


Effects of dietary protein and essential amino acid deficiencies on growth, body composition, and digestive enzyme activities of silvery-black porgy (*Sparidentex hasta*)

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Abstract A 6-week feeding trial was conducted to evaluate the effects of protein-free (PF) and essential amino-acid-deficient (EAAD) diets on the growth performance and digestive enzyme activities of silvery-black porgy (*Sparidentex hasta*) juveniles. Three experimental diets were formulated: a control diet in which 60% of dietary nitrogen was provided by intact protein (fish meal) and 40% by crystalline AA [(blends of essential amino acids (EAA) and none essential amino acids (NEAA)]; an essential amino-acid-deficient diet in which 60% of dietary N was provided by intact protein, whereas the rest was provided by NEAA and a protein-free (PF) diet, based on carbohydrate sources. Survival rates in fish fed the EAAD and PF diets were lower than in fish fed the control diet. Weight gain in fish fed with EAAD was 5.6 g lower than fish fed with the control diet. Furthermore, fish fed the PF diet lost weight and their final body weight was 12.2 g lower than the control group. Fish fed the PF diet had the highest whole-body moisture, but the lowest whole-body protein, lipid, threonine, aspartic acid, glutamic acid, and serine levels. Digestive pancreatic enzyme activities including trypsin, lipase, α -amylase, and carboxypeptidase A significantly decreased in fish fed EAAD and PF diets. The information obtained from this study testing two extreme diets (EAAD and PF) may serve for better understanding the impact of protein and EAA nutritional imbalances in fish performance.

Keywords Amino acid profile · Feed efficiency · Nitrogen retention · Sparidae · Digestive enzymes

Introduction

The optimization of the dietary protein and essential amino acid (EAA) profile according to the requirements of a given fish species maximizes dietary EAA and non-EAA utilization and increases protein synthesis resulting in a proper fish growth and economic return. An absolute requirement for the 10 EAA (arginine,

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histidine, isoleucine, leucine, lysine, methionine, phenylalanine, threonine, tryptophan, and valine) has been demonstrated in most fish species studied to date (NRC 2011). Essential amino acids are key molecules for building proteins, as well as important regulators of key metabolic pathways including cell signaling, appetite stimulation, growth and development, energy utilization, immunity, osmoregulation, ammonia detoxification, antioxidative defense, metamorphosis, pigmentation, gut and neuronal development, stress responses, reproduction, and suppression of aggressive behavior in aquatic animals (Wu et al. 2013, 2014). In addition, the previous studies have reported that different EAA have a significant role on the digestive enzyme activities in fish (Tang et al. 2009, 2013, Chen et al. 2012, Li et al. 2015). In fact, the positive effects of dietary EAA supplementation on the activities of digestive enzymes may partially be attributed to the growth and structural integrity of pancreas and intestinal brush borders as it has been reported in the above-mentioned studies.

Silvery-black porgy (*Sparidentex hasta*) is recognized as one of the most promising fish species for aquaculture diversification in the Persian Gulf and the Oman Sea regions. In this regard, during recent years, nutritional studies focused on establishing the nutritional requirements for improving diet formulation for this species (Mozanzadeh et al. 2017). Previous studies have shown that the dietary protein requirement of *S. hasta* juvenile is estimated to be 48%, being fish meal the major source of protein in the diet (Mozanzadeh et al. 2017). Moreover, the optimal dietary EAA profile for *S. hasta* juveniles was estimated to be (g 16 g/N): arginine 5.3, lysine 6.0, threonine 5.2, histidine 2.5, isoleucine 4.6, leucine 5.4, methionine + cysteine 4.0 (in a diet containing 0.6 cysteine), phenylalanine + tyrosine 5.6 (in a diet containing 1.9 tyrosine), tryptophan 1.0, and valine 4.6 (Marammazi et al. 2017). In addition, Yaghoubi et al. (2017) reported that arginine, threonine, and lysine were the most limited EAA for humoral immunity in this species. Several studies used protein-free diets (PF) for estimating nitrogen utilization and loss as well as protein and EAA requirements for maintenance in different fish species (Kaushik et al. 1995, Abboudi et al. 2006, 2009). To our knowledge, there is no available information about the effects of dietary protein and EAA deficiencies on physiological responses in this sparid species. Thus, the present study aimed to evaluate effects of protein and EAA deficiencies on the growth performance, feed utilization, body composition, and pancreatic digestive enzyme activities to clarify how protein and EAA deficiencies can affect physiological responses in this species.

Materials and methods

Experimental diets

Three diets were formulated (Tables 1, 2): a control diet (74.7 g N/kg DM), a diet deficient in essential amino acids (EAAD 76.6 g N/kg DM), and a protein-free diet (PF 7.5 g N/kg DM). In the control diet, 60% of dietary N was provided by intact protein (fishmeal and gelatine) and 40% by crystalline AAs (CAA). The CAA mixture was prepared by a blend of EAA and non-essential amino acids (NEAA), which were coated with 1% agar in order to delay their absorption and to optimize their use for protein accretion (Peres and Oliva-Teles 2006). All the EAA and NEAA incorporated in the CAA mixture of the control diet simulated the AA profile of FM from *Clupeonella* sp. and matched the EAA nutritional requirements for this species (Marammazi et al. 2017). Agar was dissolved in boiled water (10 g/kg of the diet) and cooled to 40 °C before the addition of CAAs (Peres and Oliva-Teles 2006). Regarding the EAAD diet, 60% of dietary N was provided by fish meal, whereas the rest was incorporated as a mixture of NEAA (crystalline form). The PF diet was based on wheat middling, corn starch, and fish oil. In all diets, ingredients were finely ground (particle size: < 500 µm), mixed together in a Hobart mixer, and then, the pre-coated CAA mixture was added, followed by the fish oil and sufficient distilled water to form a soft dough that was extruded to obtain pellets of 2 mm in size. Pellets were dried in a convection oven at 25 °C and stored in re-sealable plastic bags at – 20 °C until their use.

Feeding trial

Experimental procedures were conducted in compliance with the guidelines of the international council for laboratory animals science for the use of laboratory animals. The study was carried out at the Mariculture Research Station of the South Iranian Aquaculture Research Center (SIARC) (Khuzestan, Sarbandar, Iran).

Table 1 Ingredient and proximate composition of the control, essential amino-acid-deficient (EAAD), and protein-free (PF) diets

	Diets		
	Control	EAAD	PF
Dietary ingredients (g kg ⁻¹ dry diet)			
Fish meal ^a	360	360	–
Gelatin ^b	40	40	–
Wheat middling ^c	70	70	275
Corn starch ^d	205	205	540
Fish oil ^e	110	110	155
Agar ^f	10	10	10
Vitamin Premix ^g	10	10	10
Mineral Premix ^h	10	10	10
L-arginine	8.5	–	–
L-lysine-HCl	11.5	–	–
L-threonine	8	–	–
L-histidine	5	–	–
L-isoleucine	8.5	–	–
L-leucine	14	–	–
L-methionine	6	–	–
L-phenylalanine	7.5	–	–
L-tryptophan	2	–	–
L-valine	9.5	–	–
NEAA mixture ⁱ	104.5	185	–
Proximate composition (%)			
Dry matter	92.9 ± 0.4	93.2 ± 0.1	91.5 ± 0.3
Crude protein	46.7 ± 0.6	47.9 ± 0.1	4.7 ± 2.5
Crude lipid	20.1 ± 0.0	20.0 ± 0.0	19.9 ± 0.6
NFE ^j	19.1 ± 0.5	18.3 ± 0.2	64.3 ± 1.7
Ash	6.8 ± 0.1	6.1 ± 0.1	1.6 ± 0.0
Gross energy (kJ g ⁻¹) ^k	20.9 ± 0.1	20.1 ± 0.0	19.5 ± 0.0

^aFish meal (*Clupeonella* sp.); Parskilka Mazandaran, Iran (63.5% crude protein, 17.7% crude lipid)

^bGelatine; Beyza feed mill, Shiraz, Iran. (85% crude protein, crude lipid, 4.2)

^cWheat meal; Beyza feed mill, Shiraz, Iran. 12% crude protein, 3% crude lipid]

^dCorn starch, Beyza feed mill, Shiraz, Iran

^eParskilka Mazandaran, Iran (*Clupeonella* sp.)

^fMerck, Germany

^gVitamin premix (mg kg⁻¹) of premix: vitamin A, 5,000,000 IU; vitamin D3, 500,000 IU; vitamin E, 3000 mg; vitamin K3, 1500; vitamin B1, 6000; vitamin B2, 24,000; vitamin B5, 52,000; vitamin B6, 18,000; vitamin B12, 60,000; folic acid, 3000; nicotinamide 180,000; antioxidant, 500, Damloran pharmaceutical company, Broujerd, Iran

^hMineral premix (mg kg⁻¹) of premix: copper, 3000; zinc, 15,000; manganese, 20,000; Iron, 10,000; potassium iodate, 300. Microvit[®], Razak laboratories, Tehran, Iran

Crystalline amino acids: Merck, Germany, except isoleucine (Sigma-Aldrich, USA)

ⁱDispensable amino acid mixture (% mixture): L-alanine: 13; L-aspartic acid: 20; sodium glutamate: 32; L-glycine: 15; L-serine: 10; and L-proline: 10, Merck, Germany

^jNitrogen-free extract = 100 – (protein + lipid + ash)

^kGross energy content was estimated as: total carbohydrate × 17.2 J kg⁻¹; fat × 39.5 J kg⁻¹; and protein × 23.5 J kg⁻¹

Fish were adapted to the experimental conditions for 2 weeks and fed with the control diet during this period. 135 juveniles [initial body weight (BW_i) of 4.7 ± 0.1 g; mean ± standard error] were randomly distributed into 9 cylindrical polyethylene tanks (250 l of volume; 15 fish/tank; 3 replicates per diet). Fish were hand fed each of the experimental diets to visual satiation, three times per day (0800, 1200, and 1600 h), for 42 days.



Table 2 Amino acid profile of the control, essential amino acid deficient (EAAD), and protein-free (PF) diets ($n = 1$), g 100 g⁻¹ Diet

Amino acid composition	Control	EAAD	PF
Arginine	2.6	1.7	0.2
Lysine	2.2	1.3	0.2
Threonine	1.9	1.9	0.2
Histidine	1.2	0.7	0.1
Isoleucine	2.1	1.2	0.3
Leucine	2.4	1.5	0.3
Methionine	1.4	0.8	0.1
Cysteine*	0.3	0.3	0.1
Phenylalanine	1.8	1.1	0.3
Tyrosine	0.9	0.9	0.2
Tryptophan*	0.5	0.3	0.1
Valine	2.1	1.3	0.3
Alanine	3.6	4.4	0.2
Aspartate	5.5	6.7	0.3
Glutamate	7.8	9.8	0.3
Glycine	5.0	6.0	0.3
Proline	2.3	3.0	0.2
Serine	2.0	2.7	0.3

*These amounts were calculated based on the ingredient cysteine and tryptophan composition

Uneaten pellets were drained off an hour after feeding and counted every day to calculate the total feed intake (FI) per tank using a mean dry pellet weight. Tanks were supplied with filtered seawater (flow rate 1 l/min; salinity 48.0 ± 0.5 ‰; oxygen 7.6 ± 0.2 mg/l; pH 7.8 ± 0.4 ; temperature of 29.0 ± 1.5 °C) and subjected to natural photoperiod (30°32'N, 49°20'E; June and July).

Sample collection

At the end of the trial, fish were fasted for 24 h before being anaesthetized (2-phenoxyethanol at 0.5 ml/l; Merck, Schuchardt, Germany) and individually weighed (BW_f). For blood collection, three specimens from each replicate tank ($n = 9$ per diet, $n = 3$ per experimental replicate) were anesthetized and blood (ca. 250–500 μ l) was collected from the caudal vein with heparinized syringes, then centrifuged ($4000 \times g$, 10 min, 4 °C) and plasma separated. The vials containing plasma samples were transferred into liquid nitrogen and stored at -80 °C until further analysis. For the assessment of digestive enzyme activities, three fish per tank ($n = 9$ fish per diet treatment, $n = 3$ fish per replicate) were randomly sampled, euthanized with the same anaesthetic, and immediately eviscerated on ice surface. The alimentary tract was dissected, adherent adipose and connective tissues carefully removed and placed in individually marked plastic test tubes and stored at -80 °C until their analysis. For whole-body biochemical analyses, at the beginning of the trial, five fish from the initial stock were killed with an overdose of the anaesthetic and frozen at -80 °C for subsequent analysis. At the end of the feeding trial, three fish per experimental replicate were sacrificed with overdose of the same anaesthetic for subsequent biochemical analyses.

Fish growth, condition, and feed efficiency parameters

The BW_f , liver, viscera, and intraperitoneal fat weights were measured to the nearest 0.1 g and standard length (SL) to the nearest 1 mm. The following formulae were used to assess growth performance, feed utilization, and other parameters: body weight gain (WG, g) = ($BW_f - BW_i$); specific growth rate (SGR, %/day) = $[(\ln BW_f - \ln BW_i)/t] \times 100$, where t is experimental period (42 days); survival rate (SR, %) = (number of fish in each group remaining on day 42/initial number of fish) $\times 100$; feed intake (FI) = total feed intake (g)/number of fish; feed conversion ratio (FCR, %) = (feed intake (g)/weight gain (g)); protein efficiency ratio

(PER) = weight gain (g)/protein intake (g); condition factor (K) = (body weight (g)/(body length (cm))³) × 100; hepatosomatic index (HSI, %) = (liver weight (g)/whole-body weight (g)) × 100; viscerosomatic index (VSI, %) = (visceral weight (g)/whole-body weight (g)) and intraperitoneal fat index (IPF, %) = (IPF weight (g)/whole-body weight (g)) × 100. Daily N intake, gain and retention were estimated by following formulas:

$$\begin{aligned}\text{Mean metabolic weight gain (MBW)} &= 0.5 \times (\text{BW}_f^{0.75} + \text{BW}_i^{0.75}), \\ N \text{ intake (g N/kg MBW/day)} &= (D_i \times N_d) / (\text{MBW} \times t), \\ N \text{ gain (g N/kg MBW/day)} &= (\text{BW}_f \times N_f - \text{BW}_i \times N_i) / (\text{MBW} \times t), \\ N \text{ retention (\% } N \text{ intake)} &= (N \text{ gain} / N \text{ intake}) \times 100\end{aligned}$$

where MBW is the average metabolic body weight; D_i is the dry feed intake (g); N_d is the N content of the experimental diets; t is the duration of the feeding period (42 days), N_i and N_f are the initial and final body N content, and BW_i and BW_f are the mean initial and final body weight, respectively.

Biochemical analyses

Proximate analyses of ingredients, diets and whole body were determined using the standard methods (AOAC 2005). Amino acids (except for tryptophan and cysteine because of the susceptibility of these AA to acid hydrolysis) were determined after hydrolysis. Samples were hydrolysed in 6 N HCl for 24 h at 110 °C in glass vials filled with nitrogen. O-phthalaldehyde (OPA) was used as a pre-column derivatization reagent according to Lindroth and Mopper (1979). Total AA levels were determined by HPLC (Knauer, Germany), using C18 column (Knauer, Germany), at a flow rate of 1 ml/min with fluorescence detector (RF-530, Knauer, Germany). Peak identification and integration was performed by the software Waters Empower 2 (Milford, MA) using an AA standard H (Pierce, USA) as an external standard. Plasma biochemical parameters including glucose, total protein, albumin, and creatinine were analyzed by means of an auto-analyser (Mindray BS-200, China) using commercial clinical investigation kits (Pars Azmoon Kit, Tehran, Iran).

Determination of digestive enzymes

Samples were homogenized in 30 volumes (w/v) of ice-cold mannitol (50 mM), Tris–HCl buffer (2 mM) pH 7.0, at a maximum speed for 30 s (IKA, Ultra-turrax®, USA); then, 100 µl of 0.1 M CaCl₂ was added to the homogenate, which was centrifuged at 10,000× g for 10 min at 4 °C and the supernatants collected and stored in aliquots at – 80 °C for further enzyme quantification (Gisbert et al. 2009). Trypsin activity was measured with N - α -benzoyl-dlarginine-p-nitroanilide (BAPNA) as substrate. BAPNA (1 mM in 50 mM Tris–HCl, pH 8.2, 20 mM CaCl₂) was incubated with the enzyme extract at 37 °C and absorbance was recorded at 410 nm (Erlanger et al. 1961). One unit of trypsin activity corresponded to 1 µmol of 4-nitroaniline liberated in 1 min per ml of extracellular enzymatic extract, based on the extinction coefficient of the substrate ($\epsilon_{410} = 8800 \text{ M/cm}$). The activity of α -amylase was determined at 25 °C using 1% starch (Sigma-Aldrich) (diluted in a buffer at pH 6.9, 0.02 M Na₂HPO₄ and 0.006 M NaCl) as substrate by the 3, 5-dinitrosalicylic acid method (Worthington 1991) and absorbance was recorded at 540 nm. The α -amylase activity (U) was expressed as µg maltose liberated from starch per mg protein in sample per hour. Non-specific lipase activity was determined at 25 °C (absorbance: 405 nm) using 0.4 mM 4-nitrophenyl-myristate (Sigma-Aldrich) as substrate and absorbance were recorded at 405 nm (Gawlicka et al. 2000). One unit of lipase was defined as the amount of enzyme, which liberated one micromole of 4-nitrophenyl-myristate per minute. Carboxypeptidase A activity was measured using the protocol of Folk and Schirmer (1963) using HPA (hippuryl-L-phenylalanine) as substrate (dissolved in 25 mM Tris–HCl, 10 mM CaCl₂ buffer, pH 7.8). One unit of enzyme activity was defined as 1 M of hippuryl hydrolysed per minute using a coefficient of molar extinction of 0.36 at 254 nm. The protein concentration of the enzyme extracts was determined by the Bradford method (Bradford 1976) using bovine serum albumin as a protein standard. All enzymes were analyzed following the recommendations provided by Solovyev and Gisbert (2016).



Statistical analysis

Data were analyzed using SPSS version 15.0 (Chicago, IL, USA). All data are presented as mean \pm standard error of the mean. Equality of variances was tested with Levene's test and normality with Shapiro–Wilk's test. Arcsine transformations were conducted on all percentage data to achieve homogeneity of variance before statistical analysis. One-way analysis of variance was performed at a significance level of 0.05 following the confirmation of normality and homogeneity of variance. Tukey's test was used for multiple comparisons when statistical differences among groups were detected ($P < 0.05$).

Results and discussion

Fish growth, condition, and feed efficiency

In the present study, one interesting finding was the difference in the physiological responses between fish fed EAAD and PF diets. The information obtained from this study testing two extreme diets (EAAD and PF) may serve to better understand the impact of protein nutritional imbalances in fish performance, as well as generate reference values for a large series of physiological parameters under unsuitable nutritional conditions. Fish survival was significantly affected by experimental diets; thus, survival rates in fish fed the EAAD and PF diets were 8.9 and 25.6% lower, respectively, than in fish fed the control diet (Table 3). However, no external pathological signs were observed in fish, even in those fed the PF diet. Protein synthesis and accretion is a key component of the processes involved in growth response (Anthony et al. 2001). The lower survival rate in the PF and EAAD groups may have been related to disorders in different metabolic pathways, including (a) AA synthesis and catabolism, (b) generation of physiologically important low-molecular-weight peptides and nitrogenous substances, (c) intracellular protein turnover, (d) ammonia detoxification, (e) lipid and glucose metabolism, and (f) antioxidative reactions (Wu et al. 2014). In this study, weight gain in fish fed with EAAD was 5.6 g lower than fish fed with the control diet ($P < 0.05$). Furthermore, fish fed the PF diet lost weight and their final body weight was 12.2 g lower than the control group. Feed conversion ratio and PER values were negligible in fish fed with the PF diet, since those animals lost weight during the trial. All somatic indices, including the K, HSI, VSI, and IPF significantly decreased in fish fed the EAAD and PF diets ($P < 0.05$). In general, deficiency of most EAA in fish feeds leads to failure or loss of appetite, which results in a reduced FI and weight gain, as well as lower disease resistance (Wilson 2002). In addition, it has been reported that dietary deficiency in one or more EAAs results in the deamination of other AAs in the liver, which leads to an increased excretion of nitrogenous compounds and inefficient protein synthesis and somatic growth performance (Von der Decken and Lied 1993). In fact, imbalances in the dietary AA profile tend to lead to an increase in the oxidation of other EAAs and NEAAs present at normal levels in the feed, which results in reduced protein utilization (Rønnestad et al. 2007). In the current study, nitrogen retention extremely decreased in fish fed the EAAD and PF diets ($P < 0.05$). The N loss in fish fed the PF diet was 60 mg/kg MBW/day, indicating high whole-body protein breakdown in this group due to the lack of dietary AAs for protein synthesis, which impaired the turnover of tissue proteins. The highest plasma creatinine, which indicates high muscular AA catabolism for gluconeogenesis, in fish fed the PF diet also supported high whole-body AA breakdown in this group. However, fish fed the control and EAAD diets showed 240 and 60 mg/kg MBW/day N accretion, respectively. In this context, Marammazi et al. (2017) reported that deletion of a single EAA did not significantly affect N intake (g/kg MBW/day), whereas N gain (g/kg MBW/day) significantly decreased in all fish fed the EAA-deficient diets and the degree of N retention reduction was dependent upon the EAA deficiencies.

Body proximate composition and AA profile

Whole-body composition was significantly affected by different experimental diets (Table 4). As expected, fish fed the PF diet had the highest moisture, but the lowest protein, lipid, and gross energy contents ($P < 0.05$). The reduction in whole-body protein in the PF group was largely due to proteolysis, as an energy source to maintain lipid and carbohydrate utilization, and to synthesize the necessary enzymes and hormones



Table 3 Growth, feed utilization, and biometric parameters of *S. hasta* juvenile fed control, essential amino acid deficient (EAAD), and protein-free (PF) diets (mean \pm SE, $n = 3$)

	Diets		
	Control	EAAD	PF
Growth performance			
BW _i (g) ^a	4.6 \pm 0.1	4.6 \pm 0.0	4.6 \pm 0.0
BW _f (g) ^b	12.8 \pm 0.5 ^a	7.2 \pm 0.3 ^b	4.0 \pm 0.1 ^c
WG (g) ^c	8.2 \pm 0.6 ^a	2.6 \pm 0.1 ^b	-0.6 \pm 0.0 ^c
SGR (% body weight day ⁻¹)	2.4 \pm 0.1 ^a	1.0 \pm 0.1 ^b	-0.4 \pm 0.1 ^c
Survival (%) ^d	100.0 \pm 0.0 ^a	91.1 \pm 3.1 ^b	74.4 \pm 2.2 ^c
FI (g fish ⁻¹) ^e	11.4 \pm 0.2 ^a	8.4 \pm 0.2 ^b	6.9 \pm 0.1 ^c
FCR ^f	1.4 \pm 0.1 ^a	3.2 \pm 0.1 ^b	–
PER ^g	1.6 \pm 0.1 ^a	0.5 \pm 0.1 ^b	–
K ^h	2.4 \pm 0.2 ^a	2.2 \pm 0.1 ^b	1.9 \pm 0.2 ^c
HSI (%) ⁱ	2.0 \pm 0.0 ^a	1.1 \pm 0.1 ^b	1.1 \pm 0.4 ^b
VSI (%) ^j	9.5 \pm 0.4 ^a	5.8 \pm 0.3 ^b	6.4 \pm 0.5 ^b
IPF (%) ^k	1.8 \pm 0.2 ^a	0.2 \pm 0.1 ^b	0.1 \pm 0.0 ^b
<i>N</i> intake (g kg ⁻¹ MBW day ⁻¹)	0.74 \pm 0.0 ^a	0.72 \pm 0.03 ^a	0.03 \pm 0.0 ^b
<i>N</i> gain (g kg ⁻¹ MBW day ⁻¹)	0.24 \pm 0.01 ^a	0.06 \pm 0.0 ^b	-0.06 \pm 0.0 ^c
<i>N</i> retention (% <i>N</i> intake)	32.2 \pm 1.9 ^a	7.9 \pm 0.46 ^b	–

A different superscript in the same row denotes statistically significant differences ($P < 0.05$)

^a BW_i: initial body weight

^b BW_f: final body weight

^c SGR: specific growth rate = ((ln final weight – ln initial weight)/ t) \times 100, where t is experimental period = 42 days

^d Survival = (number of fish in each group remaining on day 60/initial number of fish) \times 100

^e FI: feed intake = feed intake/fish number

^f FCR: feed conversion ratio = weight gain (g)/feed intake (g)

^g PER: protein efficiency ratio = weight gain (g)/protein intake (g)

^h K: condition factor = (body weight (g)/(body length (cm))³) \times 100

ⁱ HSI: hepatosomatic index = (liver weight (g)/whole-body weight (g)) \times 100

^j VSI: viscerosomatic index = (viscera weight (g)/whole-body weight (g)) \times 100

^k IPF: intraperitoneal fat = (IPF weight (g)/whole-body weight (g)) \times 100

for intermediary metabolism, as also reported in Atlantic salmon (*Salmo salar*) fry fed a PF diet (Abboudi et al. 2006, 2009). In addition, during protein turnover, protein breakdown and resynthesis occurs, with oxidation of part of the amino acids from the protein breakdown. Thus, proteolysis must have occurred due to a lack of AAs from the diet. As intestinal metabolism may have a profound influence on whole-body AA composition, the differences in patterns of AAs between diets and body proteins have been reported to be

Table 4 Composition of whole-body proximate (% of wet weight) of *S. hasta* juvenile fed control, essential amino acid deficient (EAAD), and protein-free (PF) diets (mean \pm SE, $n = 3$)

	Diets			
	Initial	Control	EAAD	PF
Moisture	72.4 \pm 1.2	69.7 \pm 0.9 ^c	74.3 \pm 1.2 ^b	79.6 \pm 0.7 ^a
Crude protein	17.6 \pm 0.6	16.5 \pm 0.3 ^a	15.7 \pm 0.4 ^a	14.1 \pm 0.5 ^b
Crude lipid	5.1 \pm 0.3	7.6 \pm 0.2 ^a	3.3 \pm 0.4 ^b	2.5 \pm 0.5 ^c
Ash	4.3 \pm 0.2	4.7 \pm 0.2	5.1 \pm 0.3	5.4 \pm 0.1
Gross energy (MJ kg ⁻¹)	6.2 \pm 0.2	7.0 \pm 0.0 ^a	5.2 \pm 0.1 ^b	3.9 \pm 0.0 ^c

A different superscript in the same row denotes statistically significant differences ($P < 0.05$)



particularly large for various AAs (Wu et al. 2014). The whole-body AA composition of fish is shown in Table 5. Fish fed the PF diet had the highest and lowest whole-body arginine and threonine levels, respectively ($P < 0.05$). The lowest whole-body threonine in fish fed PF diet might be linked to the lowest survival rate in this group. In this regard, Yaghoubi et al. (2017) reported that *S. hasta* fed threonine-deficient diet had the lowest survival rate than fish fed other EAA-deficient diets. It has been reported that threonine is involved in many physiological and biochemical processes, including somatic growth, feed efficiency, digestive and absorptive processes, gene expression regulation, antioxidant and immune functions in different fish species (Gao et al. 2014, Habte-Tsion et al. 2015). In addition, whole-body aspartate, glutamate, and serine in fish fed the PF diet were lower than in other groups. Non-essential AAs can be synthesized de novo in adequate amounts by the animal organism from EAAs to meet the requirements for maintenance, growth, development and health (Wu et al. 2013). Thus, in the present study, dietary deficiencies in EAA may have led to the reduction of NEAA in fish fed the PF diet. Other whole-body AAs were not affected by dietary treatments ($P > 0.05$).

Plasma biochemical parameters

In the present study, plasma total protein and albumin were significantly reduced in the EAAD and PF groups in comparison with the control, suggesting a malfunctioning of the liver in these groups (Table 6), since all the plasma proteins with the exception of immunoglobulins are synthesized in this organ (Rosalki and McIntyre 1999). The results of this study showed that fish fed the EAAD and PF diets had the higher plasma glucose levels than the control group, results that might be attributed to a chronic stress condition in these groups (Barcellos et al. 1999, 2009). The highest plasma glucose in the PF group may have also been attributed to the high levels of carbohydrates in the PF diet as a consequence of limited ability of metabolizing glucose in most

Table 5 Amino acid profile of whole body (% of amino acids) of *S. hasta* juvenile fed control, essential amino acid deficient (EAAD), and protein-free (PF) diets (mean \pm SE, $n = 3$)

	Diets			
	Initial	Control	EAAD	PF
Essential amino acid				
Arginine	6.4 \pm 0.0	6.7 \pm 0.0 ^b	6.8 \pm 0.1 ^b	7.6 \pm 0.1 ^a
Histidine	0.7 \pm 0.0	0.6 \pm 0.0	0.6 \pm 0.0	0.7 \pm 0.0
Isoleucine	6.3 \pm 0.0	6.6 \pm 0.2	6.5 \pm 0.1	6.7 \pm 0.1
Leucine	7.5 \pm 0.0	7.3 \pm 0.2	7.6 \pm 0.1	7.8 \pm 0.4
Lysine	1.4 \pm 0.1	1.1 \pm 0.0	1.2 \pm 0.0	1.4 \pm 0.1
Methionine	2.8 \pm 0.0	2.9 \pm 0.1	2.8 \pm 0.0	3.1 \pm 0.1
Phenylalanine	2.9 \pm 0.0	3.0 \pm 0.0	2.9 \pm 0.0	3.0 \pm 0.1
Threonine	6.7 \pm 0.1	6.8 \pm 0.1 ^a	7.2 \pm 0.2 ^a	4.7 \pm 0.5 ^b
Valine	7.7 \pm 0.0	8.1 \pm 0.2	8.1 \pm 0.2	8.4 \pm 0.2
Non-essential amino acid composition				
Aspartate	10.9 \pm 0.4	10.2 \pm 0.0 ^a	10.2 \pm 0.3 ^a	9.3 \pm 0.4 ^b
Glutamate	18.1 \pm 0.43	17.5 \pm 0.2 ^a	17.2 \pm 0.1 ^a	16.0 \pm 0.2 ^b
Serine	6.9 \pm 0.1	7.4 \pm 0.2 ^a	7.3 \pm 0.1 ^a	6.5 \pm 0.2 ^b
Glycine	8.3 \pm 0.7	9.0 \pm 0.7	9.6 \pm 0.1	11.6 \pm 1.7
Taurine	2.1 \pm 0.1	1.7 \pm 0.1	1.6 \pm 0.0	1.7 \pm 0.1
Alanine	7.0 \pm 0.1	7.7 \pm 0.1	7.4 \pm 0.0	7.7 \pm 0.1
Tyrosine	2.0 \pm 0.0	2.0 \pm 0.0	2.0 \pm 0.0	2.1 \pm 0.1
Total EAA	42.3 \pm 0.0	43.1 \pm 0.6	41.7 \pm 0.5	43.4 \pm 0.4
Total NEAA	55.2 \pm 0.1	55.5 \pm 0.7	57.4 \pm 0.6	54.9 \pm 0.2
Total AA	97.5 \pm 0.0	98.6 \pm 0.1	99.1 \pm 0.1	98.3 \pm 0.5

A different superscript in the same row denotes statistically significant differences ($P < 0.05$)



carnivorous fish species (Booth et al. 2013). On the other hand, the increase in plasma glucose in fish fed the EAAD and PF diets was associated with an increase in plasma creatinine. Creatinine is a waste product from muscle turnover and its plasmatic levels provide an accurate marker of the hepatic and renal function. In one side, the increase in plasma creatinine level indicated a high muscular AA catabolism for gluconeogenesis, which might have resulted in higher plasma glucose levels in fish fed the PF and EAAD diets.

Digestive enzyme activities

Fish growth is mainly associated with its digestive ability that correlates with digestive enzyme activity (Rønnestad et al. 2007). The specific activity of selected pancreatic digestive enzymes in *S. hasta* juveniles fed different diets is presented in Table 7. Trypsin, non-specific lipase, α -amylase, and carboxypeptidase A activities were highest in fish fed the control diet, whereas the lowest activity levels were found in fish fed the PF diet. The activities of trypsin and carboxypeptidase A in fish fed the control diet were 10 and 30 times higher than in fish fed the EAAD and PF diets, respectively. Moreover, the activity of non-specific lipase in fish fed the EAAD and PF diets was ca. four times lower than fish fed the control diet. Regarding α -amylase activity, fish fed the control diet had 14 and 35 times higher α -amylase activity than fish fed the EAAD and PF diets, respectively ($P < 0.05$). Amino acids have a main role in digestive function through chemical sensing via the G protein-coupled receptors in the gastrointestinal tract and possibly in other tissues (i.e., liver and brain), gastrointestinal emptying, as potent stimulators of hormone (i.e., cholecystokinin), and pancreatic enzymes secretion, and they (glycine and taurine) conjugate with bile salts, which emulsify diet fats. In addition, AAs modulate growth and metabolism, and regulate intestinal microbiota (Wu et al. 2013, 2014). For instance, glutamic and aspartic acids are major metabolic fuels for intestine to maintain its digestive function and protect the mucosal integrity (Wu et al. 2013). The lowest digestive enzyme activities in fish fed the PF diet supported the above-mentioned results. Furthermore, enzyme secretion closely relates to the development of the pancreas and several studies reported that various EAA can affect the development and structural integrity of pancreas in fish (Chen et al. 2012, Tang et al. 2013, Li et al. 2015). The results of this study showed that drastic malfunction of exocrine pancreas may as a consequence of disorders in the above-mentioned mechanisms.

Conclusion

The results of this study showed that dietary protein and EAA deficiencies had negative effects on metabolic and physiological responses in *S. hasta* juveniles. The lack of dietary AAs for the protein synthesis led to lower survival and growth rates in fish fed the NEAA and PF diets. In addition, drastic changes in plasma metabolite parameters including protein, albumin, glucose, and creatinine indicated disorders in the liver and kidney function, as well in the metabolic pathways of fish fed NEAA and PF diets. Digestive enzyme secretion was lower in the NEAA and PF groups, indicating a main role of AAs in the synthesis and secretion of these enzymes from the exocrine pancreas in fish. The results of the current study may serve as a basis for establishing the optimal nitrogen content with proper AAs profile in the diet for *S. hasta* and other sparid species, to increase fish productivity while preventing N waste and environmental damage. However, more

Table 6 Plasma biochemical parameters of *S. hasta* juvenile fed control, essential amino acid deficient (EAAD), and protein-free (PF) diets (mean \pm SE, $n = 3$)

	Dietary treatments		
	Control	EAAD	PF
Total Protein (g l ⁻¹)	29.0 \pm 3.0 ^a	4.3 \pm 0.2 ^b	2.8 \pm 0.1 ^c
Albumin (g dl ⁻¹)	19.6 \pm 0.1 ^a	0.3 \pm 0.0 ^b	0.2 \pm 0.0 ^b
Glucose (mg dl ⁻¹)	134.0 \pm 7.0 ^c	145.2 \pm 8.0 ^b	231.6 \pm 14.0 ^a
Creatinine (mg dl ⁻¹)	2.1 \pm 0.1 ^c	10.8 \pm 0.6 ^b	23.4 \pm 1.1 ^a

A different superscript in the same row denotes statistically significant differences ($P < 0.05$)



Table 7 Pancreatic digestive enzyme activities (U mg protein⁻¹) of *S. hasta* juvenile fed control, essential amino acid deficient (EAAD), and protein-free (PF) diets (mean ± SE, n = 3)

	Diets		
	Control	EAAD	PF
Trypsin	3.1 ± 0.5 ^a	0.3 ± 0.2 ^b	0.1 ± 0.0 ^c
Lipase	2.2 ± 0.1 ^a	0.6 ± 0.1 ^b	0.5 ± 0.0 ^b
α-Amylase	1.4 ± 0.0 ^a	0.1 ± 0.0 ^b	0.04 ± 0.0 ^c
Carboxypeptidase A	3.0 ± 0.4 ^a	0.3 ± 0.0 ^b	0.1 ± 0.0 ^c

A different superscript in the same row denotes statistically significant differences ($P < 0.05$)

work is needed to evaluate the effects of dietary protein levels and AA profile on the utilization efficiency of limiting AAs, as well as to determine the influence of the duration of the experiments or fish size on their EAA requirements for maintenance.

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Compliance with ethical standards

Conflict of interest There is no conflict of interest between authors in the publication of this paper.

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