Tyrosine                          | 0.148  | 0.334  |
| Valine                            | 0.118  | -0.225 |
| Motility 10 s (MOT10)             | 0.266  | -0.194 |
| Curvilinear velocity 10 s (VCL10) | 0.125  | -0.230 |
| Motility 20 s (MOT20)             | 0.226  | -0.208 |
| Curvilinear velocity 20 s (VCL10) | 0.160  | -0.224 |
| Motility 30 s (MOT30)             | 0.218  | -0.193 |