

1 **Insect meals in feeds for juvenile gilthead seabream (*Sparus aurata*): effects on growth,**
2 **blood chemistry, hepatic metabolic enzymes, body composition and nutrient utilization**

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31

32 Abstract

33 Alternative and sustainable fish diets are required by modern aquaculture. We investigated the
34 possibility of using insect (*Tenebrio molitor* TM, *Hermetia illucens* HI or *Musca domestica*
35 MD) larvae meals (as 19.5% of the feed formulation) to replace 30% of the in the fish meal
36 (FM) in a gilthead seabream (*Sparus aurata*) feed formulated to contain 65% FM. The feeds
37 were isonitrogenous (ca 57% crude protein of dry matter) isolipidic (ca 17% lipid dry matter)
38 and isoenergetic (ca 22 MJ kg⁻¹ dry matter). To achieve similar energy content among the
39 experimental diets, the fish oil inclusion was adjusted. Fish (average initial weight of 29.5 g)
40 were fed up to apparent satiation three times a day, seven days per week in a 93-days trial.
41 Each diet was assigned to three 500 L tanks with fish density 2 kg m⁻³. Five fish from the initial
42 population and two fish per tank were taken for whole-body composition analysis. At the end
43 of the experimental period, nine fish per treatment were taken for the analysis of plasma
44 metabolites and liver enzyme activities. Growth performance, feed intake, feed conversion and
45 somatic indices of fish fed the different insect meal diets were similar to the FM fish. However,
46 among the insect meal fish groups, the feeding with the TM diet resulted in higher specific
47 growth rate compared to the HI diet (1.57% and 1.51% per day, respectively). The whole-body
48 proximate composition was similar among experimental groups. Fish fed HI had the lowest fat
49 retention (57.0% compared to 69.3 – 74.2%). Additionally, the HI group had also lower dry
50 matter and energy retention (30.5% and 36.6%, respectively) compared to the FM group
51 (33.8% and 41.5%, respectively). The whole-body saturated and mono-unsaturated fatty acids
52 content was similar to all the experimental groups. Fish fed diets higher in fish oil (FM and HI)
53 had higher eicosapentaenoic, docosahexaenoic and total ω-3 poly-unsaturated fatty acids
54 content. Whole-body amino acid composition was similar among all experimental groups,
55 while the amino acid retention exhibited significant differences. The plasma metabolites and
56 enzyme activities as well as the hepatic lipogenic enzyme activity were not affected by the
57 different diets. Fish fed the HI diet exhibited higher liver alanine aminotransferase (ALT)
58 activity in comparison to the TM group. Overall, this study shows that FM can be successfully
59 replaced by TM, HI or MD meals in 30% by weight in the diets of gilthead sea bream.
60 Comparing insect meals, HI meal was inferior in terms of growth performance and dry matter-
61 fat retention compared to TM and MD, respectively.

62

63 **Keywords:** dietary protein sources, amino acid catabolism, lipogenic enzymes, fatty acid
64 analysis, amino acid deposition, plasma biochemistry

65

66 **Abbreviations:** Fish meal, FM; *Tenebrio molitor*, TM; *Hermetia illucens*, HI; *Musca*
67 *domestica*, MD; Acid Detergent fiber, ADF; Specific Growth Rate, SGR; Feed Conversion
68 Ratio, FCR, alanine aminotransferase, ALT; aspartate aminotransferase, AST.

69

70 **1. Introduction**

71 Insect meals can be included in aquafeeds as valuable sources of high-quality protein, thereby
72 reducing the reliance on ingredients, such as fish meal, derived from overexploited natural resources.
73 Insect larvae have high nutritional value (Poshadri et al., 2018; Rumpold and Schlüter, 2014), they can
74 be produced in a short time due to their short life cycle (Hua et al., 2019) and they can bioconvert and
75 biotransform organic matter (Fowles and Nansen, 2019; Gasco et al., 2020) with low feed conversion
76 ratio (Ooninx et al., 2015). Therefore, insect larvae production could be performed efficiently next to
77 agri-food manufacturing facilities, by converting side-streams into products of high nutritional value
78 (Smetana et al. 2019) which contribute to limiting environmental degradation (Van Huis, 2013) in the
79 frame of circular economy (Madau et al., 2020).

80 The study of insects and insect meals in aquafeeds has a history that stretches back over 30 years
81 (Bodari and Shepard 1981), but most of the work on farmed fish has been carried out on freshwater
82 species (e.g. Alegbeleye et al., 2012; Barroso et al 2014; Sogbesan, 2014; St-Hilaire et al., 2007). More
83 recently, research has been conducted on incorporating insects and insect meals into feeds for
84 anadromous species, such as Atlantic salmon (*Salmo salar*) (Belghit et al 2018, 2019; Bruni et al 2020),
85 marine and freshwater crustaceans, such as Pacific white-leg shrimp (*Litopenaeus vannamei*) and Baltic
86 prawn (*Palaemon adspersus*) (Rahimnejad et al, 2019; Mastoraki et al, 2020b) and some marine fish
87 species, including the gilthead sea bream (*Sparus aurata*) (Piccolo et al 2017; Antonopoulou et al 2019;
88 Randazzo et al 2021; Pulido-Rodriguez et al 2021).

89 In this framework, the aim of the present study is the evaluation of the effect of 30% fish meal
90 substitution with three different insect meals derived from *Tenebrio molitor*, *Hermetia illucens* or
91 *Musca domestica* larvae on growth performance, nutrient utilization and intermediary metabolism of
92 gilthead sea bream *Sparus aurata*. Although insect meal diets have been extensively studied,
93 comparative studies, investigating the effects of different insect meals in the same trial, are scarce
94 (Józefiak et al., 2019; Melenchón et al., 2020; Fabrikov et al., 2020). Innovation-wise, the study herein
95 is the first one assessing the effects of insect larvae meal from housefly *Musca domestica* on gilthead
96 sea bream.

97 **2. Materials and methods**

98 The feeding trial was conducted at the Institute of Marine Biology, Biotechnology and
99 Aquaculture (IMBBC) of the Hellenic Centre for Marine Research (Heraklion, Greece) by accredited
100 scientists of the Federation of European Laboratory Animal Science Associations (FELASA). The
101 experiments were authorized by the ethics committee of the region of Crete, Greece (license No
102 255,340) and were conducted following the guidelines established by the EU Directive 2010/63/EU.

103 2.1. Composition of diets and experimental conditions

104 The feeds were formulated to contain fish meal (FM) as the main protein source in the control
105 feed (FM as 65% of the formulation) and insect meals from larval *Tenebrio molitor* (TM; full-fat),
106 *Hermetia illucens* (HI; defatted) and *Musca domestica* MD; full-fat) used to replace 30% of the FM, by
107 including them as 19.5% of the formulation (Tables 1 and 2). To achieve a similar amount of essential
108 amino acids between the different diets, crystalline DL-methionine and L-lysine were incorporated. In
109 addition, to ensure a similar protein and energy content between the diets, the inclusion of the different
110 ingredients, namely fish oil, wheat flour and wheat gluten, were adjusted. A mincing machine with 4
111 mm die was used for the formation of pellets. Finally, the diets were oven-dried at 40°C for 24 h and
112 were stored in a freezer.

113 Proximate compositions of insects and diets were determined by using standard methods of
114 analysis (AOAC, 1990). Dry matter was determined by drying at 90°C until constant weight and ash
115 by incineration at 700°C for 7 h. Crude fat was determined according to Folch et al. (1957) using
116 extraction with chloroform-methanol-butylated hydroxytoluene (2:1 v/v + 0.01% w/v BHT). Energy
117 was measured using a bomb calorimeter (6300, Parr Instrument Company, St. Moline, Illinois, USA).
118 Crude fiber was determined by defatting the samples with petroleum ether and sequential boiling with
119 0.13 mol l⁻¹ H₂SO₄ and 0.23 mol l⁻¹ KOH using Fibretherm (C. Gerhardt GmbH & Co., Königswinter,
120 Germany). To quantify lignin, cellulose, chitin and indigestible nitrogen (nitrogen linked to cell-walls
121 or chitin) content of the diets, the analysis of the ash-free acid detergent fiber (ADF) was performed by
122 boiling with 1N H₂SO₄ + 2% Cetyl trimethylammonium bromide (CTAB) using Fibretherm, and
123 subtracting the ash content (Bernard et al., 1997; Finke, 2007; Goering and Van Soest, 1979). Crude
124 protein was defined with a nitrogen analyzer (FP-528, Leco corporation, St. Joseph, Michigan, USA)
125 by multiplication of the nitrogen content by 6.25. Crude protein was determined, according to Dumas's
126 method, using a nitrogen analyzer (FP-528, Leco corporation, St. Joseph, Michigan, USA) by
127 multiplication of the nitrogen content by 6.25. In addition, the nitrogen linked to ADF (ADIN) was
128 determined according to Goering and Van Soest (1979) using a nitrogen analyzer. Diets' protein content
129 was adjusted by subtracting the ADIN from the total nitrogen content and then multiplying it by 6.25.

130 The amino acid composition of the diets was analyzed after acid hydrolysis (6 N HCl, 11 °C, 24
131 h), and derivatization by AccQ-Tag™ Ultra according to the amino acid analysis application solution
132 (Waters Corporation, Milford, MA, U.S.A.). DL-Norvaline (Sigma) 2.5 mM was used as an internal
133 standard. UPLC was performed on an Acquity system (Waters Corporation, Milford, MA, U.S.A.)
134 equipped with PDA detector and the detection wavelength was set at 260 nm. The column used was
135 BEH C18 column (100 mm × 2.1 mm i.d., 1.7 µm) from Waters. The flow rate was 0.7 ml min⁻¹ and
136 the column temperature was kept at 55 °C. Peak identification and integration were performed by the
137 software Empower v.2.0 (Waters Corporation, Milford, MA, U.S.A.) using Amino Acid Standard H
138 (Thermo Scientific Pierce) as an external standard. Tryptophan was not quantified due to its

139 susceptibility to acid hydrolysis, whereas cysteine reacts with cysteine to form cystine. Moreover,
140 during acid hydrolysis procedure, asparagine is converted to aspartate and glutamine to glutamate, so
141 the reported values for these amino acids (Asx and Glx) represent the sum of both amino acids.

142 For the analysis of fatty acids, lipid samples were saponified with a NaOH-methanol solution
143 and the resulting fatty acids were methylated by a Boron trifluoride-methanol solution, according to
144 AOCS (1989). The fatty acid methyl esters (FAMES) were extracted with iso-octane and analyzed
145 through a Shimadzu GC-2010 gas chromatograph (Shimadzu Corporation, Japan), equipped with a
146 flame-ionization detector (GC-FID) and a SP-2330 capillary column (30 m x 0.25 mm i.d. x 0.20 µm
147 film thickness (Supelco Inc., Bellefonte, Pennsylvania, USA). Helium was used as carrier gas at 2 ml/min
148 constant flow; the split ratio was 1:50 and the injected volume 1.0 µl. The thermal gradient was 100°C
149 to 160°C at 10 °C min⁻¹, 160°C to 220°C at 3 °C min⁻¹ and kept for 5 min, and lastly, 220°C to 250°C
150 at 10°C min⁻¹ and kept for 5 min. The injector and detector temperature were maintained at 260°C and
151 280°C, respectively. Fatty acids were identified by comparison with a known standard mixture (Supelco
152 37 Component FAME Mix). FAME contents were expressed as a percentage (%) of total FAMES basis.

153 Juvenile gilthead sea breams were provided from the IMBBC hatchery. After a light anaesthesia
154 (phenoxyethanol, 150 ppm), 360 fish were individually weighed (29.5 ± 0.7 g) and were randomly
155 divided into 12 open circulation indoor tanks (500 L). Five fish were sacrificed by anaesthesia overdose
156 (phenoxyethanol, 500 ppm) and were stored at -20°C for a whole-body proximate composition analysis.
157 The feeding trial started the following day. The water temperature throughout the experimental period
158 was 19.9 ± 0.1 °C, salinity was 35 ppt, oxygen saturation was constantly over 80%, and the photoperiod
159 was 12 h light/12 h dark. Diets were assigned to triplicate groups and fish were fed by hand until
160 apparent satiation, three times a day, seven days a week for three consecutive months. Pellets that
161 remained unconsumed were siphoned daily and dried to determine feed intake.

162 **2.2. Growth performance and somatic indices**

163 At the end of the experimental period, the sampling was carried out after a 24 h fast in order to
164 reduce handling stress and ensure an empty gastrointestinal tract. Fish were lightly anaesthetised,
165 individually weighed and measured accordingly. Three random fish per tank were sacrificed by
166 anaesthesia overdose (phenoxyethanol, 500 ppm). The liver, viscera (including liver and visceral fat)
167 and mesenteric fat were weighed, gut length was measured, while blood and liver were collected for
168 further analysis (see subchapter 2.3). Additionally, two fish per tank at the end of the experiment were
169 sacrificed by anaesthesia overdose and stored at -20°C for a whole-body proximate composition
170 analysis. Fish for the whole-body analysis were frozen, chopped, lyophilized (Telstar Cryodos,
171 Terrassa, Spain) and homogenized (Retsch ZM200, Haan, Germany). Proximate composition, amino
172 acid composition and fatty acid profile were analyzed, as described in section 2.1.

173 The following growth performance and somatic indices were calculated:

174 Survival (%) = 100 x final/ initial number of fish
 175 Weight gain (WG, %) = 100 x (FBW (final body weight, g) – IBW (initial body weight, g)) / IBW
 176 Specific growth rate (SGR, % day⁻¹) = 100 x ln (FBW/ IBW) / number of days
 177 Daily feed intake (DFI, % body weight day⁻¹) = number of days' x total dry feed intake (g) x 100 /
 178 ((IBW + FBW) x 0.5)
 179 Feed conversion ratio (FCR) = total dry feed intake (g) / weight gain (g)
 180 Condition factor (CF) = 100 x body weight (g) x total length⁻³ (cm)
 181 Hepatosomatic index (HSI) = 100 x liver weight (g) / body weight (g)
 182 Mesenteric fat index (MFI) = 100 x perivisceral fat weight (g) / body weight (g)
 183 Viscerosomatic index (VSI) = 100 x viscera weight (g) / body weight (g)
 184 Relative gut length (RGL) = gut length (cm) / fish total length (cm)
 185 Nutrient retention efficiency (%) = 100 x (final nutrient quantity in the body (g, wet basis) - initial
 186 nutrient quantity in the body (g, wet basis)) / nutrient consumed (g, dry basis),
 187 where nutrient can be dry matter, protein, lipid, energy, ash or amino acids.
 188

189 **2.3. Plasma and liver enzyme activities**

190 Blood samples from three fish per tank were taken from the caudal vein with heparin coated
 191 syringes and were stored on crushed ice until all samples were collected. Plasma was removed by
 192 centrifugation at 3,500 g for 15 min and stored at -80°C until analysis. The quantification of plasma
 193 glucose, cholesterol, triglycerides, phospholipids and lactate was performed by enzymatic colorimetric
 194 methods using commercial kits (BIOSIS Biotechnological Applications L.T.D Greece and Spinreact
 195 S.A.U., Spain). Activities of plasma alanine aminotransferase (ALT, EC 2.6.1.2) and aspartate
 196 aminotransferase (AST, EC 2.6.1.1) were determined kinetically at 340 nm using commercial kits
 197 according to the instructions of the manufacturer (BIOSIS Biotechnological Applications L.T.D,
 198 Greece).

199 After sampling, the three livers per tank were stored at -80°C. A frozen sample of each liver was
 200 homogenized using 10 volumes of buffer (4 °C) which contained 30 mM HEPES, 0.25 mM saccharose,
 201 0.5 mM EDTA, 5 mM K₂HPO₄ and 1 mM DTT (pH 7.4). After centrifugation at 1,000 g for 10 min at
 202 4°C, supernatants were sonicated for 1 min. Following a second centrifugation at 15,000 g for 20 min
 203 at 4°C, supernatants were collected for the assessment of enzyme activity. Activities of ALT and AST
 204 were measured using commercial kits (Spinreact S.A.U., Spain). Glutamate dehydrogenase (GDH, EC
 205 1.4.1.2) activity was assayed as described by Bergmeyer (1974) using 10 mM of L-glutamic acid to a
 206 reaction mixture containing 175 mM tris (pH 8.5), 100 mM semicarbazine, 1.1 mM NAD, 1mM ADP
 207 and 5 mM L-Leucine.

208 For the enzymatic activity of glucose-6-phosphate dehydrogenase (G6PD; EC 1.1.1.49), malic
209 enzyme (ME; EC 1.1.1.40) and fatty acid synthetase (FAS; EC 2.3.1.38), liver samples were
210 homogenized using 5 volumes of buffer (4°C) which contained 20 mM Tris-HCl, 0.25 M sucrose, 2
211 mM EDTA, 0.1 M sodium fluoride, 0.5 mM phenyl methyl sulphonyl fluoride (PMSF) and 10 mM β -
212 mercaptoethanol (pH 7.4). The homogenate was centrifuged at 30,000 g for 20 min at 4°C. G6PD
213 activity was measured according to Bautista et al. (1988) by adding 20 mM glucose-6-phosphate to a
214 reaction mixture containing 1000 mM Tris-HCL buffer (pH 7.8), 200 mM MgCl₂, and 10 mM NADP.
215 ME activity was measured according to Ochoa (1955) by adding 15 mM L-malate to a reaction mixture
216 containing 500 mM glycyl-glycine, 50 mM MgCl₂, and 10 mM NADP (pH 7.4). FAS activity was
217 assayed by following the method described by Chang et al. (1967) as modified by Chakrabarty and
218 Leveille (1969) by addition of 0.6 mM malonyl-CoA in a reaction mixture of 100 mM potassium
219 phosphate buffer (pH 6.5), 0.1 mM NADPH, and 0.025 mM acetyl-CoA. Enzyme activities were
220 determined at 37°C using a Multiskan GO microplate reader (model 51119200; Thermo Scientific,
221 Nanjing, China). Enzyme activities were expressed as nmol of substrate hydrolyzed per min (mU) and
222 per mg of protein. Extracts' protein concentration was determined using Sigma-Aldrich protein assay
223 kit (B6916), with bovine serum albumin as standard, according to Bradford (1976).

224 **2.4. Statistical analysis**

225 Data were tested for normality with the Kolmogorov-Smirnov test and for equality of variances
226 with the Levene's test. One-way analysis of variance (ANOVA) was performed to determine whether
227 significant differences existed among dietary treatments. Data which did not follow the ANOVA
228 assumptions were analyzed by Kruskal Wallis' tests. The results were considered statistically
229 significant at $p < 0.05$ and individual means were compared using the Tukey's test. Correlational
230 analyses were performed using the Spearman correlation. All statistical analyses were carried out using
231 SigmaStat 3.5 (Systat Software, Inc., San Jose, California, USA).

232 **3. Results**

233 **3.1. Growth performance**

234 Survival was similar in all experimental groups, and fish quadrupled their initial weight (29.5 g)
235 during the 93-day feeding trial (Table 3). Fish fed with insect meal diets had similar final body weight
236 compared to the fish meal group ($p > 0.05$, Table 3). However, the inclusion of TM resulted in
237 significantly higher weight gain and SGR ($331.3 \pm 7.5\%$ and $1.57 \pm 0.02\%$ per day, respectively)
238 compared to the HI inclusion ($305.9 \pm 5.2\%$ and $1.51 \pm 0.01\%$ per day, respectively). Insect meals
239 inclusion did not affect the daily feed intake (1.36-1.46% of body weight per day) as well as the feed
240 conversion ratio (1.04-1.12) ($p > 0.05$, Table 3).

241 Somatic indices (condition factor, hepatosomatic index, viscerosomatic index, mesenteric fat
242 index and the relative gut length) were similar to all fish groups ($p > 0.05$, Table 3).

243 **3.2. Whole-body proximate composition and nutrient retention**

244 The whole-body contents of dry matter, protein, fat, ash and energy were not affected by the
245 partial substitution of fish meal with different insect meals ($p > 0.05$, Table 4). Regarding whole-body
246 amino acid profiles, no statistically significant differences were observed in the essential and non-
247 essential amino acid content of fish fed with the different experimental diets ($p > 0.05$, Table 4).

248 Fatty acid composition of the experimental diets (Table 2) affected significantly the fish whole-
249 body fatty acid content, with differences being observed in the content of 15 of the 19 individual fatty
250 acid methyl esters studied (Table 5). A significant positive correlation between whole-body fatty acid
251 content and feed was observed for 11 out of 19 fatty acids analyzed (Table 5). Additionally, there was
252 also a significant positive correlation between dietary fish oil and whole-body fatty acid content for 11
253 fatty acids. The HI group had generally similar fatty acid profile compared to fish fed diet FM with the
254 only differences being observed in lauric acid (C12:0) and linolenic acid (C18:2 ω -6), that presented
255 higher contents in HI fed fish. Total saturated fatty acid (SFA) and mono-unsaturated fatty acid (MUFA)
256 content was not affected by the different dietary treatments (22.3-24.4% and 43.9-47.4%, respectively).
257 Fish fed with the FM and HI diets had higher poly-unsaturated fatty acid (PUFA) content compared to
258 the fish fed with TM and MD diets (32.8-33.5% compared to 27.5-28.1%), which was driven by the
259 significantly higher ω -3 content (17.8-18.6% compared to 12.0-13.5%). The highest oleic acid (C18:1
260 ω -9) of the TM diet led to a significantly higher oleic acid content in the fish fed with TM ($38.0 \pm$
261 0.8%). The ω -3 PUFA, EPA (eicosapentaenoic acid C20:5 ω -3) and DHA (docosahexaenoic acid C22:6
262 ω -3) content of the fish increased over time (higher content than the initial fish) in all experimental
263 groups. However, fish fed diets with higher fish oil inclusion (FM and HI) had a more pronounced
264 increase.

265 Protein retention (Table 6) was similar to all experimental groups (29.3-31.3%; $p > 0.05$). Fish
266 fed with HI had significantly lower dry matter, fat and energy retention compared to the FM group. In
267 addition, the HI group had significantly lower fat retention (57.0%) compared to the fish fed with TM
268 and MD (73.2% and 69.3%, respectively). Dry matter, energy and fat retention were negatively
269 correlated with the crude fiber content of the diets ($r = -0.777$, -0.820 and -0.799 , respectively; $p <$
270 0.05). Moreover, a negative correlation was found between fat retention and diet's ADF content ($r = -$
271 0.756 , $p < 0.05$). Ash retention was significantly lower in fish fed with the FM diet (22.4%) and was
272 positively correlated with SGR ($r = 0.699$, $p < 0.05$).

273 The fish meal substitution with insect meals resulted in significant differences in the amino acid
274 body retention/deposition for 10 out of 17 amino acids measured (Table 6). Within the 10 amino acids
275 that exhibited significant differences in body retention/deposition, seven of them (arginine, methionine,

276 phenylalanine, cysteine, glycine, proline and tyrosine) correlated negatively ($p < 0.05$) with the
277 corresponding dietary amino acid content (i.e., the higher the amino acid content in diet, the lower the
278 retention/deposition).

279

280 **3.3. Plasma metabolites and liver enzyme activities**

281 Plasma glucose, cholesterol, triglycerides, phospholipids, and lactate levels were not affected
282 by the different dietary treatments (Table 7). In addition, plasma ALT and AST activities were similar
283 to all the experimental groups (Table 7).

284 Liver ALT activity was significantly lower in fish fed TM (368.9 ± 5.4 mU mg protein⁻¹)
285 compared to fish fed with HI (457.9 ± 5.5 mU mg protein⁻¹; Table 8). The partial substitution of fish
286 meal with the different insect meals did not affect liver AST and GDH activities. No significant
287 differences were observed in the activities of the multi-enzyme complex of FAS and the activities of
288 ME and G6PD among the different experimental groups (Table 8).

289 **4. Discussion**

290 **4.1. Growth performance**

291 Similar daily feed intake for all the experimental groups with 30% fish meal substitution with
292 insect meals was well accepted by the fish, complying with the known nutritional requirements and
293 diets of gilthead sea bream (Tibaldi and Kaushik, 2005; Wilson, 2003). Insect meal fed fish had similar
294 growth performance compared to the fish fed the fishmeal diet. Piccolo et al. (2017) have reported an
295 improvement of SGR and FCR with 25% TM inclusion in the diet of gilthead sea bream (50% FM in
296 the control diet, 105.2 g initial body weight). In our trial, despite the similar growth performance with
297 the FM group, the inclusion of TM resulted in higher weight gain and SGR compared to HI without
298 affecting FCR and feed intake. The difference observed could be explained by the higher crude fiber
299 and ADF of the HI diet compared to the TM diet (for HI 3.9 and 8.8, respectively; for TM 2.5 and 5.6,
300 respectively) which might decrease nutrient digestibility. In addition, the HI diet had the lowest adjusted
301 crude protein content which could have negatively affected growth performance. Fabrikov et al. (2020)
302 by using TM or HI to replace 15% or 30% of fish meal in the diet of gilthead sea bream (5.1-10.9%
303 inclusion of insect meal, 36.8% FM in the control diet, 6.4 g initial body weight) have reported similar
304 growth performance and FCR. However, these experimental diets had a lower insect meal inclusion
305 compared to the present study and similar crude fiber content (1.3-1.9%).

306 *Tenebrio molitor* has been extensively used as a fish meal replacement in literature. In European
307 sea bass (*Dicentrarchus labrax*), Gasco et al. (2016) have not observed any negative effects on growth
308 performance and feed conversion with 36% fish meal substitution with full fat TM (25% inclusion),
309 while Mastoraki et al. (2020a) have reported an increased FCR with 30% fish meal substitution (19.5%

310 inclusion). It is known that growth performance and feed efficiency are not affected when up to 28% of
311 TM was included in the diets of rainbow trout (Jeong et al., 2020) and yellow catfish (*Pelteobagrus*
312 *fulvidraco*; Su et al., 2017). When defatted TM is used, a complete replacement of FM is possible in
313 larger rainbow trout (78.3 g, 20% inclusion; Chemello et al., 2020) without negative effects on growth
314 performance. However, in smaller rainbow trouts (5.01 g; Rema et al., 2019) and juvenile red seabreams
315 (*Pagrus major*; Ido et al., 2019), the inclusion of 25% and 65% of TM replacing fish meal has
316 completely improved their growth performance.

317 Regarding the substitution of fish meal with partially defatted *Hermetia illucens*, our results
318 agree with those of Abdel-Tawwab et al. (2020) and Mastoraki et al. (2020a) reporting 50% and 30%
319 replacement of fish meal (14.8% and 19.5% inclusion, respectively) in European sea bass without any
320 effect on growth performance. Furthermore, HI diets have performed in the same fashion as fish meal
321 diets in rainbow trout (21% inclusion; Cardinaletti et al., 2019) in Nile tilapia (30% inclusion to
322 complete fish meal replacement; Muin et al., 2017), in grass carp (*Ctenopharyngodon idellus*, 13.4%
323 inclusion; Lu et al., 2020), in Eurasian perch (up to 60% inclusion; Stejskal et al., 2020), and in Siberian
324 sturgeon (up to 18.5% inclusion; Caimi et al., 2020).

325 To date, this is the first study that reports the effects of *Musca domestica* inclusion in the diet
326 of gilthead sea bream. In the present study, no differences were observed in growth performance and
327 feed conversion when 30% of fish meal was substituted with MD. Successful fish meal substitution
328 with MD has also been achieved in barramundi (*Lates calcalifer*, 10% inclusion; Lin & Mui, 2017),
329 *Heteroclaris* (*Clarias x Heterobranchus*, 15-50% inclusion; Ekelemu, 2015; Omoruwou & Edema,
330 2011; Sogbesan, 2014), and in Nile tilapia (33-68% inclusion; Ezewudo et al., 2015; Ogunji et al., 2007;
331 Wang et al., 2017).

332 Regarding the somatic indices, the present study observed no differences in the hepatosomatic
333 index (values <2%). Liver is the primary metabolic tissue, and hepatosomatic indices exceeding 2%
334 can indicate an impairment of glucose and/or fat metabolism or vitamin deficiency (Chemello et al.,
335 2020). Fish from the different dietary groups of this study had similar viscerosomatic and mesenteric
336 fat indices. Dietary fat has been reported to affect fat storage in the viscera and liver of fish (Huang et
337 al., 2016). Furthermore, when fish meal is substituted with insect meals, the reduction of dietary ω -3
338 fatty acids and the increase of dietary linoleic and linolenic acids can lead to an imbalance in the ω -
339 3/ ω -6 fatty acid ratio which can result in increased liver fat deposition (Mikołajczak et al., 2020; Xu et
340 al., 2020). In the present study, the dietary fat was similar among the experimental diets, not affecting
341 visceral and liver fat deposition. The lack of differences observed in the fat deposition of the liver and
342 perivisceral cavity was further supported by similar activities of the liver lipogenic enzymes between
343 the experimental groups. Contrary to the study of Piccolo et al. (2017) who have reported significantly
344 higher relative gut length in gilthead sea bream fed diets with 50% inclusion of TM, no differences
345 were observed herein when fish meal was substituted at lower level (19.5%) with different insect meals.

346 Similarly to our study's inclusion level, the same pattern has also been observed in the relative gut
347 length of European sea bass (Mastoraki et al., 2020a).

348 **4.2. Whole-body proximate composition and nutrient retention**

349 Whole-body dry matter, protein, fat, ash, and energy contents of the experimental fish were
350 similar across dietary treatments. Our results are in line with other studies with similar or higher dietary
351 insect meal inclusion; for example, fish meal substitution with HI or MD in barramundi (Katya et al.,
352 2017; Lin and Mui, 2017), with HI in Atlantic salmon (*Salmo salar*; Belghit et al., 2018) or with TM
353 in European sea bass (Gasco et al., 2016). A general trend of increasing body fat is observed in different
354 studies when fish meal is substituted with insect meals, sometimes accompanied with a decrease in
355 protein content. The higher fat content observed in insect meal-fed fish is usually explained by the
356 higher dietary content of saturated fatty acids and by the change in the ω -3/ ω -6 ratio which can enhance
357 lipogenesis (Alves et al., 2020). On the other hand, insect fat which is rich in medium-chain fatty acids
358 is not stored but readily utilized for energy production (Tocher, 2015), leading to lower body fat
359 contents in salmon and Jian carp (Belghit et al., 2019; *Cyprinus carpio* var. *Jian*; Li et al., 2016). In this
360 study, the lack of significant differences in the whole-body fat content was corroborated by the similar
361 lipogenic enzymes activities.

362 The whole-body amino acid profile was not affected by the inclusion of the different insect
363 meals. Similarly, no effect was observed in the muscle of grass and Jian carp both fed HI (Zhou et al.,
364 2018; Lu et al., 2020). Studies have also shown that the inclusion of insect meals does not affect the
365 essential amino acid content of European sea bass fed TM, HI, or MD (Mastoraki et al., 2020a), Jian
366 carp fed silkworm meal (Ji et al., 2015), and rainbow trout fed TM (Jeong et al., 2020). Additionally,
367 Jeong et al. (2020) report no adverse effects on muscle proximate and essential amino acid composition.

368 Major differences were observed in the whole-body fatty acid content driven by the different
369 fatty acid profiles and fish oil levels of the experimental diets. Thus, due to the defatted nature of HI
370 meal used in the present study, higher percentage of fish oil was included in the HI diet to ensure similar
371 lipid and energy contents among the dietary treatments. The inclusion of insect meals did not seem to
372 affect fish total whole-body SFA content, in line with previous reports employing other fish species
373 using TM or MD (Gasco et al., 2016; Iaconisi et al., 2018; Mastoraki et al., 2020a). In contrast, an
374 increasing SFA content was observed in salmon using full fat HI, along with the increase in the HI meal
375 inclusion due to the higher content of lauric (C12:0) and myristic (C14:0) acids of the meal (Bruni et
376 al., 2020). In this study, the whole-body total MUFA content was similar across experimental groups,
377 in agreement with other studies in European sea bass fed TM, HI, or MD (Mastoraki et al., 2020a). On
378 the contrary, higher MUFA content has been reported in the fillets of rainbow trout fed TM (Iaconisi et
379 al., 2018), and lower ones in rainbow trout (Secci et al., 2018) and Eurasian perch (Stejskal et al., 2020)
380 fed HI; this is due to the differences in the respective fatty acid profiles of the insect meals used in these

381 investigations. In our study, higher inclusion of fish oil in the FM and HI diets resulted in higher total
382 ω -3 PUFA, EPA and DHA whole-body content in the respective groups. This fact is also in agreement
383 with Mastoraki et al. (2020a) employing a highly defatted HI meal in the diet of European sea bass.
384 However, the contents of ω -3 PUFA are reported to decline when the inclusion of fish oil is lower due
385 to the use of partially defatted (Stejskal et al., 2020) or full fat insect meals (present study and Sánchez-
386 Muros et al., 2016).

387 Protein retention was across the experimental groups of this study; however, dry matter and
388 energy retentions were lower in fish fed HI compared to the FM group. Moreover, the HI fed sea breams
389 had the lowest fat retention. Dry matter and energy retentions were found to be negatively correlated (p
390 < 0.05) to dietary crude fiber, while fat retention was found to be negatively correlated to dietary crude
391 fiber and ADF. Therefore, the lower fat retention of the HI group could be attributed to the presence of
392 chitin in the ADF fraction which was higher compared to the other diets and it is reported to inhibit fat
393 absorption (Kroeckel et al., 2012) and fatty acid synthesis (Coz-Rakovac et al., 2005). However, in the
394 present study, the liver lipogenic enzymes including FAS were not affected by the dietary treatments.
395 Recently, Panteli et al. (2021) using a 30% substitution of fish meal with HI meal for the diet of gilthead
396 sea bream, have observed a decrease in the abundance of the beneficial bacteria of the phylum
397 Firmucutes in the intestine of the fish fed HI compared to the control group. This decrease could
398 probably be responsible for the reduction in the fatty acid absorption effectiveness of the gut (Panteli et
399 al. 2021) and the lower fat retention; this in turn may explain the decreased fat retention of the HI-fed
400 sea breams without any differences in the lipogenic enzymes of the present study. The nutrient
401 utilization of different fish species fed with different insect meals were diversified in this research. The
402 complete replacement of FM with HI (35% inclusion) in African catfish may increase fat retention
403 without affecting protein deposition (Huda et al., 2020), while 26.4% inclusion of HI in the diet of
404 rainbow trout may increase fat retention and decrease protein deposition (Dumas et al., 2018). Enhanced
405 protein retention is observed in rainbow trout fed TM (Rema et al., 2019), whereas the 20% inclusion
406 of TM in mandarin fish (*Siniperca scherzeri*) is reported to improve protein and fat deposition (Sankian
407 et al., 2018). Several factors can be involved in these inconsistent effects, such as the species identity
408 and age of the fish involved, the different insect species used, the design and formulation of the diet or
409 even the quality of the ingredients. Moreover, since the production of insect meals is not a standardized
410 process, the different substrates and processing methods of the insects can also affect the insect meal
411 quality (Becker and Yu, 2013; Reyes et al., 2020). In the present study, the inclusion of different insect
412 meals affected the retention/deposition of 10 amino acids out of 17 studied in total. The observed
413 differences in the amino acid depositions could be attributed to the different dietary amino acid content
414 based on the detected negative correlations.

415

416 4.3. Plasma and liver enzyme activities

417 Regarding plasma metabolites, fish meal substitution had no negative effects on plasma glucose
418 and lactate, and the levels observed were in line with the up to date reported values, for unstressed
419 gilthead sea bream (Peres et al., 2013; Rotllant et al., 2001). Cholesterol and triglycerides were higher
420 in the fish fed FM and HI diets (differences not statistically significant though) and the same is true for
421 European sea bass cholesterol levels (Mastoraki et al., 2020a). This trend could be attributed to the
422 higher content of these energetic metabolites in the fish oil (Mastoraki et al., 2020a). Additionally, it
423 has been reported that insect linoleic acid promotes the breakdown of cholesterol and triglycerides
424 (Song et al., 2018). Therefore, the higher linoleic content of the TM and MD diets could have led to
425 lower plasma cholesterol and triglycerides in fish fed with those two insect meals. Fish meal
426 substitution with insect meals in the diets of Japanese sea bass (*Lateolabrax japonicus*, HI), Korean
427 rockfish (*Sebastes schlegeli*, TM), mirror carp (*Cyprinus carpio* var. *specularis*, hydrolyzed silkworm),
428 African catfish (TM), pearl gentian grouper (*Epinephelus lanceolatus* x *E. fuscoguttatus*, TM),
429 European sea bass (HI) and mandarin fish (TM) are reported to result in lower plasma cholesterol and/or
430 triglycerides, an effect attributed to the presence of chitin (Fawole et al., 2020; Khosravi et al., 2018;
431 Li et al., 2017; Magalhães et al., 2017; Sankian et al., 2018; Song et al., 2018; Wang et al., 2019; Xu et
432 al., 2018).

433 Increased plasma aminotransferase activities are related to liver-cell function impairment in
434 cases of severe steatosis, and this is positively correlated with the degree of tissue necrosis (Lemaire et
435 al., 1991). Plasma ALT and AST activities did not differ among the studied groups herein. The levels
436 of both aminotransferases were within the reported range for healthy gilthead sea bream (Peres et al.,
437 2013). In the liver, the ALT activity of fish fed HI was significantly higher compared to the TM group,
438 whereas the activities of other amino acid catabolizing enzymes (GDH and AST) were not affected by
439 the dietary treatment. The amino acid catabolizing enzyme activities are positively correlated with
440 growth performance and therefore an increase in the activity can indicate better protein utilization
441 (Kumar et al., 2017; Lin, Luo, 2011). In addition, high dietary protein content promotes amino acid
442 catabolism (Ballantyne, 2001; Fynn-Aikins et al., 1995). On the other hand, a diet rich in fat may
443 decrease the activity of amino acid catabolizing enzymes due to the sparing effect of fat on amino acids
444 used for energy production (Ballantyne, 2001). In the present study, the growth performance exhibited
445 converse results, with the HI group presenting lower SGR compared to TM. Moreover, the HI diet had
446 the lowest protein content and the highest fat content which would have lowered liver ALT activity. A
447 possible explanation for this could be the higher dietary fiber content, which may have lowered even
448 further the digestible energy of a diet with an already lower energy content. A lower energy availability
449 might have led to a higher demand for amino acids so as to be catabolized for energy production (Kumar
450 et al., 2010), and consequently, this may have increased the activity of ALT in the liver of HI-fed fish.
451 Given that the biological significance of the hepatic ALT activity is still unclear, it has been suggested

452 that the substitution of fish meal with defatted TM in rainbow trout (Chemello et al., 2020), with HI in
453 meagre (Guerreiro et al., 2020), as well as hydrolysed TM in sea trout (Hoffmann et al., 2020) do not
454 actually affect the activity of liver amino acid catabolic enzymes. On the contrary, in rainbow trout, the
455 inclusion of TM may result in higher AST activity compared to the inclusion of HI, without affecting
456 ALT and GDH (Melenchón et al., 2020).

457 Lipogenic enzymes' activity can be affected by the level and quality of dietary protein (Alvarez
458 et al., 1998; Dias et al., 2005; Wacyk et al., 2012) and fat (Alvarez et al., 1998; Jordal et al., 2007).
459 Menoyo et al. (2004) have reported a decrease in liver lipogenesis with vegetable oils' replacement
460 (80%) in the diets of gilthead sea bream. Peng et al. (2017) have attributed the increase in FAS activity
461 of turbot (*Scophthalmus maximus*) fed plant oils to the increase of the saturated and mono-unsaturated
462 fatty acid content of the diets. Furthermore, Menoyo et al. (2003) have reported a lowering effect of
463 dietary ω -3 fatty acids on G6PD and ME activity of Atlantic salmon. In a feeding trial with rainbow
464 trout, the dietary inclusion of up to 20% partially defatted TM have led to comparable dietary ω -3 fatty
465 acid contents, resulting in no differences of the lipogenic enzymes' activities (Chemello et al., 2020).
466 Despite the 44% replacement of fish oil by insect meals in the present study, no differences were
467 observed in the liver lipogenic enzyme activities; this may imply an adequate fish oil supply or a non-
468 detrimental effect of insect meal fat at this rate of inclusion.

469 **Conclusion**

470 Under the experimental conditions examined herein, all three insect meal diets performed
471 equally well compared to the fish meal diet, in terms of growth performance, feed consumption and
472 feed conversion. However, among the insect meal diets, HI was slightly inferior compared to TM and
473 MD. The whole-body composition was not affected by the different diets. Protein retention was similar
474 to all experimental groups, however fat retention was lower in the fish fed HI probably due to the higher
475 fiber content. Differences were observed in the amino acid depositions among the experimental groups
476 which could be attributed to the differences in the dietary amino acid composition. In addition, the TM
477 and MD majorly affected the whole-body fatty acid composition, due to the lower fish oil inclusion
478 compared to the FM and HI diets. Plasma metabolites and liver lipogenic enzymes were not affected
479 by the different diets. Overall, fish meal can be successfully substituted by *Tenebrio molitor*, *Hermetia*
480 *illucens* and *Musca domestica* meals in the diets of gilthead sea bream at a rate of 30%.

481

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498

499 **Author statement:**

500 The authors here declare their individual contributions:

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502 **Katsika** Investigation. **Paula Enes:** Investigation, Formal analysis, Writing - review & editing. **Inês**

503 **Guerreiro:** Investigation, Writing - review & editing. **Yannis P. Kotzamanis:** Investigation, Formal

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882

Table 1: Ingredients and proximate composition of insect meals and experimental diets

	FM	TM	HI	MD
Ingredients (%)				
Fish meal (Peru, prime)	65	45.5	45.5	45.5
Insect larvae meal	0	19.5	19.5	19.5
Fish oil	9	5	9.7	6.3
Wheat	17.2	16.6	14.8	16.8
Wheat gluten meal	6	9	6	8.5
Vitamin & mineral mix ^a	2.5	2.5	2.5	2.5
DL-methionine	0.3	0.7	0.7	0
L-lysine	0	1.2	1.3	0.9
Proximate composition of the different insect meals (dry basis) ^b				
Crude Protein (%)		61.0	67.0	58.5
Crude Lipid (%)		28.6	5.7	23.1
Ash %		4.1	7.8	7.4
Gross energy (MJ kg ⁻¹)		26.9	21.4	24.8
Proximate composition of the experimental diets (dry basis) ^b				
Crude Protein (%)	58.0	57.6	55.8	56.6
Adjusted Crude Protein (%) ^c	56.7	55.5	51.9	54.5
Crude Lipid (%)	17.7	16.1	18.1	16
Ash (%)	12.6	9.2	10.9	10.7
Crude fiber (%)	1.7	2.5	3.9	2.9
Acid detergent fiber (%)	6.2	5.6	8.8	7.1
NFE (%) ^d	10.1	14.5	11.3	13.9
Gross energy (MJ kg ⁻¹)	22.1	22.4	22.0	22.1
EAA (%)				
Arginine	3.04	2.60	2.69	2.80
Histidine	1.07	1.12	1.03	1.12
Isoleucine	2.28	2.19	2.14	2.18
Leucine	4.15	3.97	3.86	3.99
Lysine	3.81	4.19	4.10	4.38

Methionine	1.54	2.09	1.59	1.60
Phenylalanine	2.16	1.92	1.95	2.38
Threonine	2.36	2.21	2.21	2.34
Valine	2.63	2.73	2.59	2.59
NEAA (%)				
Alanine	3.20	3.42	3.12	3.17
Asx	4.13	4.06	3.97	4.70
Cysteine	0.28	0.24	0.23	0.26
Glx	8.32	8.11	7.83	8.97
Glycine	3.01	2.65	2.67	2.61
Proline	2.88	3.07	2.90	2.83
Serine	2.45	2.29	2.33	2.41
Tyrosine	1.52	1.81	1.61	1.91

Abbreviations: FM, Fish meal; TM, *Tenebrio molitor*; HI, *Hermetia illucens*; MD, *Musca domestica*; EAA, essential amino acids; NEAA, non-essential amino acids; Asx, sum of asparagine and aspartate; Glx, sum of glutamine and glutamate

^a Premix (kg⁻¹): Choline 90,000 (mg) Vitamin A 0.3 (MIU), Vitamin D3 0.1 (MIU), Vitamin E 20,000 (IU), Vitamin K 1030 (mg), Vitamin B1 390 (mg), Vitamin B 960 (mg), Nicotinic acid 2600 (mg), Pantothenic acid 4400 (mg), Vitamin B6 890 (mg), Vitamin B12 15 (mg), Folic acid 290 (mg), Biotin 14 (mg), Vitamin C (Stay C 35% MONO) 20,300 (mg), Inositol 15,600 (mg), Total Mn 1200 (mg), Total Ca 72,000 (mg), Total Zn 7,000 (mg), Total Cu 450 (mg), Total Se 14 (mg), Total I 100 (mg), Betaine (mg) 71,250 (mg), BHA (E320) 3000 (mg)

^b Mean of triplicate analyses

^c Protein adjusted for the nitrogen linked to acid detergent fiber

^d Nitrogen-free extract, NFE = 100 - % crude protein - % crude lipid - % ash - % crude fiber

Table 2: Fatty acid composition (% of total fatty acids) of the experimental diets in which 30% of the fish meal (FM) was substituted with *Tenebrio molitor* (TM), *Hermetia illucens* (HI) or *Musca domestica* (MD) larvae meal

	FM	TM	HI	MD
12:0	0.12	0.14	2.71	ND
14:0	3.61	3.84	4.48	3.21
16:0	13.38	13.20	14.21	16.84
16:1 ω -7	4.81	4.68	4.94	9.08
17:0	0.78	0.87	1.68	1.50
17:1	0.65	0.76	0.63	0.69
18:0	2.08	2.27	2.52	2.93
18:1 ω -9	23.37	26.57	23.64	25.09
18:2 ω -6	8.03	12.10	9.52	13.10
20:1 ω -9	2.03	1.61	1.94	1.86
18:3 ω -3	6.46	4.20	6.15	4.20
21:0	2.04	1.75	1.79	1.14
20:2 ω -6	0.57	0.72	0.59	0.33
22:1 ω -9	0.64	0.42	0.52	0.54
20:3 ω -3	8.29	5.79	7.74	5.01
20:4 ω -6	0.70	0.51	0.58	0.39
22:2 ω -6	5.70	5.46	4.55	3.56
20:5 ω -3	1.43	1.07	1.13	0.85
22:6 ω -3	9.21	7.50	7.26	5.77
SFA	22.85	22.95	28.44	26.51
MUFA	32.76	35.14	32.95	38.87
PUFA	40.72	37.65	37.87	33.50
ω -3	25.39	18.56	22.28	15.82
ω -6	15.33	19.09	15.59	17.68

Abbreviations: SFA, saturated fatty acids; MUFA, mono-unsaturated fatty acids; PUFA, poly-unsaturated fatty acids; ND, not detected

Means of duplicate analyses

Table 3: Effect of 30% substitution of fish meal with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*) on growth performance and somatic indices of gilthead sea bream

	FM	TM	HI	MD
Survival (%)	100	99.1±1.0	100	100
IBW (g)	29.7±0.2	29.1±0.6	29.4±0.5	29.8±0.2
FBW (g)	121.4±2.1	125.3±0.8	119.2±2.3	126.6±0.4
WG (%)	308.3±4.3 ^{ab}	331.3±7.5 ^a	305.8±5.2 ^b	324.8±4.4 ^{ab}
SGR (% day ⁻¹)	1.51±0.01 ^{ab}	1.57±0.02 ^a	1.51±0.01 ^b	1.56±0.01 ^{ab}
DFI (% BW day ⁻¹)	1.36±0.03	1.41±0.02	1.46±0.01	1.38±0.02
FCR	1.04±0.03	1.06±0.03	1.12±0.02	1.04±0.01
Somatic indices				
CF	1.55±0.01	1.66±0.02	1.59±0.02	1.61±0.05
HSI (%)	1.24±0.04	1.45±0.06	1.41±0.12	1.40±0.04
VSI (%)	5.40±0.01	5.70±0.10	5.68±0.20	5.66±0.12
MFI (%)	0.90±0.06	0.97±0.04	1.12±0.08	1.01±0.07
RGL	1.69±0.07	1.85±0.03	1.83±0.05	1.77±0.02

Abbreviations: IBW, initial body weight; FBW, final body weight; WG, weight gain; SGR, specific growth rate; DFI, daily feed intake; FCR, feed conversion ratio; CF, condition factor; VSI, viscerosomatic index; HSI, hepatosomatic index; MFI, mesenteric fat index; RGL, relative gut length

Mean ± standard error, n = 3 tanks per diet. Different letters in the same row denote statistically significant difference ($p < 0.05$)

Table 4: Effect of 30% substitution of fish meal with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*) on whole-body proximate and amino acid composition

	Initial	FM	TM	HI	MD
Dry matter	27.4	33.3±0.3	32.7±0.6	32.5±0.4	32.4±0.4
Crude protein	16.3	17.6±0.1	17.7±0.2	17.8±0.1	17.8±0.0
Crude fat	6.5	11.9±0.1	11.1±0.6	10.4±0.4	10.3±0.3
Ash	4.8	3.4±0.1	3.6±0.1	3.6±0.1	3.6±0.1
Gross energy (MJ kg ⁻¹)	6.1	8.7±0.1	8.5±0.3	8.3±0.2	8.3±0.1
EAA					
Arginine	3.21	3.03±0.07	3.02±0.11	3.25±0.12	3.12±0.03
Histidine	1.21	1.35±0.05	1.32±0.07	1.49±0.06	1.40±0.04
Isoleucine	2.27	2.15±0.01	2.15±0.09	2.24±0.05	2.19±0.04
Leucine	4.27	3.93±0.04	3.93±0.17	4.06±0.09	4.01±0.06
Lysine	4.80	4.54±0.05	4.58±0.18	4.40±0.01	4.61±0.15
Methionine	1.56	1.53±0.05	1.50±0.06	1.66±0.07	1.55±0.01
Phenylalanine	2.02	2.01±0.08	1.92±0.10	2.18±0.09	2.04±0.06
Threonine	2.63	2.39±0.03	2.41±0.10	2.47±0.06	2.44±0.02
Valine	2.65	2.48±0.01	2.49±0.10	2.57±0.06	2.54±0.04
NEAA					
Alanine	3.62	3.31±0.04	3.29±0.09	3.37±0.03	3.45±0.03
Asx	5.38	5.16±0.08	4.94±0.30	5.12±0.02	5.36±0.10
Cysteine	0.29	0.28±0.01	0.27±0.01	0.30±0.01	0.27±0.01
Glx	7.81	7.15±0.06	7.02±0.35	7.08±0.06	7.38±0.13
Glycine	3.48	3.41±0.24	3.32±0.13	3.85±0.12	3.65±0.17
Proline	2.43	2.24±0.08	2.25±0.05	2.39±0.06	2.35±0.05
Serine	2.51	2.19±0.03	2.21±0.08	2.30±0.06	2.27±0.02
Tyrosine	1.41	1.59±0.08	1.52±0.09	1.78±0.09	1.64±0.05

Abbreviations: EAA, essential amino acids; NEAA, non-essential amino acids; Asx, sum of asparagine and aspartate; Glx, sum of glutamine and glutamate.

Percentage (%) on wet basis unless otherwise stated. Mean ± standard error, n = 3 tanks per diet. No statistically significant differences were observed.

Table 5: Fatty acid composition of whole-body of gilthead sea bream fed diets in which 30% of the fish meal was substituted with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*) and correlational analysis with dietary fatty acids and dietary fish oil inclusion

	Initial	FM	TM	HI	MD	r feed	r fish oil
12:0	0.13	0.06±0.01 ^b	0.06±0.01 ^b	1.00±0.01 ^a	0.08±0.01 ^b	0.389	0.518
14:0	3.66	3.11±0.07 ^{ab}	2.82±0.13 ^{ab}	3.49±0.15 ^a	2.66±0.24 ^b	0.605 [*]	0.756 ^{**}
16:0	17.75	12.90±0.15	15.72±0.45	12.36±0.32	14.98±1.01	-0.086	-0.842 ^{***}
16:1 ω-7	5.87	6.70±0.10 ^c	5.76±0.11 ^d	6.82±0.07 ^{bc}	9.30±0.30 ^a	0.907 ^{***}	0.324
17:0	0.82	1.07±0.03	0.72±0.10	1.00±0.02	1.24±0.34	-0.043	0.367
17:1	1.56	0.79±0.06 ^a	0.45±0.04 ^b	0.62±0.00 ^{ab}	0.68±0.07 ^{ab}	-0.626 [*]	0.626 [*]
18:0	4.21	2.24±0.08 ^{ab}	2.77±0.10 ^a	1.97±0.06 ^b	2.71±0.23 ^a	0.065	-0.885 [*]
18:1 ω-9	43.08	32.36±0.30 ^b	38.04±0.76 ^a	32.87±0.20 ^b	33.89±1.18 ^b	0.734 ^{**}	-0.626 [*]
18:2 ω-6	2.69	9.04±0.17 ^c	12.63±0.25 ^a	10.52±0.22 ^b	11.66±0.30 ^a	0.799 ^{***}	-0.756 ^{**}
20:1 ω-9	0.14	1.92±0.07 ^{ab}	1.68±0.02 ^{bc}	2.01±0.06 ^a	1.56±0.06 ^c	0.648 [*]	0.756 ^{**}
18:3 ω-3	3.37	5.04±0.12 ^a	3.28±0.02 ^c	4.97±0.05 ^a	3.82±0.17 ^b	0.907 ^{***}	0.842 ^{***}
21:0	0.87	1.80±0.08 ^a	1.40±0.09 ^{ab}	1.51±0.08 ^{ab}	0.71±0.37 ^b	0.907 ^{***}	0.475
20:2 ω-6	6.76	0.50±0.03	0.23±0.12	0.45±0.03	0.34±0.01	-0.022	0.713 ^{**}
22:1 ω-9	ND	0.53±0.01 ^a	0.36±0.01 ^c	0.53±0.01 ^a	0.42±0.01 ^b	0.561	0.885 ^{***}
20:3 ω-3	1.76	4.83±0.11 ^a	2.93±0.03 ^b	4.78±0.02 ^{ab}	3.54±0.02 ^{ab}	0.713 ^{**}	0.842 ^{***}
20:4 ω-6	ND	1.67±0.98	0.39±0.04	0.68±0.03	0.30±0.12	0.820 ^{***}	0.734 ^{**}
22:2 ω-6	1.33	3.36±0.03 ^a	2.06±0.02 ^b	3.19±0.10 ^a	2.16±0.13 ^b	0.475	0.734 ^{**}
20:5 ω-3	0.77	1.19±0.01 ^a	0.87±0.01 ^c	1.10±0.02 ^{ab}	1.01±0.05 ^b	0.691 [*]	0.691 [*]
22:6 ω-3	0.72	7.55±0.07 ^a	4.88±0.06 ^b	6.90±0.12 ^{ab}	5.12±0.32 ^{ab}	0.600 [*]	0.713 ^{**}

SFA	28.79	22.28±0.26	24.41±0.87	22.33±0.44	23.41±2.07	0.000	-0.518
MUFA	51.67	43.92±0.52	47.30±0.72	44.44±0.21	47.40±1.52	0.691*	-0.691**
PUFA	17.92	33.46±0.57 ^a	27.50±0.27 ^b	32.77±0.37 ^a	28.14±0.96 ^b	0.756**	0.799***
ω-3	6.62	18.60±0.30 ^a	11.97±0.08 ^c	17.75±0.09 ^a	13.50±0.43 ^b	0.777**	0.777**
ω-6	11.31	14.86±0.87	15.53±0.20	15.02±0.31	13.27±1.13	0.410	-0.626*

Abbreviations: SFA, saturated fatty acids; MUFA, mono-unsaturated fatty acids; PUFA, poly-unsaturated fatty acids; ND, not detected

Percentage (%) of total fatty acids. Mean ± standard error, n = 3 tanks per diet. Different letters in the same row denote statistically significant difference ($p < 0.05$). In the correlational analysis an asterisk (*) indicates significance at the 0.05 level, ** at the 0.01 level and *** at the 0.001 level.

Table 6: Nutrient retention, essential amino acid retention efficiency, non-essential amino acid deposition and correlational analysis with dietary amino acids of gilthead sea bream fed diets in which 30% of the fish meal was substituted with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*)

	FM	TM	HI	MD	r
Dry matter	33.8±0.6 ^a	32.5±0.1 ^{ab}	30.5±0.4 ^b	32.8±0.6 ^a	nd
Crude protein	29.9±0.6	29.8±1.0	29.3±0.2	31.3±0.4	nd
Crude fat	74.2±1.2 ^a	73.2±3.0 ^a	57.0±2.3 ^b	69.3±2.6 ^a	nd
Ash	22.4±1.3 ^c	33.3±1.4 ^a	26.1±1.1 ^b	29.1±0.8 ^a	nd
Gross Energy	41.5±0.9 ^a	39.3±0.8 ^{ab}	36.6±0.8 ^b	39.3±0.2 ^{ab}	nd
EAA					
Arginine	27.8±0.7 ^b	32.0±0.9 ^a	29.2±0.4 ^{ab}	30.4±0.4 ^{ab}	-0.756 ^{**}
Histidine	29.7±0.8	28.1±0.8	28.9±0.4	28.6±0.4	-0.324
Isoleucine	26.3±0.7	26.9±0.8	26.0±0.4	27.6±0.4	-0.043
Leucine	27.1±0.7	28.0±0.8	27.1±0.4	28.4±0.4	0.108
Lysine	33.2±0.8 ^a	29.8±0.8 ^b	28.7±0.4 ^b	29.1±0.4 ^b	-0.453
Methionine	26.6±0.7 ^a	19.3±0.6 ^c	24.0±0.3 ^b	25.8±0.3 ^{ab}	-0.691 ^{**}
Phenylalanine	24.7±0.6 ^{bc}	27.4±0.8 ^a	25.4±0.4 ^{ab}	22.5±0.3 ^c	-0.907 ^{***}
Threonine	29.3±0.7	31.0±0.9	29.2±0.4	29.7±0.4	0.000
Valine	26.6±0.7	25.3±0.7	25.0±0.4	27.1±0.4	0.043
NEAA					
Alanine	29.9±0.6 ^{ab}	27.5±0.8 ^b	28.4±0.4 ^{ab}	30.3±0.4 ^a	-0.173
Asx	34.3±0.9 ^a	34.4±1.0 ^a	33.1±0.5 ^{ab}	30.3±0.4 ^b	-0.453
Cysteine	27.0±0.7 ^b	31.1±0.9 ^a	30.9±0.4 ^a	29.6±0.4 ^{ab}	-0.820 ^{***}
Glx	24.7±0.6	25.0±0.7	24.4±0.3	23.1±0.3	-0.518
Glycine	30.5±0.8 ^c	34.2±1.0 ^{ab}	31.9±0.5 ^{bc}	35.3±0.5 ^a	-0.885 ^{***}
Proline	22.3±0.6 ^{ab}	20.6±0.6 ^b	20.5±0.3 ^b	22.8±0.3 ^a	-0.756 ^{**}
Serine	27.0±0.7	28.4±0.8	26.4±0.4	27.6±0.4	-0.108
Tyrosine	24.3±0.6 ^a	20.2±0.6 ^b	21.4±0.3 ^b	19.6±0.3 ^b	-0.864 ^{**}

Abbreviations: EAA, essential amino acids; NEAA, non-essential amino acids; Asx, sum of asparagine and aspartate; Glx, sum of glutamine and glutamate; nd, not determined.

Mean ± standard error, n = 3 tanks per diet. Different letters in the same row denote statistically significant difference ($p < 0.05$). In the correlational analysis an asterisk (*) indicates significance at the 0.05 level, ** at the 0.01 level and *** at the 0.001 level.

Table 7: Effect of 30% substitution of fish meal with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*) on plasma metabolites of gilthead sea bream

	FM	TM	HI	MD
Glucose (mg dl ⁻¹)	93.8±15.8	103.4±11.7	98.2±9.1	112.0±8.4
Cholesterol (mg dl ⁻¹)	332.6±5.6	279.4±5.2	320.1±22.1	262.9±3.5
Triglycerides (mg dl ⁻¹)	338.8±26.9	337.6±84.8	466.0±48.2	312.8±6.9
Phospholipids (mg dl ⁻¹)	887.2±31.1	894.8±12.2	879.8±42.9	877.1±37.8
Lactate (mg dl ⁻¹)	12.2±0.9	17.4±4.4	15.0±1.0	13.8±2.1
ALT (u l ⁻¹)	29.8±3.1	19.8±2.7	65.4±26.6	28.6±11.1
AST (u l ⁻¹)	31.5±5.3	59.5±12.4	56.3±13.9	32.4±2.0

Abbreviations: ALT, Alanine aminotrasferase; AST, Aspartate aminotransferase. Mean ± standard error, n = 3 tanks per diet. Different letters in the same row denote statistically significant difference ($p < 0.05$).

Table 8: Effect of 30% substitution of fish meal with different insect meals (TM: *Tenebrio molitor*, HI: *Hermetia illucens* or MD: *Musca domestica*) on liver amino acid catabolism and lipogenic enzymes of gilthead sea bream

	FM	TM	HI	MD
Alanine aminotrasferase (ALT)	432.4±19.6 ^{ab}	368.9±5.4 ^b	457.9±25.5 ^a	432.6±6.9 ^{ab}
Aspartate aminotransferase (AST)	1539.9±8.3	1570.3±52.9	1656.5±85.8	1688.6±54.4
Glutamate dehydrogenase (GDH)	65.7±2.5	64.0±2.8	61.7±1.6	66.0±2.1
Glucose-6-phosphate dehydrogenase (G6PD)	136.1±4.4	163.5±14.9	143.3±10.4	167.9±7.9
Fatty acid synthase (FAS)	17.1±2.9	22.5±1.7	21.8±1.7	22.1±1.0
Malic enzyme (ME)	9.8±1.3	11.8±1.3	11.6±0.4	12.9±0.5

Expressed as mU mg protein⁻¹ (nmoles min⁻¹ mg protein⁻¹). Mean ± standard error, n = 3 tanks per diet. Different letters in the same row denote statistically significant difference ($p < 0.05$).