



A blend of plant proteins as a potential fishmeal substitute in the diet of Asian seabass *Lates calcarifer* (Bloch, 1790): Effect on growth, digestive enzymes and fatty acid composition

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ABSTRACT

A blend of plant proteins (soybean meal, groundnut oil cake, sunflower oil cake, wheat gluten meal and corn gluten meal at 4:2:1:6.5:6.5) was used to substitute fishmeal in the diet of Asian seabass *Lates calcarifer* (Bloch, 1790). Five iso-nitrogenous (401.13-407.33 g kg⁻¹) diets were prepared by substituting fishmeal at 0 (control), 25, 50, 75 and 100% (FM-R0, FM-R25, FM-R50, FM-R75 and FM-R100, respectively) using the above mix. Average final weight of the fish increased ($p < 0.05$) by three-folds (20.77-29.99 g) after 45 days compared to their initial weight (6.09±0.25 g) in all the treatments, however, significant ($p < 0.05$) variation was observed between the dietary treatments ($p \leq 0.002$). The results indicated that 50% of dietary fishmeal can be substituted and beyond this level resulted in significant reduction of growth performance. Broken-line analysis indicated optimal fishmeal substitution as 46.7%. The highest activity of pepsin, trypsin, chymotrypsin, carboxypeptidases-A and B was observed in fish fed with diets FM-R0, FM-R25 and FM-R50. Dietary change significantly ($p < 0.05$) influenced the carcass lipid composition of Asian seabass ($p \leq 0.001$). Fatty acids like C20:4, C20:5 and C22:6 were significantly ($p < 0.05$) low in fish carcass fed with higher levels of plant proteins (FM-R75 and FM-R100). The results concluded that dietary fishmeal level can be partially substituted using a blend of plant proteins in the diet of Asian seabass juveniles.

Keywords: Asian seabass, Digestive enzymes, Fatty acid, Fishmeal, Growth, Plant proteins

Introduction

Asian seabass *Lates calcarifer* (Bloch, 1790) is an economically important food fish and is generally cultured in tropical and subtropical regions of the Pacific and Indian Ocean and South-East Asian countries. In general, most marine carnivorous fish, including seabass require high dietary protein compared to omnivorous and herbivorous species. Therefore, fishmeal is being used as a major protein source in commercial formulations due to its balanced nutrients, in particular essential amino acids and fatty acids, higher palatability and digestibility (Jannathulla *et al.*, 2019a). However, the global fishmeal production has reduced from 6.2 million t to 4.3 million t, in the past two decades due to various climatic events, which resulted in the escalation of its price from 452 to 2169 USD per ton (Jannathulla *et al.*, 2019b; USDA, 2020). This scenario prompted researchers towards substituting a considerable quantity of fishmeal by using a potential alternative from various animal and plant origins. Though animal proteins have been extensively evaluated as a protein source in fish feeds, their utility has not been explored commercially due to the higher content of lipid

and ash, which negatively influences the palatability and digestibility (Nandakumar *et al.*, 2013). Besides, European Community stated that the usage of rendered animal protein sources, in particular meat and bone meal have been restricted in various countries due to the spread of Bovine Spongiform Encephalitis (Title 21 Part 589.2000 of the US Food and Drug Administration, Code of Federal Regulations). Therefore, researchers prefer plant protein sources as an alternative to fishmeal due to wide availability, reasonable price and to a certain extent desirable nutrient content (Jannathulla *et al.*, 2018a).

Plant protein constituents and their byproducts such as soybean meal (Tomas *et al.*, 2009), groundnut meal (Sanchez-Lozano *et al.*, 2011) and other oil seed meals/cakes (Martínez-Llorens *et al.*, 2012) have been successfully substituted by dietary fishmeal from 12 to 40% in different aquatic species. Similarly, glutes, in particular wheat gluten meal and corn gluten meal have shown to be potential protein sources, containing 60 to 80% of crude protein, which is reported to be a vital alternative to fishmeal (Nandakumar *et al.*, 2017). Major problems associated with plant proteins are deficiencies

of certain essential amino acids (methionine, lysine and tryptophan) and higher content of anti-nutrients (Jannathulla *et al.*, 2017a). In order to resolve these issues, researchers used a combination of plant protein sources with varying levels of success (Burr *et al.*, 2012; Dayal *et al.*, 2020). Fish are capable of modulating their digestive enzyme pattern in response to the quality and concentration of dietary nutrients and their sources (Gonzalez-Felix *et al.*, 2010). These responses are the basis of the adaptation of fish to dietary changes, and knowing the mechanisms involved in this process can provide useful information to improve fish growth performance and health status (Eusebio and Coloso, 2002). However, the effect of utilisation of various plant protein sources together as a fishmeal substitute in *L. calcarifer* is scarce in the literature. Hence, present investigation is aimed to study the effect of inclusion of graded levels of various plant proteins on growth, digestive enzyme and fatty acid composition in *L. calcarifer* in relation to fishmeal substitution.

Materials and methods

Experimental diets

Five different plant protein sources such as soybean meal, groundnut oil cake, sunflower oil cake, wheat gluten meal and corn gluten meal, were purchased from the local markets in Chennai, India and were mixed in the ratio of 4:2:1:6.5:6.5 based on our earlier results (Madhubabu, 2019) to substitute fishmeal in the diet of Asian seabass.

Five iso-nitrogenous (401.13-407.33 g kg⁻¹) and iso-lipidic (150.30-155.15 g kg⁻¹) diets were formulated using locally available ingredients listed in Table 1. The experimental diets were prepared by replacing fishmeal (w/w) at the rate 0 (control), 25, 50, 75 and 100% (FM-R0, FM-R25, FM-R50, FM-R75 and FM-R100, respectively). In preparing experimental diets, the coarse ingredients were powdered in a micro pulveriser and passed through a 0.5 mm sieve. The ground materials were thoroughly mixed by hand and to which, additives (vitamin, mineral mix and binder) followed by oil sources (fish oil and soy lecithin) were subsequently added. They were homogenised in an electric blender for 20 min. The homogenised mash was hydrated with deionised water (500 ml kg⁻¹ of mash) and kneaded into a dough. It was steamed for 5 min at atmospheric pressure and pelleted in a table top pelletiser with a 2 mm diameter die. The pellets were dried in a forced-air oven at 60°C to bring down the moisture content to <10% and stored in a refrigerator until use. Proximate, essential amino acids and fatty acid composition of experimental diets are presented in Table 2.

Experimental feeding

Juveniles of Asian seabass procured from the hatchery of ICAR-Central Institute of Brackishwater Aquaculture (ICAR-CIBA), Chennai, India were used in the present study. They were acclimatised to the experimental conditions for two weeks by feeding a formulated feed

Table 1. Ingredient and proximate composition of experimental diets containing graded levels of plant protein sources by replacing fishmeal (g kg⁻¹ as fed basis)

| Particulars | Experimental diets | | | | |
|---------------------------------|--------------------|--------|--------|--------|---------|
| | FM-R0 | FM-R25 | FM-R50 | FM-R75 | FM-R100 |
| Ingredient composition | | | | | |
| Fishmeal ¹ | 350.0 | 262.5 | 175.0 | 87.5 | - |
| Plant protein mix ² | - | 100.0 | 200.0 | 300.0 | 400.0 |
| Mantis shrimp meal ³ | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 |
| Soybean meal | 200.0 | 200.0 | 200.0 | 200.0 | 200.0 |
| Groundnut oil cake | 40.0 | 40.0 | 40.0 | 40.0 | 40.0 |
| Sunflower oil cake | 15.0 | 15.0 | 15.0 | 15.0 | 15.0 |
| Whole wheat | 121.0 | 102.5 | 82.0 | 63.5 | 45.0 |
| Broken rice | 30.0 | 30.0 | 30.0 | 30.0 | 30.0 |
| Fish oil ¹ | 74.0 | 78.0 | 82.0 | 86.0 | 90.0 |
| Soy-lecithin ⁴ | 40.0 | 42.0 | 46.0 | 48.0 | 50.0 |
| Pre-mix ⁵ | 20.0 | 20.0 | 20.0 | 20.0 | 20.0 |
| Binder ⁶ | 10.0 | 10.0 | 10.0 | 10.0 | 10.0 |

¹Bismi Fisheries, Mayiladuthurai, Tamil Nadu, India

²Plant protein mix contains soybean meal, groundnut oil cake, sunflower oil cake, wheat gluten meal and corn gluten meal at the ratio of 4:2:1:6.5:6.5

³Ismail haji Abdullah and Co., Chennai, Tamil Nadu, India

⁴Real Soy Enterprises, Madhya Pradesh, India

⁵Pre-mix (per 100 g): Vitamin A-200,000 IU, Vitamin D-40,000 IU, Vitamin E-30 U, Vitamin K-40 mg, Riboflavin-80 mg, Capantothenate-100 mg, Nicotinamide-400 mg, Vitamin B12-0.24 mg, Choline chloride-6 g, Ca-30 g, Mn-1.1 g, I-40 mg, Fe-300 mg, Zn-0.6 g, Cu-80 mg, Co-18 mg (Sarabhai Zydus Animal Health Ltd., Vadodara, Gujarat, India)

⁶Pegabind, Bentoli Agri-Nutrition Asia Pvt. Ltd., Singapore

Other feed ingredients were purchased from local markets around Chennai, Tamil Nadu, India

Table 2. Proximate, essential amino acid and fatty acid composition of experimental diets containing graded levels of plant protein sources by replacing fishmeal

| Particulars | Experimental diets | | | | |
|--|--------------------|--------|--------|--------|---------|
| | FM-R0 | FM-R25 | FM-R50 | FM-R75 | FM-R100 |
| Proximate composition (% as fed basis) | | | | | |
| Moisture | 92.12 | 91.30 | 91.50 | 91.70 | 92.40 |
| Crude protein | 401.13 | 407.33 | 405.20 | 402.68 | 406.31 |
| Ether extract | 155.15 | 150.30 | 150.75 | 151.97 | 151.48 |
| Crude fibre | 30.72 | 36.64 | 38.53 | 39.64 | 41.09 |
| NFE ¹ | 185.79 | 197.90 | 207.50 | 218.80 | 224.54 |
| Total ash | 135.09 | 116.53 | 106.52 | 95.21 | 84.18 |
| Essential amino acids (g per 16 g N) | | | | | |
| Arg | 2.86 | 2.58 | 2.34 | 2.24 | 1.93 |
| His | 1.31 | 1.10 | 1.10 | 1.06 | 1.07 |
| Ile | 1.98 | 1.88 | 1.85 | 1.84 | 1.79 |
| Leu | 2.89 | 3.30 | 3.73 | 4.01 | 4.28 |
| Lys | 2.54 | 2.27 | 1.98 | 1.68 | 1.37 |
| Met | 1.03 | 0.96 | 0.94 | 0.87 | 0.84 |
| Phe | 1.81 | 1.81 | 1.86 | 1.96 | 2.23 |
| Thr | 1.56 | 1.62 | 1.50 | 1.49 | 1.40 |
| Trp | 0.51 | 0.45 | 0.42 | 0.40 | 0.38 |
| Val | 1.94 | 1.87 | 1.76 | 1.74 | 1.68 |
| Fatty acids (% total fatty acids) | | | | | |
| C14:0 | 6.43 | 6.42 | 6.31 | 6.31 | 6.44 |
| C15:0 | 0.73 | 0.70 | 0.66 | 0.62 | 0.59 |
| C16:0 | 19.66 | 19.29 | 18.11 | 16.47 | 15.03 |
| C17:0 | 0.87 | 0.71 | 0.69 | 0.71 | 0.67 |
| C18:0 | 5.69 | 5.53 | 5.30 | 5.22 | 5.09 |
| C20:0 | 0.48 | 0.52 | 0.49 | 0.53 | 0.50 |
| C24:0 | 0.23 | 0.20 | 0.23 | 0.22 | 0.19 |
| C16:1 | 4.78 | 4.63 | 4.57 | 4.44 | 4.35 |
| C18:1n9 | 14.59 | 14.63 | 14.86 | 15.85 | 16.63 |
| C18:1n7 | 3.38 | 3.07 | 2.95 | 2.82 | 2.55 |
| C20:1 | 0.65 | 0.58 | 0.49 | 0.52 | 0.59 |
| C18:2n6 | 16.47 | 15.42 | 15.61 | 16.07 | 16.55 |
| γC18:3n6 | 0.69 | 0.71 | 0.72 | 0.69 | 0.63 |
| αC18:3n3 | 2.11 | 2.16 | 2.24 | 2.29 | 2.36 |
| C20:2n6 | 0.21 | 0.20 | 0.20 | 0.21 | 0.22 |
| C20:3n6 | 0.32 | 0.38 | 0.43 | 0.46 | 0.53 |
| C20:4n6 | 1.76 | 1.82 | 1.84 | 1.85 | 1.88 |
| C20:5n3 | 9.42 | 9.04 | 8.61 | 8.37 | 8.03 |
| C22:6n3 | 6.48 | 6.36 | 6.25 | 6.13 | 6.05 |

All the values are the mean of three replicates.

¹Nitrogen free extract (Calculated by difference)

(400 g kg⁻¹ crude protein and 150 g kg⁻¹ ether extract) developed at ICAR-CIBA, Chennai, India. Post-acclimatisation, a total of 300 healthy and uniform sized fish (6.09±0.25 g) were randomly stocked in fifteen 500 l (1.31 x 0.64 x 0.73 m) oval-shaped fiberglass reinforced plastic (FRP) tanks. All the experimental tanks were provided with sand-filtered seawater and continuous aeration by air diffuser stones, were closed with fibremat to prevent escape of animals and to avoid external disturbances. Each treatment had three replicates and

each replicate contained twenty fish. Individual fish length and weight were recorded at the beginning of the experiment and feeding trial was conducted for 45 days. Fish were hand fed to satiation for an hour, twice daily (09:30 and 16:00 hrs). After feeding, uneaten feed pellets (if any) were siphoned out from the tanks daily, using a clean Falcon tube and dried at 60°C in a hot air oven to assess the feed intake. Animals were group weighed at fortnightly intervals for adjusting the feed ration. During the experimental period, about 80%

of water was exchanged before the first feeding, every day and water quality parameters such as temperature ($28.47 \pm 1.58^\circ\text{C}$), salinity ($26.14 \pm 1.57\text{‰}$), pH (8.24 ± 0.63), dissolved oxygen ($8.01 \pm 0.71 \text{ mg l}^{-1}$) and ammonia-N ($0.08 \pm 0.01 \text{ mg l}^{-1}$) were maintained at optimal levels. At the end of the experiment, the fish were anaesthetised using 2-phenoxyethanol at a dose of 0.3 ml l^{-1} (Nandakumar *et al.*, 2017) and weight was measured individually to determine the growth parameters *viz.* weight gain (WG), average daily growth (ADG), specific growth rate (SGR), daily growth coefficient (DGC), feed conversion ratio (FCR), protein efficiency ratio (PER), apparent protein utilisation (APU) and survival as follows:

$$\text{WG (\%)} = \frac{\text{Final body weight (g)} - \text{Initial body weight (g)}}{\text{Initial body weight (g)}} \times 100$$

$$\text{ADG (mg day}^{-1}\text{)} = \frac{\text{Final body weight (g)} - \text{Initial body weight (g)}}{\text{Days of experiment}}$$

$$\text{SGR} = \frac{[\ln \text{Final body weight}] - [\ln \text{Initial body weight}]}{\text{Days of experiment} \times 100}$$

$$\text{DGC} = \frac{[\text{Final body weight}^{1/3}] - [\text{Initial body weight}^{1/3}]}{\text{Days of experiment} \times 100}$$

$$\text{FCR} = \frac{\text{Feed intake (g)}}{\text{Weight gain (g)}}$$

$$\text{PER} = \frac{\text{Weight gain (g)}}{\text{Protein intake (g)}}$$

$$\text{APU} = \frac{\text{Protein gain (g)}}{\text{Protein intake (g)}}$$

$$\text{Survival (\%)} = \frac{\text{Final number of animals}}{\text{Initial number of animals}} \times 100$$

Fish from each replicate treatment (total 15 fish per treatment) were dissected to separate the digestive tract such as stomach, pyloric caeca, liver, anterior and posterior intestine to assess the activity of digestive enzymes. In addition, another fifteen fish from each treatment (five fish per replicate) that were not used for the measurement of digestive enzyme activity, were analysed for carcass proximate and fatty acid composition.

Biochemical analysis

Proximate composition was determined by AOAC (1997) method. Amino acid profiles were analysed using a pre-column derivatisation HPLC gradient system (Shimadzu Corp, LC-30AD) after hydrolysing the samples with 6 N hydrochloric acid in a sealed tube filled with nitrogen for 22 h at 110°C in an oven (Finlayson, 1964; Jannathulla *et al.*, 2017b). Tryptophan, being liable to acid hydrolysis, was measured after alkali hydrolysis by spectrophotometric method at 500 nm (Sastry and Tammuru, 1985). Fatty acid methyl esters (FAMES) were prepared according to Metcalfe *et al.* (1966) and was quantified by a gas chromatograph (GC-2014 Shimadzu) (Khan *et al.*, 2018). The total soluble protein was determined as per Lowry *et al.* (1951) in a UV-VIS spectrophotometer (UV-1800, Shimadzu, Japan). All the digestive

enzymes, including pepsin, trypsin, chymotrypsin, amylase, carboxypeptidase-A, carboxypeptidase-B, acid phosphatase, alkaline phosphatase and lipase were measured as per the methods described by Anson (1938), Erlanger *et al.* (1961), Hummel (1959), Bernfeld (1955), Folk and Schirmer (1963), Folk *et al.* (1960), Walter and Schutt (1974) and Winkler and Stuckmann (1979) respectively and their activity was expressed as U mg protein⁻¹.

Statistical analysis

Experimental data on growth and body composition were statistically analysed by one-way analysis of variance (ANOVA). Two-way ANOVA was performed using two different factors *viz.* level of fishmeal substitution (0, 25, 50, 75 and 100%) and different parts of the digestive tract (stomach, pyloric caeca, liver, anterior and posterior intestine) to find difference in digestive enzyme activity. The statistical package for social science (SPSS 17.0) was used to analyse data obtained in the present investigation and means were compared using the Tukey's test at significance level of $p < 0.05$. Prior to statistical evaluation, data were checked for determining the homogeneity of variance after ascertaining the normal distribution. Broken-line regression was carried out to find an optimal substitution level of fishmeal using a combination of plant proteins in the diet of Asian seabass.

Results and discussion

Several studies conducted earlier to explore the utilisation of various plant proteins such as soybean meal (Tantikitti *et al.*, 2005), lupin concentrate, wheat gluten meal (Glencross *et al.*, 2011), canola meal (Plaipetch and Yakupitiyage, 2012) and corn gluten meal (Nandakumar *et al.*, 2017) in Asian seabass showed <50% fishmeal substitution. Jannathulla *et al.* (2018b) suggested that this could partly be attributed to the deficiency of limiting amino acids, in particular methionine and lysine.

Boonyaratpalin *et al.* (1998) reported that the solvent extracted soybean meal could replace 10 to 15% dietary fishmeal without having any deleterious effect in Asian seabass and also suggested that the limitation in soybean meal usage is mainly attributed to the presence of anti-nutritional factors, especially trypsin inhibitor. Though the meals of gluten from wheat and corn were identified as a potential protein source (60 to 80% protein), beyond the inclusion of 10% by replacing an equal quantity of fishmeal, significantly ($p < 0.05$) affected digestibility and growth of Asian seabass (Nandakumar *et al.*, 2017). Hence, a combination of plant protein sources with soybean meal, wheat gluten meal, corn gluten meal, groundnut oil cake and sunflower oil cake was used to substitute fishmeal in our study. The results showed that

fish fed with diet containing up to 50% plant proteins by substituting fishmeal (w/w) had no negative effect on growth rate in terms of WG, ADG, SGR and DGC (Table 3). Our results corroborated the findings of De Francesco *et al.* (2007), who reported that plant proteins could replace >50% of fishmeal in the diet of gilthead seabream. This improved performance with a combination of plant proteins were mainly attributed to the compensation of deficient amino acids. It is noteworthy that methionine and lysine are the most limiting amino acids in the selected plant protein sources and the same was also available in fishmeal. Though level of methionine and lysine were reduced from 1.18 to 0.91 g per 16 g N and 2.75 to 1.47 g per 16 g N, respectively in the experimental diets with increasing the inclusion of plant proteins, the growth performance of fish fed diet with plant protein sources up to 50% was on par with control group (FM-R0). This clearly indicates that a combination of plant proteins would be better in compensating deficient amino acids rather than individual source. However, broken-line analysis (Fig. 1) indicated that optimal fishmeal substitution using a combination of plant protein was 46.7% in the diet of Asian seabass.

On the other hand, when the replacement of fishmeal was further increased (75 to 100%), growth indices of experimental fish exhibited significant ($p < 0.05$) decrease which could be attributed to the higher content of anti-nutritional factors (Jannathulla *et al.*, 2017a). The SGR obtained in the present study was in the range of 2.74-3.48 whereas, comparatively lower SGR was reported by Kissil and Lupatsh (2004) in gilthead sea bream (1.28), by Tomas *et al.* (2011) in Dentex (0.69) and by Guroy *et al.* (2013) in catfish when fish fed a diet with 100% plant proteins as fishmeal substitute. However, the results obtained in the present study are similar to the earlier reports in Asian seabass fed corn gluten meal as substitute

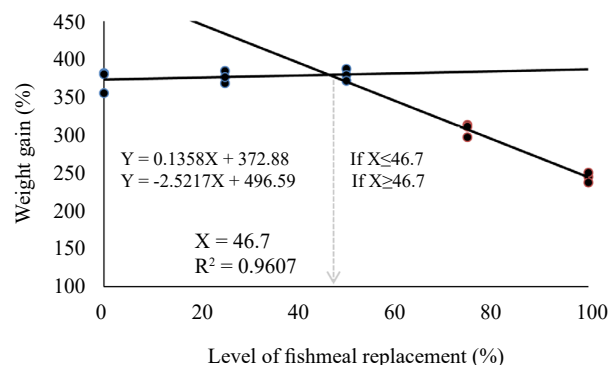


Fig. 1. Estimation of optimal replacement level of fishmeal with plant proteins in the diet of *L. calcarifer* using broken-line analysis

for fishmeal at moderate level (Nandakumar *et al.*, 2017). The difference noticed between the studies could be due to the variation in species, age, size and culture conditions. But in contrast, Kissil and Lupatsh (2004) and Tomas *et al.* (2011) reported a higher SGR in fish fed 100% fishmeal substituted diet using plant proteins than those fed control diet (no replacement). However, the present findings are in concurrence with rainbow trout (Gomes *et al.*, 1995), European seabass (Kaushik *et al.*, 2004), Turbot (Fournier *et al.*, 2004) and Atlantic salmon (Espe *et al.*, 2006) fed plant ingredients as a major protein source in their diets. The FCR gradually increased ($p < 0.05$) from 1.82 to 2.26 with increasing fishmeal substitution. The result is in agreement with the finding of Nandakumar *et al.* (2017). On the contrary, the FCR was not significantly ($p < 0.05$) affected due to the inclusion of plant proteins in a previous study with European seabass (Kaushik *et al.*, 2004). Survival of the fish fed with diets having 50% plant proteins did not differ significantly (88.33 to 96.66%) whereas, it was reduced ($p < 0.05$) to 70% with FM-R75

Table 3. Growth performances of *L. calcarifer* fed with experimental diets having graded levels of plant protein sources by replacing fishmeal (n=3)

| Particulars | Experimental diets | | | | | SEM (\pm) | CV (%) | p value |
|--|---------------------|---------------------|---------------------|---------------------|---------------------|---------------|--------|---------|
| | FM-R0 | FM-R25 | FM-R50 | FM-R75 | FM-R100 | | | |
| Initial weight (g) | 6.35 ^a | 6.01 ^a | 5.95 ^a | 6.11 ^a | 6.04 ^a | 0.150 | 8.357 | 0.887 |
| Final weight (g) | 29.99 ^a | 28.62 ^a | 28.52 ^a | 24.88 ^b | 20.77 ^c | 1.989 | 6.988 | 0.002 |
| Weight gain (%) | 372.65 ^a | 376.72 ^a | 379.44 ^a | 307.46 ^b | 244.42 ^c | 62.172 | 3.087 | <0.001 |
| ADG ¹ (mg day ⁻¹) | 525.40 ^a | 502.59 ^a | 501.63 ^a | 417.25 ^b | 327.55 ^c | 544.971 | 6.754 | <0.001 |
| SGR ² | 3.45 ^a | 3.47 ^a | 3.48 ^a | 3.12 ^b | 2.74 ^c | 0.002 | 1.595 | <0.001 |
| DGC ³ | 2.78 ^a | 2.75 ^a | 2.75 ^a | 2.42 ^b | 2.06 ^c | 0.002 | 2.540 | <0.001 |
| FCR ⁴ | 1.82 ^c | 1.86 ^c | 1.84 ^c | 2.07 ^b | 2.26 ^a | 0.002 | 2.596 | <0.001 |
| PER ⁵ | 1.37 ^a | 1.34 ^a | 1.35 ^a | 1.20 ^b | 1.10 ^c | 0.001 | 2.064 | <0.001 |
| APU ⁶ | 26.40 ^{ab} | 25.42 ^b | 26.44 ^a | 23.49 ^c | 21.59 ^d | 0.164 | 2.158 | <0.001 |
| Survival (%) | 96.66 ^a | 91.66 ^a | 88.33 ^a | 70.00 ^b | 53.33 ^c | 32.957 | 9.444 | 0.001 |

Means bearing different superscript letters within in the row significantly ($p < 0.05$) differ each other

¹Average daily growth; ²Specific growth rate; ³Daily growth coefficient; ⁴Feed conversion ratio; ⁵Protein efficiency ratio; ⁶Apparent protein utilisation

and to 53.33% with FM-R100. This is attributed to the poor palatability and intake due to higher inclusion level of plant proteins.

Utilisation of feed by the fish mainly depends on nutrient levels, ingredients in the feed, feeding time, amount of feed and activity of the digestive enzymes in the fish. Of all the dietary treatments, fish fed with diets FM-R0, FM-R-25 and FM-R50 showed significantly ($p < 0.05$) higher pepsin activity (14.76 - 14.99 U mg protein⁻¹) compared to other groups. Trypsin, amylase and chymotrypsin were found in all the analysed digestive parts except stomach and was higher in the anterior intestine (6.62 U mg protein⁻¹), posterior intestine (4.33 U mg protein⁻¹) and pyloric caeca (2.36 U mg protein⁻¹), respectively. The activity of trypsin and chymotrypsin decreased with increasing fishmeal substitution and the reverse was true for amylase (Table 4). Krogdahl *et al.* (2003) studied the effects on digestive enzymes with the incorporation of graded levels of soybean meal in Atlantic salmon and showed a lower digestive enzymatic activity with increase of dietary soybean level. Lin and Luo (2011) studied the replacement of fishmeal with soybean meal on growth, digestive enzymes and transaminase activities in juvenile tilapia. Significant decrease in protease activities were observed in both intestine and hepatopancreas, but not in the amylase activity in hepatopancreas among the dietary treatments. All the digestive parts had an activity of carboxypeptidase-A, carboxypeptidase-B and

acid phosphatase except liver. Posterior intestine had the highest activity of both carboxypeptidase-A and B (5.43 and 8.51 U mg protein⁻¹), while acid phosphatase was high in pyloric caeca (0.87 U mg protein⁻¹). There was no significant difference in carboxypeptidase-A up to 75% fishmeal substitution and was limited to 50% for carboxypeptidase-B and acid phosphatase. The highest activity of alkaline phosphatase was found in the anterior intestine (5.16 U mg protein⁻¹) followed by pyloric caeca and posterior intestine, while it was not detected in both stomach and liver. However, the dietary change had no influence ($p = 0.046$) on alkaline phosphatase activity irrespective of the digestive tract (4.48 - 4.75 U mg protein⁻¹). Activity of lipase was detected in all the digestive parts and was high in the pyloric caeca (1.32 U mg protein⁻¹). Fish fed with FM-R25 diet had the highest activity of lipase compared to other dietary groups (0.97 - 1.09 U mg protein⁻¹). Jalili *et al.* (2012) studied the effects of replacement of fishmeal with graded levels of plant protein sources on feed utilisation and digestive enzyme activities in carnivorous feeding habit of rainbow trout. Complete replacement of fishmeal resulted in significant reduction of alkaline protease and lipase activities, but not amylase activity in rainbow trout. Yu *et al.* (2013) studied the effects of replacement of fishmeal with soybean meal on growth, body composition and digestive enzyme activities in the intestine and hepatopancreas of Chinese sucker. Digestive enzyme activities showed decreasing trend with increasing dietary

Table 4. Digestive enzymes activities (U mg protein⁻¹) in various digestive tissues of *L. calcarifer* fed with experimental diets having graded levels of plant protein sources by replacing fishmeal (n=3)

| Particulars | Digestive enzymes | | | | | | | | |
|------------------------|--------------------|-------------------|-------------------|--------------------|---------------------|---------------------|---------------------|----------------------|--------------------|
| | Pepsin | Trypsin | Chymotrypsin | Amylase | Carboxy-peptidase A | Carboxy-peptidase B | Acid phosphatase | Alkaline phosphatase | Lipase |
| Experimental diets (A) | | | | | | | | | |
| FM-R0 | 14.99 ^a | 4.34 ^a | 2.12 ^a | 3.47 ^b | 3.32 ^a | 5.56 ^a | 0.62 ^a | 4.63 ^a | 1.08 ^{ab} |
| FM-R25 | 14.81 ^a | 4.41 ^a | 2.16 ^a | 3.86 ^a | 3.35 ^a | 5.60 ^a | 0.59 ^{abc} | 4.66 ^a | 1.12 ^a |
| FM-R50 | 14.76 ^a | 4.55 ^a | 2.08 ^a | 4.01 ^a | 3.32 ^a | 5.55 ^a | 0.60 ^{ab} | 4.75 ^a | 1.09 ^{ab} |
| FM-R75 | 9.75 ^b | 3.55 ^b | 2.01 ^a | 3.94 ^a | 3.29 ^a | 5.17 ^b | 0.55 ^{bc} | 4.63 ^a | 1.00 ^{bc} |
| FM-R100 | 8.20 ^c | 2.97 ^c | 1.50 ^b | 3.70 ^{ab} | 2.93 ^b | 4.65 ^c | 0.52 ^c | 4.48 ^a | 0.97 ^c |
| Digestive parts (B) | | | | | | | | | |
| Stomach | 20.55 ^a | - | - | - | 1.27 ^d | 1.69 ^d | 0.40 ^d | - | 0.59 ^c |
| Pyloric caeca | 4.45 ^b | 2.53 ^c | 2.36 ^a | 3.89 ^b | 2.58 ^c | 3.96 ^c | 0.87 ^a | 4.87 ^b | 1.32 ^a |
| Liver | - | 2.19 ^d | 1.98 ^b | 2.65 ^c | - | - | - | - | 1.03 ^b |
| Anterior intestine | - | 6.65 ^a | 1.94 ^b | 4.32 ^a | 3.70 ^b | 7.06 ^b | 0.55 ^b | 5.16 ^a | 1.06 ^b |
| Posterior intestine | - | 4.48 ^b | 1.61 ^c | 4.33 ^a | 5.43 ^a | 8.51 ^a | 0.47 ^c | 3.87 ^c | 1.25 ^a |
| p value | | | | | | | | | |
| A | <0.001 | <0.001 | <0.001 | 0.022 | <0.001 | <0.001 | 0.046 | 0.051 | 0.022 |
| B | <0.001 | <0.001 | <0.001 | <0.001 | <0.001 | <0.001 | <0.001 | <0.001 | <0.001 |
| A x B | <0.001 | 0.113 | <0.001 | 0.006 | 0.487 | <0.001 | 0.077 | 0.024 | 0.075 |
| SEM (±) | 0.586 | 0.108 | 0.024 | 0.097 | 0.017 | 0.056 | 0.004 | 0.018 | 0.011 |
| CV (%) | 8.059 | 10.873 | 10.366 | 10.755 | 5.343 | 5.875 | 14.520 | 3.789 | 12.866 |

Mean bearing different superscript letters in a column within the main effects between the categories significantly ($p > 0.05$) differ

soybean meal. When fishmeal was replaced beyond 60%, the protease and lipase activities were reduced significantly in both intestine and hepatopancreas.

The carcass proximate composition of Asian seabass fed with experimental diets containing varying levels of plant proteins showed significant ($p < 0.05$) difference in lipid and was high in FM-R100 fed group. This observation is similar to the result obtained in other fish species fed diets having plant protein blend (Cabral *et al.*, 2011). However, other proximate principles were not affected due to the dietary change in our study (Table 5). A similar trend was observed by Kaushik *et al.* (2004) in European seabass fed diet with almost complete replacement of fishmeal. In the present study, the dietary change has a remarkable impact on carcass fatty acids and among them, C16:0, C18:0, C16:1, C18:1n-9, C18:2n-6

and C22:6 were particularly abundant and contributes more than 50% of the total fatty acids. This agrees with the observation on other fish like Atlantic salmon (Pratoomyot *et al.*, 2010), Senegalese sole (Cabral *et al.*, 2011) and European seabass (Messina *et al.*, 2013). Fatty acids like C16:1, C20:4, C20:5 and C22:6 were found significantly ($p < 0.05$) decreased with increasing fishmeal substitution, while the reverse trend was observed for C18:1n-9, C18:2n-6. In most cases, fatty acid composition of experimental diets was reflected in the body composition. Similar findings were observed in the present study, however, certain fatty acids appeared to be retained or actively synthesised, though their levels were lower in the diet. Gonzalez-Felix *et al.* (2002) suggested that this could be due to the sparing and retention phenomena of the cultured species. On the other hand, the level of certain

Table 5. Carcass proximate and fatty acid composition of *L. calcarifer* fed with experimental diets having graded levels of plant protein sources by replacing fishmeal (n=3)

| Particulars | Experimental diets | | | | | SEM (\pm) | CV (%) | p value |
|---|---------------------|---------------------|---------------------|---------------------|---------------------|---------------|--------|---------|
| | FM-R0 | FM-R25 | FM-R50 | FM-R75 | FM-R100 | | | |
| Proximate composition (g kg ⁻¹ wet weight basis) | | | | | | | | |
| Moisture | 695.07 ^a | 698.60 ^a | 697.03 ^a | 692.27 ^a | 692.27 ^a | 10.330 | 0.607 | 0.339 |
| Crude protein | 167.90 ^a | 166.03 ^a | 168.17 ^a | 168.07 ^a | 168.63 ^a | 6.737 | 2.032 | 0.893 |
| Crude lipid | 76.37 ^c | 77.03 ^c | 77.13 ^c | 85.10 ^b | 87.00 ^a | 0.518 | 1.172 | <0.001 |
| Crude fibre | 9.53 ^a | 9.37 ^a | 8.47 ^a | 9.27 ^a | 9.50 ^a | 0.578 | 10.844 | 0.714 |
| NFE ¹ | 6.37 ^a | 3.47 ^a | 2.87 ^a | 1.73 ^a | 0.73 ^a | 3.484 | 80.763 | 0.152 |
| Total ash | 44.80 ^a | 45.57 ^a | 46.33 ^a | 43.50 ^a | 41.83 ^a | 3.733 | 5.723 | 0.301 |
| Fatty acid composition (% total fatty acids) | | | | | | | | |
| C14:0 | 3.23 ^{ab} | 3.27 ^a | 3.19 ^{bc} | 3.15 ^c | 3.23 ^{ab} | 0.001 | 1.199 | 0.038 |
| C15:0 | 0.60 ^{bc} | 0.58 ^c | 0.61 ^b | 0.63 ^{ab} | 0.66 ^a | <0.001 | 2.834 | 0.005 |
| C16:0 | 16.74 ^a | 17.00 ^a | 16.98 ^a | 17.11 ^a | 17.10 ^a | 0.025 | 1.233 | 0.214 |
| C17:0 | 0.59 ^a | 0.58 ^a | 0.61 ^a | 0.57 ^a | 0.65 ^a | 0.001 | 5.894 | 0.129 |
| C18:0 | 6.07 ^c | 6.10 ^{bc} | 6.17 ^b | 6.25 ^a | 6.30 ^a | 0.001 | 0.688 | 0.001 |
| C20:0 | 0.79 ^a | 0.71 ^c | 0.78 ^a | 0.74 ^{bc} | 0.75 ^b | <0.001 | 2.083 | 0.002 |
| C22:0 | 0.70 ^{ab} | 0.64 ^{bc} | 0.71 ^{ab} | 0.60 ^c | 0.77 ^a | 0.001 | 6.388 | 0.011 |
| C24:0 | 0.16 ^{ab} | 0.19 ^a | 0.17 ^{ab} | 0.14 ^b | 0.15 ^b | <0.001 | 9.776 | 0.043 |
| C16:1 | 6.57 ^a | 6.58 ^a | 6.42 ^b | 6.32 ^c | 6.27 ^c | 0.001 | 0.675 | 0.000 |
| C17:1 | 0.30 ^a | 0.30 ^a | 0.28 ^a | 0.29 ^a | 0.30 ^a | <0.001 | 5.603 | 0.585 |
| C18:1n9 | 12.30 ^d | 12.48 ^c | 12.54 ^c | 12.66 ^b | 12.80 ^a | 0.001 | 0.278 | 0.000 |
| C18:1n7 | 3.31 ^a | 3.29 ^a | 3.32 ^a | 3.33 ^a | 3.35 ^a | 0.002 | 1.603 | 0.725 |
| C20:1 | 0.65 ^a | 0.53 ^{bc} | 0.57 ^b | 0.50 ^c | 0.55 ^{bc} | <0.001 | 4.480 | 0.001 |
| C22:1 | 0.14 ^a | 0.17 ^a | 0.16 ^a | 0.15 ^a | 0.16 ^a | <0.001 | 8.501 | 0.089 |
| C24:1 | 0.21 ^a | 0.19 ^a | 0.22 ^a | 0.19 ^a | 0.19 ^a | <0.001 | 11.803 | 0.457 |
| C18:2n6 | 15.95 ^c | 16.57 ^b | 16.84 ^{ab} | 17.04 ^{ab} | 17.26 ^a | 0.048 | 1.721 | 0.004 |
| C20:2n6 | 0.12 ^a | 0.13 ^a | 0.13 ^a | 0.13 ^a | 0.11 ^a | <0.001 | 14.374 | 0.604 |
| γ C18:3 | 0.80 ^b | 0.85 ^a | 0.80 ^b | 0.88 ^a | 0.86 ^a | <0.001 | 2.018 | 0.002 |
| α C18:3n3 | 1.18 ^c | 1.22 ^b | 1.19 ^c | 1.26 ^a | 1.23 ^b | <0.001 | 1.052 | 0.000 |
| C20:3n6 | 0.44 ^a | 0.42 ^a | 0.44 ^a | 0.45 ^a | 0.44 ^a | <0.001 | 2.935 | 0.181 |
| C20:4n6 | 1.59 ^b | 1.60 ^b | 1.66 ^a | 1.53 ^c | 1.38 ^d | <0.001 | 1.671 | 0.000 |
| C20:5n3 | 2.91 ^a | 2.82 ^{ab} | 2.74 ^{bc} | 2.67 ^c | 2.39 ^d | <0.001 | 1.845 | 0.000 |
| C22:6n3 | 7.05 ^a | 6.87 ^b | 6.92 ^{ab} | 5.85 ^c | 5.34 ^d | 0.003 | 1.151 | 0.000 |

Means bearing different superscript letters within the row differ significantly ($p < 0.05$).

¹Nitrogen free extract (Calculated by difference)

fatty acids was low in carcass than the experimental diet due to the selective utilisation of such fatty acids for energy production (Caballero *et al.*, 2002). Fish fed diet with FM-R100 (complete replacement) showed ($p < 0.05$) a low level of eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA) compared to other treatments, however, the sum of polyunsaturated fatty acids (PUFAs) content was not affected due to the higher content of C18:2 n-6. The result is in agreement with the findings obtained with gilthead seabream (Izquierdo *et al.*, 2003). Gonzalez-Felix *et al.* (2002) reported that a plant-based diet, in which 90% of fish oil was replaced by soy oil, had no adverse effects on growth performance, but dramatically reduced body PUFAs content. This result is in agreement with the findings of Montero *et al.* (2005) in *D. labrax* while replacing fish oil using various plant oils *viz.* rapeseed, linseed and palm oils. The authors suggested to switch from plant oil to fish oil containing diet (finisher diet) prior to harvesting for an appropriate time to restore both EPA and DHA as they are most important fatty acids in human nutrition. A study with juveniles of *Penaeus monodon* (Khan, 2013) showed that a period of 30-days was required to restore both EPA and DHA by using a finisher diet (control diet with no fishmeal replacement) in both field and laboratory conditions, whereas finishing diet phase was 16 to 20 weeks in salmon fed a blend of rapeseed, linseed and fish oil (Bell *et al.*, 2004). The variation in time phase between the studies could be due to species difference, weight of species, rearing and environmental conditions. Our data demonstrated that fatty acids like C16:0, C18:1n-9 and C18:2n-6 were abundant in fish carcass and formed nearly 40% of the total fatty acids. Fish fed post-experimental diets had a higher content of PUFAs (30.48 to 32.55%) and saturated fatty acids (SFAs) (29.62 to 31.70%) compared to mono-unsaturated fatty acids (MUFAs) (23.48 to 23.65%). This result is contrary to the findings in other species (Paleari *et al.*, 1997). Chen *et al.* (1995) reported a higher content of MUFAs instead of SFAs. Among the n-6 series of PUFAs, C18:2 was predominant and gradually increased ($p < 0.05$) with increasing inclusion level of plant protein sources. In n-3 series of PUFAs, C22:6 dominated and gradually decreased ($p < 0.05$) due to the inclusion of plant proteins. The sum of n-6 fatty acids increased in fish carcass with increasing inclusion level of plant proteins, while the reverse trend was noticed in n-3 series, hence, the ratio of n-3/n-6 reduced from 0.72 to 0.50.

The present investigation demonstrated that a combination of plant protein sources could substitute 50% of fishmeal in the diet of Asian seabass without having any adverse effect on growth performance. The findings of digestive enzyme profiles have given an insight into how the fish is adapting to fishmeal substitution by modulating

enzyme profiles. Feeding fish with finisher diet (no fishmeal replacement) is imperative to restore the reduced PUFAs, in particular EPA and DHA when fishmeal has been replaced beyond 50%. From the present results, it can be concluded that a blend of plant protein sources could be a potential alternative to fishmeal. Further studies on supplementation of crystalline amino acids, phosphorus, phytase and customised digestive enzymes could help to maximise utilisation of these ingredients by reducing inclusion level of fishmeal in the diet of Asian seabass.

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