



Effects of Dietary Protein Source and Feeding Regime on Growth Performance, Nutrient Digestibility, Fatty Acids and Fillet Quality Traits in Rainbow Trout (*Oncorhynchus mykiss*)

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Review

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3 1 **Effects of Dietary Protein Source and Feeding Regime on Growth Performance,**
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7 3 **(Oncorhynchus mykiss)**
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3 26 **Abstract**
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5 27 **The aim of this study was to investigate the effects of feeding regime and of**
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7 28 **bacterial protein meal (BPM), pea protein concentrate (PPC) and a mixture thereof**
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9 29 **(MIX) compared to a control fish meal-based diet on growth performance, nutrient**
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11 30 **digestibility, fatty acid (FA) profile and fillet quality traits in rainbow trout. A stock of**
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13 31 **1200 juvenile rainbow trout were individually weighed (mean weight 114.6 ± 0.2 g) and**
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15 32 **randomly distributed into 24 fibre-glass tanks (4 diets x 3 replications x 2 feeding**
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17 33 **regimes). Statistical differences appeared among the diets in terms of crude protein**
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19 34 **digestibility, while no differences appeared for dry matter, ether extract and gross**
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21 35 **energy digestibility. Growth performance and somatic indexes were significantly**
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23 36 **affected by the diet effect, while only the condition factor was influenced by the feeding**
24
25 37 **rate effect. None of the parameters appeared to be affected by the interaction effects.**
26
27 38 **Differences appeared between the FA profiles of the dorsal muscle. Oleic, linoleic, α -**
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29 39 **linolenic, and docosahexaenoic acid contents were influenced by diet, while only minor**
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31 40 **FAs were influenced by feeding regime. Consumer tests showed that fillets of trout fed**
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33 41 **the MIX diet ad libitum were the most preferred. A similar ranking was obtained with**
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35 42 **the trout fed rationed diets.**
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3 44 The use of alternative protein sources in aquaculture as fishmeal substitutes is an
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5 45 extensively studied subject (Bakke-McKellep and Refstie 2008; Médale and Kaushik 2008),
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7 46 since fishmeal will be a limited resource for fish feedstuff production in the future. In the past
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9 47 decade, in fact, a great deal of research into aquaculture nutrition has dealt with fishmeal and
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11 48 fish oil substitution with alternative sources. Various microbes (algae, fungi and bacteria)
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13 49 have been used to produce a wide range of single cell protein varieties (Anupama and
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15 50 Ravindra 2000). They can be used for fish or shellfish as a substitute for fish meal (4-5%
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17 51 substitution) and have been investigated as a feed ingredient in diets for rainbow trout
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19 52 (Øverland et al. 2006; Aas et al. 2006), Atlantic halibut (Aas et al. 2007), and Atlantic salmon
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21 53 (Storebakken et al. 2004).

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25 54 Among the protein concentrates derived from fermentation bacteria, noted for its use
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27 55 as an attractant in fish food and shellfish, Protorsan is a bacterial protein meal (BPM) which is
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29 56 a by-product of fermentation conducted by Corynebacterium melassecola that led to the
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31 57 production of L-glutamic acid fermentation carried out using plant substrates, usually from
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33 58 beet molasses and/or starch hydrolysates. Fermentation takes place anaerobically, under
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35 59 optimal pH and temperature conditions for the growth of Corynebacterium melassecola, for
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37 60 about 36 hours, followed by heat treatment of fermentation broth at 75 C for 30 minutes to
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39 61 deactivate the bacteria. The bacterial mass is then separated from the liquid phase by
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41 62 centrifugation and subjected to washing and drying. The resulting product is used for animal
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43 63 feed. Protorsan contains 12% L-glutamic acid, around 7-7.5% of total nucleotides and 4.5%
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45 64 betaine, a methyl donor with high palatability. It also contains high levels of peptidoglycan as
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47 65 components of the bacterial cell wall. Protorsan has been tested as a feed stimulant in diets for
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49 66 sea bream (Chatzifotis et al. 2009), while no studies have been performed in rainbow trout
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51 67 (Oncorhynchus mykiss).

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3 68 As far as alternative protein sources are concerned, the best growth performances have
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5 69 been achieved using plant protein concentrates and plant protein mixture (De Francesco et al.
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7 70 2004, 2007). Among the plant protein sources, field peas (Pisum sativum) have reported some
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9 71 success for different fish species such as Atlantic salmon (Øverland et al. 2009), rainbow trout
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11 72 (Thiessen et al. 2003), hybrid sturgeon (Sicuro et al. 2012), common carp (Davies and
12
13 73 Gouveia 2010), gilthead seabream (Sánchez-Lozano et al. 2009, 2011), sea bass (Tibaldi et al.
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15 74 2005, Tulli et al. 2007), African catfish (Davies and Gouveia 2008), and Nile tilapia (Schulz
16
17 75 et al. 2007).

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19
20 76 Øverland et al. (2009) showed that 20% air-classified pea protein concentrate (PPC)
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22 77 could replace 20% of high-quality fish meal protein in feed without any adverse effect on
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24 78 growth performance, carcass composition or distal intestine histology in Atlantic salmon. By
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26 79 contrast, in another study, as PPC at high inclusion levels was shown to induce enteropathy in
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28 80 the distal intestine of Atlantic salmon, the authors concluded that caution should be used
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30 81 when including PPC in formulated feeds for Atlantic salmon (Penn et al. 2011).

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33 82 Feeding PPC has been reported to support acceptable weight gain, feed intake, and
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35 83 feed conversion in both Atlantic salmon (Carter and Hauler 2000) and rainbow trout
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37 84 (Thiessen et al. 2003).

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40 85 The aim of this study was to investigate the effects both of BPM, PPC and a mixture
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42 86 thereof compared to a control fish meal-based diet and of feeding regime on growth
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44 87 performance, nutrient digestibility, fatty acid (FA) profile and fillet quality traits in rainbow
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46 88 trout (Oncorhynchus mykiss).

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50 51 52 90 **Materials and Methods**

53 54 91 Experimental plan

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3 92 Three experimental diets were obtained by including BPM, PPC or a mixture of both
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5 93 protein concentrates (MIX), respectively, replacing fish meal. These experimental diets were
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7 94 tested against a control fish meal (FM)-based diet; all the diets were isonitrogenous (CP 45
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9 95 %) and isoenergetic (22 MJ/kg DM).

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11 96 The feeds were manufactured in the laboratory at the Experimental Station of the
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13 97 Department of Agriculture, Forestry, and Food Sciences of the University of Torino by means
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15 98 of a pelleting process using a 3.5 mm diameter. Pellets were dried in a stove overnight at 50 C
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17 99 and then refrigerated at 6 C until utilization.
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23 101 Digestibility trial

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25 102 A stock of juvenile rainbow trout was obtained from a private hatchery (Bassignana,
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27 103 Cuneo, Italy) and transferred to the facility at the Department of Agriculture, Forestry, and
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29 104 Food Sciences at the University of Torino. An in vivo digestibility experiment was performed
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31 105 in order to determine the apparent digestibility coefficient (ADC) of the diets following the
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33 106 experimental design adopted in a previous study reported by Palmegiano et al. (2006). The
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35 107 ADCs were measured using the indirect acid-insoluble ash (AIA) method; 1% celite® (Fluka,
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37 108 Switzerland) was added to the diets as an inert marker. The faeces were collected from each
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39 109 tank using a continuous automatic device, as reported by Palmegiano et al. (2006), six days
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41 110 per week. The faeces were collected daily and frozen (-20 C) for three consecutive weeks.
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43 111 The faeces were then dried in a stove in order to determine the dry matter (DM) content.
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47 112 The ADC of the DM was calculated as follows:

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$$\text{ADC}_{\text{DM}} (\%) = (1 - A/B) \times 100$$

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52 114 in which A and B represent the AIA concentrations in the feed and faeces, respectively.

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54 115 The ADCs of the crude protein (CP), ether extract (EE) and gross energy (GE) were
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56 116 calculated as follows:
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$$\text{ADCs (\%)} = [1 - (A/B) \times (SB/SA)] \times 100$$

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5 118 in which SA and SB represent the CP, EE or GE concentrations in the feed and faeces,
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7 119 respectively.
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12 121 Growth trial

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14 122 A selection of 1200 juvenile rainbow trout (initial mean body weight 114.6 ± 0.2 g)
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16 123 were individually weighed to obtain a homogeneous stock of fish and randomly distributed
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18 124 into 24 fibre-glass tanks (0.5 m^3) supplied by an open-water circuit with a water flow rate of
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20 125 25 l/min and a temperature of 13 ± 1 C while dissolved oxygen was 7.0 ± 0.5 mg/l.
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23 126 The adopted experimental design was balanced, bi-factorial with four diets x three
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25 127 replicates x two feeding regimes (4x3x2). The feeding trial lasted 77 days, after a 2-week
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27 128 period of acclimatisation to the tanks and diets. The feedstuff was distributed by hand, 6 days
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29 129 per week, twice a day with a daily feeding rate of 1.4% of the wet biomass or ad libitum,
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31 130 respectively. Feed intake was checked each time and no feed reject events were recorded
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33 131 during the trial. The biomass tanks were weighed in bulk every 15 days, in order to update the
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35 132 daily feeding rate.
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40 134 Sampling and chemical analysis

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43 135 At the end of the feeding trial, the fish were starved for one day, then the fish tanks
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45 136 were weighed for final mean body weight. In order to determine the somatic indexes, five
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47 137 trout per tank, with a body weight close to the mean body weight, were sampled and killed.
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49 138 The gut and liver were separated from the rest of the body and weighed. The dorsal muscle
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51 139 tissues from the same fish body were sampled and frozen until the subsequent chemical
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53 140 determinations. The diet and fish muscle samples were freeze-dried before analysis. All the
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55 141 diets were analyzed to determine proximate composition and AIA concentration according to
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3 142 standard methods (AOAC 1990). GE content was determined using an adiabatic calorimetric
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5 143 bomb (IKA C7000, Staufen, Germany).

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9 145 Gas-chromatographic analysis of the fatty acids

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11 146 FA composition was determined on the diets and fish flesh samples. The lipid
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13 147 extraction of the samples was performed according to Peiretti and Meineri (2008); the extract
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15 148 was expressed as crude fat and used for the trans-methylation of the FAs. The FA methyl
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17 149 esters in hexane were then injected into a gas chromatograph (Dani Instruments S.P.A.
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19 150 GC1000 DPC; Cologno Monzese, Italy) equipped with a flame ionization detector (FID). The
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21 151 separation of the FA methyl esters was performed using a Famewax™ fused silica capillary
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23 152 column (30m×0.25mm [i.d.], 0.25 µm) (Restek Corporation, Bellefonte, PA, USA). The peak
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25 153 area was measured using a Dani Data Station DDS 1000. Each peak was identified and
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27 154 quantified on the basis of pure methyl ester standards (Restek Corporation, Bellefonte, PA,
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29 155 USA).

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35 157 pH and Color flesh measurements

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37 158 pH (pH₂₄) was measured on muscles by means of a Crison MicropH 2001 (Crison
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39 159 Instruments, Barcelona, Spain) equipped with a combined electrode and an automatic
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41 160 temperature compensator.

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43 161 The flesh colour measurements were taken on the inside fillet portion using a bench
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45 162 colorimeter Chroma Meter CR-400 Konica Minolta Sensing (Minolta Sensing Inc, Osaka,
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47 163 Japan) in the CIELAB colour space (CIE 1976). The lightness (L*), redness (a*) and
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49 164 yellowness (b*) were recorded. Three readings were taken on each portion of the fillet and
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3 167 Consumer tests
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5 168 A sensory panel of 36 untrained PhD students and staff members from the campus of
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7 169 the University of Torino and of the Italian National Research Council of Torino, 21 males and
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9 170 15 females, ranging in age from 25 to 60 years, participated in this study. Panelists were
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11 171 regular consumers of fish flesh and were already involved in surveys on fish flesh
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13 172 preference/acceptability tests. Consumer tests were carried out in 6 distinct evaluation
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15 173 sessions over three days in the Sensory Evaluation Facility of the Department of Agriculture,
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17 174 Forestry and Food Science of Torino. In each session, a preference ranking test was
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19 175 performed to evaluate the preference of cooked fillets from trout fed with the four
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21 176 experimental diets offered ad libitum or rationed. Between sessions, panelists took a 15 min
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23 177 break. Sixteen trout (two fish from each diet), homogeneous for size and weight, were filleted.
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25 178 The fillets were wrapped in aluminum foil and cooked without additives in an air convection
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27 179 oven at 200 C until the core temperature reached 70 C (about 15 min). After cooking, the
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29 180 fillets were cut into equal portions.
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34 181 Each panelist received four warm samples corresponding to the 4 diets. Samples were
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36 182 labelled with three-digit numbers, and were offered using a Williams design to balance the
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38 183 order of presentation (MacFie et al. 1989).
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40 184 Panelists were asked to rank the samples from trout fed with the four diets in order of
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42 185 preference (most preferred =1; least preferred =4). Tap water was offered to the panelists to
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44 186 rinse their mouths between samples.
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49 188 Statistical analysis
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52 189 Statistical analyses were performed using the SPSS software package (version 11.5.1
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54 190 for Windows, SPSS Inc., USA). Growth performance, FA profile and fillet quality traits were
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56 191 analysed by two-way ANOVA by considering dietary protein source, feeding regime and their
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3 192 interaction as the main effects. The data were presented as the means for each group, together
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5 193 with the significance levels of the main effects and interactions.
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7 194 Analysis of variance was used to evaluate the diet effect on ADC. These data were
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9 195 presented as the means for each group and the standard deviation (SD). Significance was
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11 196 established at $P < 0.05$ for all data.
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14 197 The results of the sensory analysis were analysed by Friedman's test. The Friedman
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16 198 rank sum was performed to determine whether the panellists were able to discriminate
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18 199 between samples. Then, the least significant ranked difference values were calculated to
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20 200 ascertain which samples were significantly preferred to the others (Meilgaard et al. 1991). The
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22 201 ranking data were analysed by box-plots and correspondence analysis.
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25 203 **Results and Discussion**

26 204 Composition and fatty acid profile of the diets

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29 205 The ingredients and chemical composition of the four diets are shown in Table 1,
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31 206 while the FA patterns for the four experimental diets are reported in Table 2.
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34 207 The experimental diets were similar as concerned CP, crude fibre and GE, while the
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36 208 dietary concentration of EE was lower in the BPM diet than in the other diets. Ash and
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38 209 nitrogen-free extracts were higher and lower, respectively, in the FM diet.
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41 210 The concentration of crude fat in the bacterial protein resembles that of fish meal,
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43 211 while the composition of the lipid is different (Storebakken et al. 2004). Phospholipids are the
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45 212 main lipid components of bacterial protein, consisting mainly of phosphatidylethanolamine
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47 213 and phosphatidylglycerol (Müller and Skrede, 2003) with predominantly SFA and MUFA and
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49 214 no PUFA.
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52 215 The experimental diets showed a similar FA profile with slightly high values of
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54 216 C20:5n-3 (EPA) and C22:6n-3 (DHA) in the FM diet. This diet also showed a slightly higher
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3 217 saturated FA (SFA) and PUFA content than the other diets, while the lowest content of
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5 218 MUFA was found in the MIX diet.

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9 220 Digestibility of the experimental diets

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11 221 As far as digestibility is concerned (Table 3), statistical differences appeared among
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13 222 the diets for CP, while no differences appeared for DM, EE and GE. The lowest CP
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15 223 digestibility coefficients were recorded in the BPM and MIX groups, both fed diets containing
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17 224 the bacterial protein meal. Similar ADC of nitrogen was found in studies carried out by
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19 225 Storebakken et al. (2004) and Øverland et al. (2006) in Atlantic salmon and rainbow trout,
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21 226 respectively. These authors recorded lower nitrogen digestibility with increasing BPM
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23 227 inclusion compared to a fish meal-based diet, with values of 87% and 83% in fish fed diets
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25 228 containing 193 and 147 g of bacterial protein meal per kg of diet, respectively. In contrast,
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27 229 even though this was lower than fish meal-based diets, higher values (91 and 88%) of ADC of
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29 230 nitrogen were found in trials with rainbow trout and Atlantic halibut fed diets containing 270
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31 231 and 180 g of bacterial protein meal per kg of diet, respectively (Aas et al. 2006, 2007). The
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33 232 lower nitrogen digestibility values recorded in the BPM and MIX groups of this study and in
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35 233 the other trials utilising bacterial protein meal diets, could be due to a negative effect of
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37 234 bacterial membrane and cell wall components on protein digestibility, as observed in previous
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39 235 studies in rainbow trout fed single-cell proteins from brewer's yeast (Rumsey et al. 1991;
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41 236 Kiessling and Askbrandt 1993). Burel et al. (2000) found that extruded peas showed lower
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43 237 protein digestibility in trout (88%) than in turbot (92%).

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47 239 Growth performance and somatic indexes

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49 240 As far as the growth performance traits and the somatic indexes reported in Table 4
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51 241 are concerned, all the parameters investigated were significantly affected by the diet effect
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3 242 while only the condition factor (CF) was influenced by the feeding rate effect. None of the
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5 243 parameters showed to be affected by the interaction between diet and feeding rate effect.
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7 244 Compared to the FM group, the trout fed alternative protein source diets had lower
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9 245 weight gain and specific growth rates (SGR). Similar results were found by de Francesco et
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11 246 al. (2004) in a long-term feeding study where large rainbow trout were fed with a plant
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13 247 protein mixture-based diet. In contrast, de Francesco et al. (2007) found similar weight gain
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15 248 and SGR in gilthead sea bream fed with a plant protein high-level fish meal replacement diet.
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17 249 In our trial, trout fed alternative protein source diets were characterised by a higher viscerosomatic
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19 250 and hepato-somatic index but a lower carcass yield. A decrease in dressed carcass and fillet
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21 251 yield was also observed in a study carried out in trout fed plant proteins and guar gum as fish
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23 252 meal replacements (Brinker and Reiter, 2011).
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27 253 A similar effect on CF was observed in a study carried out in Atlantic salmon fed to
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29 254 satiation or moderately reduced rations of high or low energetic feeds, in which the rationed
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31 255 fish showed the lowest CF values (Johnsen et al. 2011).
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36 257 Fatty acid profile of the fillet

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38 258 While it is common to see changes in the FA profile when dietary fat is modified
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40 259 through changes in dietary lipid sources, there is little information as regards the effects of
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42 260 changes in FA content as affected by dietary protein sources. Indeed, total replacement of fish
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44 261 meal by plant protein ingredients modifies FA profiles to a certain extent with the consequent
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46 262 changes seen in muscle FA profiles (Tables 5 and 6). As far as FA composition of different
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48 263 fillets is concerned, C18:1n-9, C18:2n-6, C18:3n-3, and DHA contents and some minor FAs
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50 264 (such as C15:0, C16:2n-4, C17:0, C16:3n-4, C18:3n-6, C20:1n9, C20:4n-6, C21:0 and
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52 265 C22:1n-9) were influenced by diet, while only minor FAs (such as C16:1n-9, C16:2n-4,
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3 266 C16:3n-4, C18:3n-4, C20:1n9 and C22:1n-9) were influenced by feeding regime (Table 5).
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5 267 An interaction was found only for two minor FAs (C16:2n-4 and C22:1n-9).
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7 268 In common with other studies on rainbow trout (De Francesco et al. 2004; Morris et al.
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9 269 2005) the FA content of trout muscle in the present study was significantly influenced by that
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11 270 of the feed. In fact, it is well known that dietary FA composition strongly influences flesh FA
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13 271 composition in fish (Sargent et al. 2002). As shown by Palmegiano et al. (2006), who
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15 272 evaluated the use of rice protein concentrate as a potential substitute of fish meal in rainbow
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17 273 trout, fillet FA profile reflects diet composition, but some FAs are not present in the same
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19 274 proportion and this induces one to suppose that an elongation and desaturation process has
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21 275 occurred. Numerous FAs were present at higher proportions in the flesh lipids than in the
22
23 276 feeds, including C18:1n-9 and DHA. However, C18:2n-6, C18:3n-3 and EPA were all present
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25 277 at lower relative percentages in the flesh than in the feeds. Preferential accumulation and/or
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27 278 retention of selected FAs, including C18:2n-6, C20:4n-6 and DHA, has previously been
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29 279 recorded in rainbow trout (Greene and Selivonchick 1990).
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34 280 In the present study, MUFA and PUFA content (total PUFA, PUFA n-3, PUFA n-6
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36 281 and their ratio) were influenced by diet treatment, while only MUFA content was influenced
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38 282 by feeding regime without interaction between factors (Table 6), as previously demonstrated
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40 283 in rainbow trout fed with a plant protein mixture-based diet (De Francesco et al. 2004). They
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42 284 reported that SFA, MUFA and PUFA n-3 and the n-3/n-6 ratio were significantly higher in
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44 285 trout fed diet based on fish meal, while PUFA n-6 (above all in C18:2n-6) were significantly
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46 286 higher in trout fed diet based on mixture of plant protein sources (corn gluten meal, wheat
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48 287 gluten, extruded peas, and rapeseed meal), while the main difference observed in single FA
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50 288 was the higher incidence and content of C18:3n-3 in fillet of trout fed diet based on mixture of
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52 289 plant protein sources in comparison to those fed the fish meal, no differences were found for
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54 290 eicosapentaenoic (EPA) and the docosahexaenoic (DHA) acid levels. This result was
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3 291 consistent with the data obtained by Gomes et al. (1993), who observed an increased level of
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5 292 PUFA n-6, in particular of C18:2n-6, in muscle of rainbow trout fed diets with increasing
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7 293 levels (5%, 10%, 15% and 45%) of a co-extruded plant protein (rapeseed and peas). It is
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9 294 reassuring to note that at the same lipid level, EPA and DHA were no different between the
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11 295 two groups of trout.

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14 296 Morris et al. (2005) reported that, with the exception of five individual FAs (C16:2n-6;
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16 297 C20:4n-6; C20:4n-3; DHA and C24:1n-9), the FA profile of the rainbow trout flesh responded
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18 298 linearly to changing proportions of individual FAs in the feed formulated with extracted soya
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20 299 (7.5%) and full-fat soya (0–25%). Although the percentage of DHA and PUFA n-3 in the
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22 300 fillet was not significantly influenced by the level of soya in the feeds, the relative proportions
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24 301 of the fish and soybean derived PUFA n-3 shifted towards the latter, i.e. higher relative
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26 302 percentages of C18:3n-3, in response to a higher proportion of soya-derived fat in the feeds.

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29 303 The challenges of creating new plant-based feed ingredients for salmonid diets are
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31 304 providing high-quality protein and providing a source of PUFA n-3 (Drew et al. 2005). Fish
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33 305 oil is the most widely-used source of PUFA n-3, required by salmonids to maximize growth
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35 306 potential and maintain the PUFA n-3 content of the fish carcass desired by consumers.
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37 307 Rainbow trout can elongate and desaturate C18:3n-3 into EPA and DHA (Owen et al. 1975),
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39 308 but most plant oils are poor sources of C18:3n-3.

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44 45 310 Fillet quality traits

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47 311 Parameters of fillet pH and colour are reported in Table 7; all the parameters
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49 312 investigated were significantly affected by the treatments except redness for the diet effect
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51 313 and yellowness for the feeding rate effect, respectively. None of the parameters appeared to
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53 314 be affected by the interaction between diet and feeding rate effect.
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3 315 The fillet pH level after 24h was slightly higher for fish fed BPM and MIX diets while
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5 316 similar values were found for the FM and PPC groups. Brinker and Reiter (2011) observed a
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7 317 reduction in pH 24h post mortem in fillets of trout fed a mixture of fish meal and plant
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9 318 protein-based feeds and they attribute the observed reduction in pH to the differences in fillet
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11 319 energy stores. The same authors found that the pure plant diet appears to increase undesirable
12
13 320 yellowness in the trout fillets, in agreement with the results of the present trial where we
14
15 321 found a similar trend in fish fed the PPC and MIX diets.

16
17 322 In our trial, the feeding rate modified the redness values of the trout fillets with the
18
19 323 higher values recorded in rationed fish while satiation feeding induced higher red colour
20
21 324 intensity (a*-value) compared to restricted feeding in a study carried out in Atlantic salmon
22
23 325 (Johnsen et al. 2011). The same authors did not find any differences in the lightness values, in
24
25 326 contrast to the findings from the present trial where lightness values decreased in fillets from
26
27 327 the rationed groups.

28
29 328 Fillet composition in terms of DM, CP, EE, and ash content is reported in Table 8. The
30
31 329 results for fillet composition showed that only CP content was affected by the diet effect with
32
33 330 an increased content in fish fed fish meal-alternative protein sources. A similar increase was
34
35 331 also observed by De Francesco et al. (2004) in large rainbow trout fed with plant protein
36
37 332 mixtures in replacement of fish meal. The same effect was also reported in rainbow trout fed
38
39 333 diets where 25, 50, 75, or 100% of the fish meal protein was replaced with a mixture of
40
41 334 rendered animal protein ingredients (Lesiow et al. 2009). DM and ash content were affected
42
43 335 by feeding rate effect, fillets from rationed fish groups showed an increased content of DM
44
45 336 and a decreased ash content compared to ad libitum groups.

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Consumer tests

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3 339 The results of the preference ranking test concerning the trout fillets fed ad libitum
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5 340 with the four diets are reported in Fig. 1. Fillets of trout fed the MIX diet ad libitum obtained
6
7 341 the highest number of most preferred votes as well as the best median value (Fig. 2). Fillets of
8
9 342 trout fed the FM diet ad libitum obtained the largest number of least preferred and the
10
11 343 smallest number of most preferred votes. Fillets of trout fed the BPM and PPC diets ad
12
13 344 libitum had the majority of votes in 2nd and 3th preference votes, 75% and 53% respectively.

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15
16 345 The Friedman's test showed that there was a significant difference ($P \leq 0.05$) in
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18 346 preference between the fillets (Table 9). Fillets of trout fed the MIX diet ad libitum were the
19
20 347 most preferred (rank sum = 72), followed by the BPM diet ad libitum (rank sum = 89), PPC
21
22 348 diet ad libitum (rank sum = 90) and FM diet ad libitum (rank sum = 109).

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24
25 349 A similar ranking was obtained with the rationed trout, although no significant
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27 350 difference in preference between the four fillets was observed (Table 9).

28
29 351 Fillets of trout fed the rationed MIX diet obtained the highest number of most
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31 352 preferred votes (Fig.3). Fillets of trout fed the rationed PPC diet obtained the same number of
32
33 353 most and least preferred votes. The rationed BPM diet had the majority of 2nd preference
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35 354 votes while the rationed FM diet received the largest number of least preferred votes. The
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37 355 rationed FM and PPC diets showed the worst median values (Fig. 2).

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40 356 The overall results can be represented in the correspondence analysis plot (Fig. 4).
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42 357 Dimensions 1 and 2 explain 58% and 39% of the inertia, respectively.

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45 358 Fillets of trout fed the MIX diet ad libitum and 1st preference votes, and, fillet of trout
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47 359 fed the rationed FM diet and 4th preference votes, showing the highest deviation from the
48
49 360 origin, gave the main contribution to the inertia of dimension 1 and dimension 2, respectively.
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51 361 According to the distribution of samples in the plane, it can be seen that the samples preferred
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53 362 by the consumers are found to the right in the plot and near the 1st preference votes. The next
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55 363 two groups, rationed BPM diet and BPM diet ad libitum, are found on the left near the 2nd

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3 364 preference votes. Less- preferred fillets were from FM diet ad libitum and rationed FM diet
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5 365 near the 4th preference votes, while rationed PPC diet and PPC diet ad libitum, near the 3th
6
7 366 preference votes, are found between the 1st and 4th preference votes.
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506 TABLE 1. Ingredients and proximate composition of experimental diets.

Diets	FM	BPM	PPC	MIX
<u>Ingredients (%)</u>				
Herring fish meal ^a	50.00	25.00	25.00	25.00
Protorsan ^b	0.00	25.00	0.00	12.50
Pea protein concentrate ^c	0.00	0.00	30.00	15.00
Corn meal	23.00	24.00	12.00	17.50
Fish oil	10.00	11.00	12.50	12.00
Corn gluten	8.00	6.00	12.00	9.00
Lignum sulphate	6.00	6.00	6.00	6.00
Mineral mixture ^d	1.50	1.50	1.25	1.50
Vitamin mixture ^e	1.50	1.50	1.25	1.50
<u>Proximate composition (%DM)</u>				
Dry matter (% fresh matter)	96.6	92.5	95.9	96.1
Crude protein	45.4	45.9	45.2	45.2
Ether extract	17.3	14.6	16.5	17.0
Ash	12.0	8.9	9.2	9.2
Crude fiber	1.9	2.0	1.9	1.9
Nitrogen free extracts ^f	23.4	28.6	27.2	26.7
Gross energy (MJ/kg DM) ^g	21.5	21.8	22.2	22.1

507 ^a Mangimi Monge, Torre San Giorgio, Italy: DM 91.2%, CP 69%, EE 8.2%, ash 9.6%, CF 0.5%.

508 ^b Mazzoleni Prodotti Zootecnici, Cologno al Serio, Italy: DM 92%, CP 67%, EE 6%, ash 3.8%, CF 1%.

509 ^c AgriMarin Nutrition, Stavanger, Norway: DM 90%, CP 55%, starch 9%, EE 2%, ash 6%.

510 ^d Mineral mixture (g or mg/kg diet): bicalcium phosphate 500 g, calcium carbonate 215 g, sodium salt
 511 40 g, potassium chloride 90 g, magnesium chloride 124 g, magnesium carbonate 124 g, iron sulphate 20 g, zinc

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3 512 sulphate 4 g, copper sulphate 3 g, potassium iodide 4 mg, cobalt sulphate 20 mg, manganese sulphate 3 g,
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5 513 sodium fluoride 1g, (Granda Zootechnica, Cuneo, Italy).

6 514 ^e Vitamin mixture (IU or mg/kg diet): DL-a tocopherol acetate, 60 IU; sodium menadione bisulphate, 5
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8 515 mg; retinyl acetate, 15000 IU; DL-cholecalciferol, 3000 IU; thiamin, 15 mg; riboflavin, 30 mg; pyridoxine, 15
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10 516 mg; B12, 0.05 mg; nicotinic acid, 175 mg; folic acid, 500 mg; inositol, 1000 mg; biotin, 2.5 mg; calcium
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12 517 panthotenate, 50 mg; choline chloride, 2000 mg (Granda Zootechnica, Cuneo, Italy).

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14 518 ^f Calculated as 100-(%Crude protein +%Ether extract +%Ash +%Crude fiber).

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16 519 ^g Determined by calorimetric bomb.

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For Peer Review

521 TABLE 2. Dietary main fatty acid (% of total fatty acid) composition.

	FM	BPM	PPC	MIX
C14:0	4.70	4.66	4.19	4.56
C15:0	0.34	0.32	0.30	0.32
C16:0	13.39	13.15	12.08	13.00
C16:1n-9	0.20	0.21	0.18	0.21
C16:1n-7	5.13	5.23	4.79	5.17
C16:2n-4	0.62	0.23	0.15	0.15
C17:0	0.27	0.24	0.24	0.24
C16:3n-4	0.69	0.69	0.71	0.61
C17:1n-7	0.18	0.17	0.19	0.22
C18:0	3.57	3.27	3.14	3.15
C18:1n-9	19.53	21.90	20.79	22.09
C18:1n-7	2.86	2.81	2.70	2.81
C18:2n-6	9.56	10.85	11.03	11.20
C18:3n-6	0.21	0.23	0.34	0.19
C18:3n-4	0.16	0.18	0.52	0.18
C18:3n-3	2.36	2.62	2.99	2.83
C18:4n-3	1.66	1.69	1.70	1.71
C20:0	0.26	0.26	0.25	0.30
C20:1n-9	2.24	2.49	2.55	2.66
C20:2n-6	0.44	0.48	0.49	0.47
C20:3n-3	0.14	0.15	0.41	0.11
C20:4n-6	0.67	0.59	0.62	0.62

C21:0	0.22	0.23	0.24	0.11
C20:4n-3	0.81	0.85	0.81	0.82
C20:5n-3	9.55	9.00	8.51	8.73
C22:1n-9	1.68	1.90	1.87	1.90
C22:2n-6	0.45	0.43	1.69	0.70
C22:5n-3	1.91	1.88	1.79	1.84
C22:6n-3	11.01	8.80	9.02	8.80
SFA ^a	22.74	22.14	20.44	21.69
MUFA ^b	31.83	34.72	33.07	35.05
PUFA ^c	39.81	38.24	39.11	38.27
PUFA n-3 ^d	27.44	24.99	25.23	24.85
PUFA n-6 ^e	11.34	12.58	14.19	13.17
n-3/n-6	2.42	1.99	1.78	1.89

522 ^a Saturated fatty acids.

523 ^b Monounsaturated fatty acids.

524 ^c Polyunsaturated fatty acids.

525 ^d Polyunsaturated fatty acids serie n-3.

526 ^e Polyunsaturated fatty acids serie n-6.

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528 TABLE 3. Apparent digestibility coefficients (%) of nutrients and gross energy of the
529 experimental diets (means \pm SD; n=9).

Diets	FM	BPM	PPC	MIX
Dry matter	69.97 \pm 0.67	69.23 \pm 0.25	70.33 \pm 1.29	67.33 \pm 1.92
Crude protein	90.30 \pm 0.28 ^a	84.20 \pm 0.28 ^b	88.90 \pm 0.28 ^a	85.40 \pm 1.13 ^b
Ether extract	97.35 \pm 0.78	96.90 \pm 0.28	95.70 \pm 0.28	96.35 \pm 0.21
Gross energy	79.93 \pm 1.62	77.00 \pm 2.26	78.50 \pm 3.04	74.87 \pm 2.97

530 In the row, different letters mean statistical difference at $P \leq 0.05$.

531 TABLE 4. Growth performance, feed utilization and whole body composition (n=5) in rainbow trout fed experimental diets.

	<u>ad libitum</u>				Rationed				Significance		
	FM	BPM	PPC	MIX	FM	BPM	PPC	MIX	Diet effect	F.R. effect	Interaction
WG ^a	113.5	39.5	99.4	102.5	100.4	40.6	99.8	91.8	0.000	0.082	0.250
SGR ^b	0.90	0.40	0.80	0.80	0.85	0.40	0.80	0.77	0.000	0.360	0.815
PER ^c	1.63	0.67	1.75	1.63	1.60	0.80	1.77	1.57	0.000	0.790	0.459
FCR ^d	1.40	3.53	1.25	1.33	1.40	2.70	1.23	1.40	0.000	0.246	0.213
VSI ^e	10.8	12.4	11.7	13.0	9.8	10.8	11.5	12.3	0.025	0.095	0.790
HSI ^f	1.02	1.17	1.08	1.40	1.04	1.13	1.20	1.25	0.007	0.803	0.418
CF ^g	1.20	1.15	1.26	1.26	1.09	0.99	1.18	1.22	0.002	0.005	0.577
CY ^h	88.2	86.4	87.2	85.6	89.1	88.1	87.3	86.5	0.015	0.107	0.774

532 ^a Weight gain, (g) =[FBW (final body weight, g) –IBW (initial body weight, g)].533 ^b Specific growth rate (%) =[lnFBW–lnIBW)/number of feeding days]*100.534 ^c Protein efficiency ratio =[WG (weight gain,g)/total protein fed (g DM)].535 ^d Feed conversion ratio=[total feed supplied (g DM)/WG (weight gain, g)].536 ^e Viscerosomatic index, (%) =[gut weight (g)/ fish weight (g)]*100.

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537 ^f Hepatosomatic index, (%) = [liver weight (g)/ fish weight (g)]*100.

538 ^g Condition factor, (%) = [fish weight (g)/ fish length (cm)³]*100.

539 ^h Carcass yield, (%) = [carcass weight (g)/ fish weight (g)]*100.

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For Peer Review

541 TABLE 5. Fatty acids in fillets (n=5): composition (% of total fatty acid).

	<u>ad libitum</u>				Rationed				Significance		
	FM	BPM	PPC	MIX	FM	BPM	PPC	MIX	Diet effect	F.R. effect	Interaction
C14:0	3.35	3.33	2.94	3.47	3.17	3.11	3.15	3.23	0.375	0.374	0.523
C15:0	0.28	0.29	0.24	0.28	0.27	0.27	0.25	0.27	0.026	0.410	0.482
C16:0	14.60	14.70	14.06	15.06	14.56	14.34	14.82	14.50	0.618	0.783	0.085
C16:1n-9	0.25	0.24	0.24	0.26	0.14	0.25	0.23	0.19	0.332	0.032	0.136
C16:1n-7	4.28	4.20	3.83	4.39	4.01	4.06	4.01	3.94	0.590	0.231	0.484
C16:2n-4	0.27	0.21	0.39	0.41	0.21	0.15	0.14	0.17	0.000	0.000	0.000
C17:0	0.28	0.30	0.21	0.28	0.28	0.27	0.26	0.27	0.005	0.663	0.053
C16:3n-4	0.25	0.24	0.27	0.29	0.18	0.14	0.12	0.21	0.010	0.000	0.142
C18:0	3.29	3.26	3.14	3.37	3.20	3.40	3.21	3.33	0.279	0.805	0.611
C18:1n-9	19.23	20.06	18.96	20.30	18.19	19.57	17.59	19.95	0.022	0.082	0.849
C18:1n-7	2.80	2.68	2.54	2.05	2.68	2.64	2.53	2.63	0.300	0.514	0.387
C18:2n-6	12.01	13.79	16.13	12.88	12.41	14.71	15.82	13.11	0.001	0.603	0.904
C18:3n-6	0.27	0.34	0.35	0.29	0.30	0.29	0.34	0.31	0.005	0.714	0.117

C18:3n-4	0.34	0.38	0.40	0.37	0.29	0.28	0.29	0.31	0.731	0.000	0.576
C18:3n-3	1.69	2.11	1.93	2.00	1.70	2.05	1.71	1.91	0.000	0.068	0.400
C18:4n-3	0.99	1.14	0.95	1.05	0.93	1.04	1.00	1.04	0.101	0.441	0.586
C20:0	0.08	0.19	0.14	0.19	0.18	0.19	0.16	0.12	0.371	0.640	0.158
C20:1n-9	2.11	2.02	1.64	2.19	1.69	2.02	1.34	1.94	0.001	0.019	0.482
C20:2n-6	0.69	0.67	0.76	0.71	0.65	0.74	0.64	0.71	0.688	0.454	0.082
C20:3n-3	0.38	0.37	0.50	0.39	0.56	0.47	0.48	0.38	0.554	0.259	0.503
C20:4n-6	0.70	0.63	0.62	0.65	0.70	0.63	0.64	0.64	0.001	0.796	0.909
C21:0	0.21	0.21	0.18	0.22	0.15	0.17	0.04	0.21	0.006	0.003	0.162
C20:4n-3	0.95	0.93	0.91	1.01	0.96	0.92	0.84	0.85	0.299	0.049	0.188
C20:5n-3	5.59	5.77	5.17	5.59	5.48	5.39	5.56	5.44	0.749	0.727	0.479
C22:1n-9	1.04	1.13	0.76	1.10	0.82	0.29	0.21	0.28	0.000	0.000	0.001
C22:2n-6	0.46	0.48	0.34	0.40	0.91	0.38	0.37	0.38	0.210	0.466	0.366
C22:5n-3	2.15	2.11	2.07	2.11	2.15	2.03	2.24	1.98	0.293	0.810	0.139
C22:6n-3	18.60	15.44	17.38	16.31	19.77	16.08	18.32	17.46	0.013	0.189	0.993

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543 TABLE 6. Relationship of fatty acids in fillets (n=5): composition (% of total fatty acid) to nutritional quality.

	<u>ad libitum</u>				Rationed				Significance		
	FM	BPM	PPC	MIX	FM	BPM	PPC	MIX	Diet effect	F.R. effect	Interaction
SFA ^a	22.09	22.28	20.90	22.87	21.82	21.73	21.90	21.91	0.147	0.504	0.126
MUFA ^d	29.71	30.34	27.97	30.30	27.52	28.82	25.91	28.94	0.049	0.019	0.971
PUFA ^c	44.87	44.12	47.84	44.05	46.30	44.92	48.16	44.51	0.004	0.299	0.945
PUFA n-3 ^d	30.34	27.87	28.91	28.46	31.56	27.97	30.15	29.05	0.037	0.294	0.934
PUFA n-6 ^e	14.13	15.91	18.21	14.93	14.96	16.76	17.82	15.14	0.002	0.536	0.868
n-3/n-6	2.18	1.76	1.64	1.91	2.15	1.69	1.70	1.93	0.003	0.952	0.962

544 ^a Saturated fatty acids.545 ^b Monounsaturated fatty acids.546 ^c Polyunsaturated fatty acids.547 ^d Polyunsaturated fatty acids series n-3.548 ^e Polyunsaturated fatty acids series n-6.

550 TABLE 7. Parameters of fillet pH (n=12) and colour (n=24) in rainbow trout fed experimental diets.

	<u>ad libitum</u>				Rationed				Significance		
	FM	BPM	PPC	MIX	FM	BPM	PPC	MIX	Diet effect	F.R. effect	Interaction
pH ₂₄	6.56	6.76	6.58	6.63	6.65	6.74	6.67	6.75	0.000	0.000	0.063
L ^a	44.9	46.4	47.8	48.9	44.2	45.5	45.6	44.8	0.004	0.000	0.051
a ^b	1.14	1.84	1.00	0.84	1.84	1.83	1.33	1.84	0.066	0.007	0.234
b ^c	5.08	3.12	6.71	6.33	4.33	3.14	7.19	5.53	0.000	0.435	0.460

551 ^a Lightness.

552 ^b Redness.

553 ^c Yellowness.

555 TABLE 8. Composition (% dry matter) of fillets (n=5).

	<u>ad libitum</u>				Rationed				Significance		
	FM	BPM	PPC	MIX	FM	BPM	PPC	MIX	Diet effect	F.R. effect	Interaction
Dry matter ^a	24.1	23.0	23.4	24.5	25.9	24.3	26.9	24.8	0.480	0.017	0.451
Crude protein	89.7	92.3	90.3	90.2	88.5	92.0	88.1	88.9	0.021	0.085	0.836
Ether extract	8.0	14.5	10.8	8.6	9.8	7.7	10.1	10.6	0.440	0.357	0.018
Ash	6.7	7.6	7.5	7.3	6.3	6.5	6.1	6.8	0.056	0.000	0.076

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557 TABLE 9. Preferences of flesh expressed as Rank sums.

	<u>ad libitum</u>				Rationed			
	MIX	BPM	PPC	FM	MIX	BPM	PPC	FM
Rank	1	2	3	4	1	2	3	4
Rank sum	72a	89ab	90ab	109b	79	90	91	100

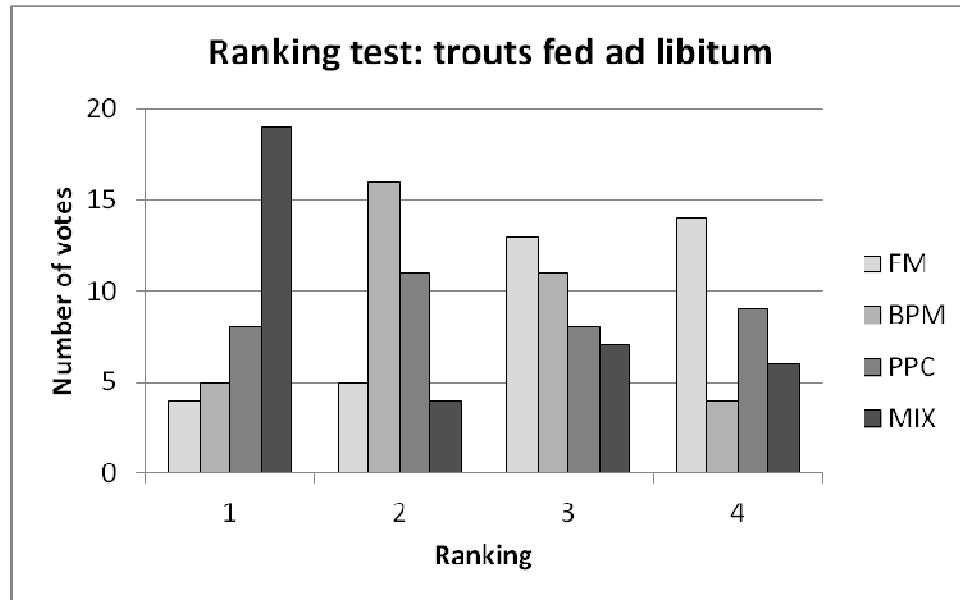
558 Rank sum of the preference ranking test for each trout flesh.

559 Rank sums with different superscripts indicate significant differences among

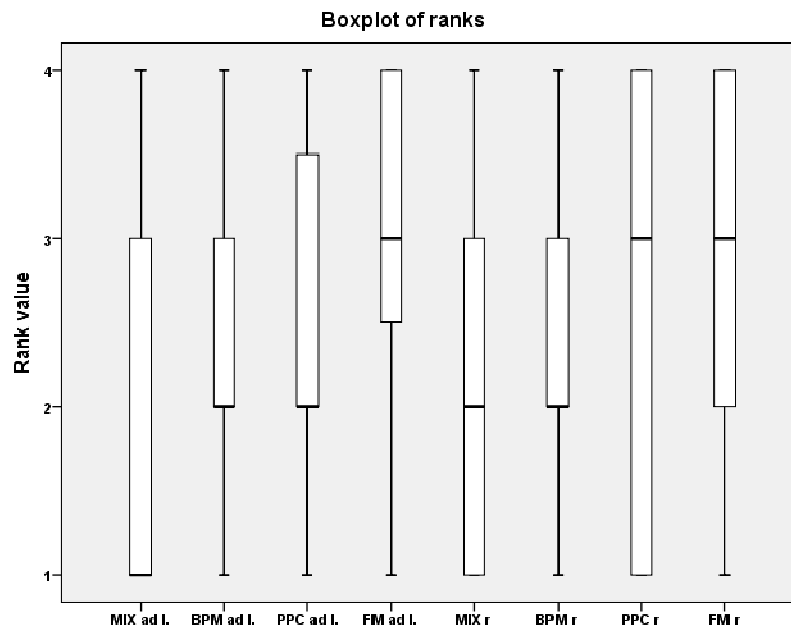
560 treatments ($P \leq 0.05$).

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3 562 FIGURE 1. Results from ranking test of fillets from trout fed ad libitum (where 1 = most
4 preferred and 4 = least preferred). Legend: FM = Fish meal diet; BPM = Bacterial protein
5 563 preferred and 4 = least preferred). Legend: FM = Fish meal diet; BPM = Bacterial protein
6 diet; PPC = Pea protein diet; MIX = Bacterial protein diet+Pea protein diet.
7 564 diet; PPC = Pea protein diet; MIX = Bacterial protein diet+Pea protein diet.
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3 567 FIGURE 2. Box-plot of ranks. Legend: Feeding regime ad libitum: FM ad l. = Fish meal diet;
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5 568 BPM ad l. = Bacterial protein diet; PPC ad l. = Pea protein diet; MIX ad l. = Bacterial protein
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7 569 diet+Pea protein diet. Rationed feeding regime: FM r =Fish meal diet; BPM r =Bacterial
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9 570 protein diet; PPC r = Pea protein diet; MIX r = Bacterial protein diet+Pea protein diet.
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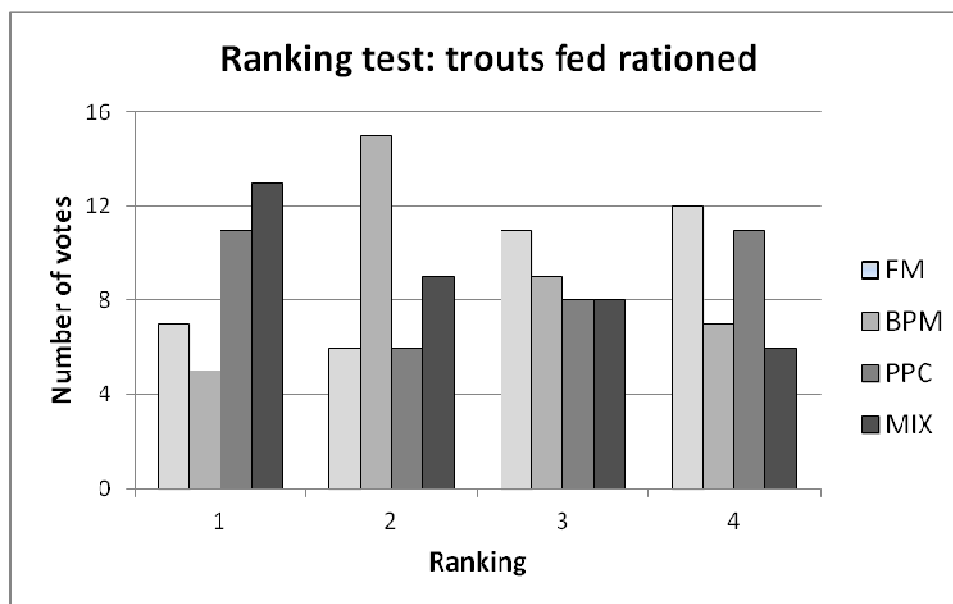


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573 FIGURE 3. Results from ranking test of fillets from trout fed rationed diets (where 1 = most
574 preferred and 4 = least preferred). Legend: FM = Fish meal diet; BPM = Bacterial protein
575 diet; PPC = Pea protein diet; MIX = Bacterial protein diet+Pea protein diet.

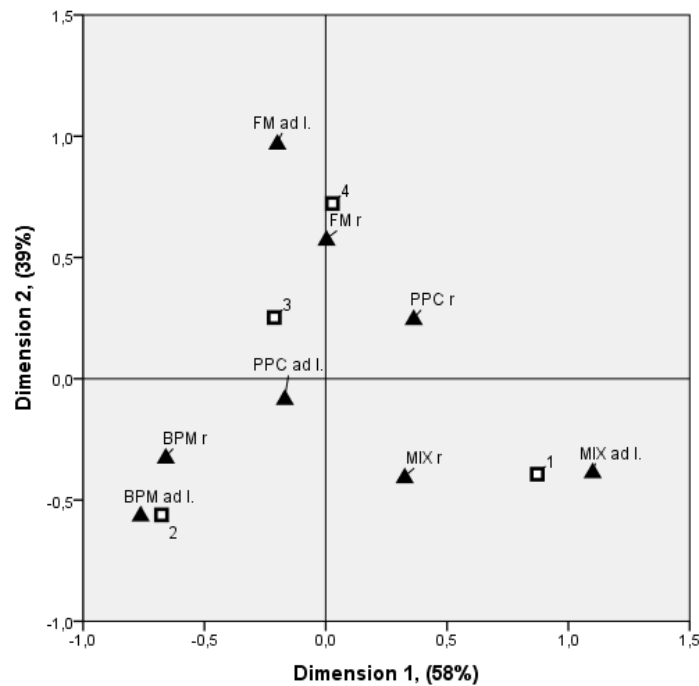
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579 FIGURE 4. Correspondence analysis of sensory data. Position of the fillets of trout and ranks
 580 in the plane formed by the first two dimensions. . Legend: Feeding regime ad libitum: FM ad
 581 l. = Fish meal diet; BPM ad l. = Bacterial protein diet; PPC ad l. = Pea protein diet; MIX ad l.
 582 = Bacterial protein diet+Pea protein diet. Rationed feeding regime: FM r =Fish meal diet;
 583 BPM r =Bacterial protein diet; PPC r = Pea protein diet; MIX r = Bacterial protein diet+Pea
 584 protein diet.



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