



A Challenge to Conventional Fish Meal: Effects of Soy Protein Peptides on Growth, Histomorphology, Lipid Metabolism and Intestinal Health for Juvenile Pompano *Trachinotus ovatus*

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This experiment was conducted to assess the possibility of replacing fish meal (FM) with soy protein peptide (SPP) at different levels—0% (FM), 14.29% (S5), 28.57% (S10), 57.14% (S20), 71.42% (S25)—and its effects on growth, histology, gene expression related to liver lipid metabolism and intestinal immunity in juvenile pompano *Trachinotus ovatus* (initial mean weight = 39.88 ± 0.15 g). 600 healthy and uniformed-size fish were distributed to five groups of three replicates, each with 40 fish in each floating cage and fed twice daily for 8 weeks. Results showed that no significant difference in the growth was observed with SPP replacing FM ($P > 0.05$). Serum glutathione peroxidase activity in the S10 group was significantly higher than that in the FM group, and serum malondialdehyde content significantly decreased ($P < 0.05$). SPP significantly improved intestinal immunity by increasing alkaline phosphatase and lysozyme activities and up-regulating interleukin 10 and complement 4 mRNA levels while simultaneously decreasing triglyceride and total cholesterol content and down-regulating interleukin 1β mRNA expression. Villus length and muscle thickness in the S10 group were significantly higher than those in the FM group ($P < 0.05$). SPP significantly improved liver fat metabolism by increasing carnitine palmitoyl transferase I mRNA levels, and down-regulating fatty acid synthesis mRNA expression ($P < 0.05$). In summary, SPP substitution for FM promoted intestinal health, liver lipid metabolism and reduced liver fat accumulation for juvenile pompano *T. ovatus*, with no significant effect on growth performance. Based on the second-order polynomial analysis model of LYZ activity, the optimal replacement SPP level for juvenile pompano *T. ovatus* was 11.82%.

Keywords: *Trachinotus ovatus*, soy protein peptide, fish meal, histological morphology, lipid metabolism

HIGHLIGHTS

- The replacement of FM with partial SPP can significantly improve disease resistance without affecting growth performance for juvenile pompano *Trachinotus ovatus* under experimental conditions.
- By analyzing serum immune indexes, intestinal histology and intestinal immune-related genes, the replacement of 10% FM with SPP significantly promotes the intestinal health for juvenile pompano *Trachinotus ovatus*.
- Based on the broken-line regression analysis model of LYZ activity in serum, the SPP optimal replacement level for juvenile pompano *Trachinotus ovatus* is 12.80%.

INTRODUCTION

Fish meal (FM) has high amounts of proteins, essential amino acids, N-3 polyunsaturated fatty acids, and vitamins and has been the most used protein source in the compound feed of pompano *Trachinotus ovatus* (Zhang et al., 2018). However, the declining world fish meal production and the rising prices lead to an increasingly prominent conflict between supply and demand (Tacon and Metian, 2008). The increasing cost of feed has severely limited the sustainable development of the aquafeed industry. Therefore, nutrient-rich, efficient, and environmentally friendly plant protein sources as replacements for FM has become a major topic in the aquatic feed industry (Food and Agriculture Organization [FAO], 2014).

Alternative protein sources are essential to the growth of the aquaculture industry (Tacon, 2003) and can be used to reduce protein sources from FM (Naylor et al., 2000). The final products of protein digestion in the digestive tract mainly originate from small peptides rather than from free amino acids, and enter the blood circulation in the form of two or three peptides (Liu et al., 2007; Zhou et al., 2011). Peptides as feeding materials have been widely studied. Peptides can strengthen the immune system, reduce feed coefficient rate, and accelerate protein synthesis (Liang et al., 2020a; Wang J.X. et al., 2020). In *Gadus morhua* (Aksnes et al., 2006a), *Salmo salar* (BØGwald et al., 1996), *Dicentrarchus labrax* (Kotzamanis et al., 2007), and *Litopenaeus vannamei* (Gyan et al., 2020), similar improvements in weight gain rate, feed efficiency, and survival were observed (Savoie et al., 2006). In addition, small peptides preferentially act as energy substrates for the intestinal mucosa and can effectively promote the development of intestinal mucosal tissue (Zheng et al., 2006). Soybean protein peptide (SPP) is a mixed oligopeptide prepared through protease hydrolysis or microbial fermentation (Zhang J. et al., 2020). It has no anti-nutritional factors and is rich in amino acids, low-molecular-weight small peptide and special nutrients (Puchalska et al., 2014). In addition, it is easy to digest and absorb, and has physiologically active substances that promote fat metabolism (Huang, 2015). Previous studies showed that SPP had many advantages, including antioxidant (Ma et al., 2016), cholesterol-lowering and immune-enhancing activities (Zhao Z. X. et al., 2016; Cheng et al., 2017; Zhang Y. J. et al., 2020).

Pompano *T. ovatus*, a valuable food fish with tender flesh and delicious flavor, is mainly distributed in the warm tropical waters of the Pacific Ocean, Indian Ocean, and Atlantic Ocean (Tutman et al., 2004; Li M. M. et al., 2020). This fish thrives in tropical and subtropical climates due to its delicious flesh, rapid growth, simple feeding habits and high survival rate (Qu et al., 2014; Zhang et al., 2019). The annual production of *T. ovatus* in China has exceeded 100,000 tons in 2020, promising candidates for intensive aquaculture in Southern China with annual production of around 120,000 tons (Li M. et al., 2020). Plant protein sources in Pompano *T. ovatus* feed, including soybean meal (Wu et al., 2010; Niu et al., 2016), fermented soybean meal (Lin et al., 2012), rapeseed meal (Kou et al., 2015), and cottonseed protein concentrate (Shen et al., 2019), have attracted considerable interest. Meanwhile, SPP has not been reported. In this experiment, juvenile Pompano *T. ovatus* was used as a research target, and different proportions of FM were substituted with SPP to prepare iso-nitrogenous and iso-lipid test feeds. This study aimed to investigate the effects of the replacement of FM with SPP on growth, histomorphology, liver fat metabolism, and intestinal immune function for juvenile Pompano *T. ovatus*.

MATERIALS AND METHODS

Diet Formulation and Preparation

The experimental diet was formulated to provide 41.00% of crude protein (Table 1), and the dietary amino acid composition was shown (Table 2). The proximate compositions and amino acid composition of FM and SPP used in this study were presented in Table 3. The proportion of SPP with a relative molecular mass of less than 1,000 Da was 100%, provided by Yisheng Biotechnology Company (Yangjiang, Guangdong, China). Five iso-nitrogenous and iso-lipidic experiment diets were formulated to contain 0, 5.45, 10.90, 21.78, and 27.23% of SPP by replacing 0, 14.29, 28.57, 57.14, and 71.42% of FM, respectively. Reduction in feed by 0% (FM, control group), 5% (S5), 10% (S10), 20% (S20), and 25% (S25) FM, respectively. FM and soybean meal were the main sources of protein in the diet. Fish oil, soybean oil, and soybean lecithin oil were the lipid sources.

All raw material was crushed through a 60-mesh sieve, mixed thorough a V-type vertical mixer (JS-14S; Zhejiang Zhengtai Electric Co., Ltd.), followed by adding with oil and water, and then pelleted (2.5 mm diameter) making use of a double screw extruder (F-75; South China University of Technology). After the prepared experimental feed was naturally dried to about 10% moisture, it was sealed in a vacuum-packed bag and stored at -20°C until it was fed (He et al., 2021; Lin et al., 2021).

Experimental Animals and Breeding Management

The experiment was conducted at an experimental site in Zhanjiang, Guangdong, China. Juvenile *T. ovatus* was procured from a seedling farm in Hainan Province for this investigation. Live fish were transported from Hainan to Guangdong using live fish transport techniques and cultured in a floating cage

TABLE 1 | Formulation and proximate composition of experimental diet (% dry matter).

Ingredients	Diets				
	FM	S5	S10	S20	S25
Brown fish meal	35.00	30.00	25.00	15.00	10.00
Soybean protein peptide	0.00	5.45	10.90	21.78	27.23
Soybean meal	20.00	20.00	20.00	20.00	20.00
Peanut meal	4.00	4.00	4.00	4.00	4.00
Corn gluten meal	4.00	4.00	4.00	4.00	4.00
Wheat flour	20.00	20.00	20.00	20.00	20.00
Calcium dihydrogen phosphate	1.50	1.50	1.50	1.50	1.50
Vitamin C	0.05	0.05	0.05	0.05	0.05
Choline chloride	0.30	0.30	0.30	0.30	0.30
Soybean lecithin oil	1.50	1.50	1.50	1.50	1.50
Soybean oil + fish oil (1:1)	4.18	4.35	4.53	4.89	5.07
Vitamin premix ^a	0.50	0.50	0.50	0.50	0.50
Mineral premix ^b	0.50	0.50	0.50	0.50	0.50
Methionine	0.41	0.47	0.53	0.65	0.71
Lysine	0.24	0.34	0.44	0.64	0.74
Threonine	0.00	0.03	0.06	0.13	0.16
Arginine	0.13	0.12	0.12	0.11	0.10
Microcrystalline cellulose	7.69	6.89	6.07	4.46	3.64
Total	100.00	100.00	100.00	100.00	100.00
Nutrient levels					
Crude protein ^c	40.48	40.72	40.43	40.71	40.21
Crude lipid ^c	7.97	8.27	8.26	8.16	8.02
Crude ash ^c	8.76	8.37	7.84	6.97	6.41

^aThe vitamin premix: vitamin A 500,000 IU/kg; vitamin D₃ 100,000 IU/kg; vitamin E 4,000 mg/kg; vitamin K₃ 1,000 mg/kg; vitamin B₁ 500 mg/kg; vitamin B₂ 1,000 mg/kg; vitamin B₆ 1,000 mg/kg; vitamin B₁₂ 2.0 mg/kg; nicotinic acid 4,000 mg/kg; D-calcium pantothenate 2,000 mg/kg; folic acid 100 mg/kg; biotin 10.0 mg/kg; vitamin C 15,000 mg/kg.

^bMineral mixture: Fe 10,000 mg/kg; Cu 300 mg/kg; Zn 5,000 mg/kg; Mn 1,200 mg/kg; I 80 mg/kg; Se 30 mg/kg; Co 20 mg/kg.

^cCrude protein, crude lipid and ash contents were measured values.

(length: 5 m; width: 5 m; and height: 5 m) (Harmon, 2009; Zhang et al., 2021). Before the experiment, all fish were fed commercial diets (Zhanjiang Yuehai Feed Co. Ltd., Guangdong, China; crude protein 42%, crude lipid 8%) for a week to acclimatize to the conditions. At this experiment, the total number of the experimental fish was 600, and there were 40 fish (initial body weight: 39.88 ± 0.15 g) in each floating cage (length: 1 m; width: 1 m; and height: 2 m), respectively. The 15 groups were randomly assigned to the five test diets, with three replicates each. All fishes were fed twice daily (7:00 and 17:00) to visual satiety for 8 weeks. During the experiment, water temperature ranged from 29.0 to 31.0°C, the salinity was 24–26‰ and the dissolved oxygen was not less than 6 mg/L.

All procedures involving live animals were approved by the Guangdong Ocean University Institutional Animal Care and Use Committee.

Sample Collection

Fish samples were kept fast for 24 h before collection at the end of trial and anesthetized with MS-222 (1:10,000; Zhou

TABLE 2 | Amino acid composition of the experimental diet (% dry matter).

	Experimental diets				
	FM	S5	S10	S20	S25
EAA ^a					
Threonine	1.48	1.51	1.50	1.56	1.52
Valine	1.82	1.80	1.79	1.72	1.74
Methionine	1.05	1.03	0.98	0.81	0.82
Leucine	3.04	3.07	3.09	3.06	3.08
Phenylalanine	1.76	1.78	1.79	1.87	1.84
Lysine	2.59	2.62	2.65	2.64	2.63
Histidine	1.11	1.10	1.07	1.08	1.02
NEEA ^b					
Aspartic acid	3.49	3.57	3.61	3.75	3.74
Serine	1.59	1.66	1.64	1.78	1.68
Glutamic acid	6.68	6.85	7.01	7.37	7.39
Glycine	1.98	1.93	1.86	1.75	1.70
Alanine	2.19	2.14	2.03	1.96	1.84
Isoleucine	1.59	1.59	1.62	1.58	1.64
Tyrosine	1.18	1.20	1.10	1.18	1.16
Proline	2.41	2.44	2.44	2.51	2.52
Arginine	2.03	2.08	2.07	2.14	2.07
Cystine	0.41	0.45	0.46	0.45	0.45

^aEAA, essential amino acids; ^bNEEA, non-essential amino acids.

et al., 2019). The fish samples of each experimental group were calculated and weighed, and it was determined that weight gain ratio, the survival rate, specific growth rate and feed conversion ratio. Afterward, three fish samples of each floating cage were randomly used to test the body length and body weight to make a calculation of the hepatosomatic index, condition factor and viscerosomatic index, and then stored at −20°C to detect the whole-body composition (Wang et al., 2018; Wu et al., 2021). Blood samples were gotten from the tail veins of seven randomly selected fish samples from each floating cage, and then stored at 4°C for 12 h (Chen et al., 2016; Cai et al., 2020). After centrifuged (4,000 × g, 4°C, 10 min), the serum of fish samples was obtained and immediately preserved at −80°C for analyzing serum biochemical (Wang J. et al., 2020). Furthermore, the fish liver and intestine samples were quickly removed, one part was stored in 4% paraformaldehyde solution for histological analysis (Cai et al., 2021), whereas the others were stored in RNA-later at −80°C before RNA isolation.

Formula for Calculations

$$\text{Weight gain rate (WGR, \%)} = 100 \times \frac{[\text{final body weight (g)} - \text{initial body weight (g)}]}{\text{initial body weight (g)}}$$

$$\text{Feed coefficient rate (FCR)} = \frac{\text{feed intake (g)}}{[\text{final body weight (g)} - \text{initial body weight (g)}]}$$

TABLE 3 | Proximate composition of the fishmeal and soy protein peptide (% dry matter).

Ingredient	FM	SPP
Nutrients		
Moisture	8.15	8.00
Crude protein	70.03	60.87
Crude lipid	7.57	3.68
Amino acids		
Aspartate	5.80	4.80
Threonine	2.76	1.54
Serine	2.51	2.01
Glutamic	8.73	7.75
Glycine	3.99	1.68
Alanine	4.22	1.82
Cystine	0.57	0.00
Valine	3.04	1.74
Methionine	1.94	0.15
Isoleucine	2.73	1.67
Leucine	4.82	2.99
Tyrosine	2.24	1.30
Phenylalanine	2.66	1.97
Lysine	5.18	2.43
Histidine	2.28	1.21
Arginine	3.75	2.54
Proline	2.51	1.96

$$\text{Specific growth rate (SGR, \%)} = 100 \times [\ln(\text{final body weight(g)}) - \ln(\text{initial body weight (g)})] / \text{days};$$

$$\text{Survival rate (SR, \%)} = 100 \times (\text{final fish number} / \text{initial fish number});$$

$$\text{Hepatic somatic indices (HSI, \%)} = 100 \times \text{hepatic weight (g)} / \text{body weight (g)};$$

$$\text{Condition factor (CF, g/cm}^3\text{)} = 100 \times [\text{body weight (g)}] / [\text{body length (cm)}]^3;$$

$$\text{Daily feed intake (DFI, \% / days)} = 100 \times \text{feed intake} / [(\text{initial body weight} + \text{final body weight}) / 2 \times \text{experimental period}].$$

Chemical Analyses

The ingredients of the experimental diets and fish samples (crude protein, crude lipid, moisture, and ash) were measured by using standard methods AOAC (Association of Official Analytical Chemists [AOAC], 2005).

The total protein (TP), triglyceride (TG), total cholesterol (T-CHO), and glucose (GLU) in serum were assayed using an automatic blood analyzer (Hitachi 7020, Hitachi Science Systems, Japan), following a previously described method of Gyan et al. (2020).

The immune enzyme activities were determined by using a detection kit (Nanjing Jian Cheng Bioengineering Institute, China). The total superoxide dismutase (T-SOD) was measured by the xanthine oxidase method according to Wang et al. (2011). The activity of glutathione peroxidase (GSH-Px) was determined using the xanthine oxidase method (Ma et al., 2014). The total antioxidant capacity (T-AOC) was measured by Yang et al. (2017). The malondialdehyde (MDA) content was determined as described by Liang et al. (2020a) and Lin et al. (2020). The acid phosphatase (ACP) and alkaline phosphatase (AKP) were determined following Zhu et al. (2012) and Zhu et al. (2021). The lysozyme (LYZ) activity was assayed according to Liu et al. (2021).

Real-Time PCR Analysis of the Organization

Total RNA from the liver and hind intestines was extracted with an RNA extraction kit (TransZol Up Plus RNA Kit, Beijing, China). PrimeScriptTM RT-PCR Kit (TaKaRa, Kusatsu, Japan) was used to synthesize complementary DNA (cDNA) according to the manufacturer's instructions. The PCR primers were listed in Table 4. The PCR cycling protocol by Liang et al. (2020b) was used, and all the real-time PCR reactions were performed on a Roche LightCycler480II (Switzerland) using an SYBR @ Premix Ex TaqTM Kit (Takara). The relative mRNA expressions were calculated using the $2^{-\Delta\Delta CT}$ method.

Histological Morphology

The intestinal tracts and livers of the fish were quickly removed, and one part was stored in 4% paraformaldehyde solution for histological analysis using hematoxylin-eosin (H&E) (Martínez-Llorens et al., 2012). The tissue sections were observed and photographed under an electron microscope scanner (VS 120-S6, Olympus, Norway) with villi length and muscle thickness measured according to the method of Wang J.X. et al. (2020). Sections from each floating cage were randomly measured for villi lengths and muscle layer thicknesses (Lin et al., 2019).

Statistical Analysis

All data were subjected to ANOVA using SPSS 21.0 (SPSS Inc., Chicago, IL, United States). Tukey's HSD multiple comparisons were performed in the case of a significant overall difference between the experimental group and the control ($P < 0.05$). The results were presented as the mean \pm SEM (standard error of the mean).

TABLE 4 | Sequences of primers used for real-time quantitative PCR.

Gene name	Primer sequence (5'-3')	References
<i>c4</i>	F-TGGAGAAAAAGTTAAAGGGGC R-CAGGAAGGAAGTATGAGCGAGT	Tan et al., 2018
<i>nf-κb</i>	F-TGCGACAAAGTCCAGAAAGAT R-CTGAGGGTGGTAGGTGAAGGG	Zhou et al., 2020
<i>il-8</i>	F-GAGAAGCCTGGGAATGGA R-GAGCCTCAGGGTCTAAGCA	Zhou et al., 2020
<i>il-10</i>	F-CTCCAGACAGAAGACTCCAGCA R-GGAATCCCTCCACAAAACGAC	Tan et al., 2017
<i>il-1β</i>	F-CGGACTCGAACGTGGTCACATTC R-AATATGGAAGGCAACCGTGCTCAG	Xie et al., 2020
<i>cptI</i>	F-CTTTAGCCAAGCCCTTCATC R-CACGGTTACCTGTTCCCTCT	Liu et al., 2018
<i>fsan</i>	F-GAAGGAGAGGGGGTGGAGTC R-GTGTGAAGGTGGAGGGTGTG	Liu et al., 2018
<i>apob100</i>	F-AAAAGCCACAAGACGAAAGCA R-GAAGCAGCAAAAAGGCAGAGC	Liu et al., 2018
<i>srebp-1</i>	F-GAGCCAAGACAGAGGAGTGT R-GTCTCTTGTCTCCAGCTT	Li et al., 2020
<i>fabpI</i>	F-AGTCATTGTCTGGGGAGGG R-GTCAAGGCGGTGGTTCA	Liu et al., 2018
β -actin	F-TACGAGCTGCCTGACGGACA R-GGCTGTGATCTCCTTCTGC	Xie et al., 2019

c4, complement 4; *nf- κ b*, nuclear factor kappa B; *il-8*, interleukin 8; *il-10*, interleukin 10; *il-1 β* , interleukin 1 β ; *cptI*, carnitine palmitoyl transferase I; *fsan*, fatty acid synthesis; *apob100*, apolipoprotein B-100; *srebp-1*, sterol-regulatory element binding protein-1; *fabpI*, fatty acid binding protein I.

RESULTS

Growth Performance and Whole-Body Composition

No significant differences in IBW, WGR, SGR, SR, FCR, CF, and DFI were found among the groups ($P > 0.05$; **Table 5**). As SPP increased, VSI firstly decreased and then increased, and the S10 group was significantly lower than the FM group ($P < 0.05$). As SPP level increased, HSI gradually decreased and was significantly lower in the S10, S20, and S25 groups than in the FM group ($P < 0.05$).

The results of the whole-body composition for juvenile pompano *T. ovatus* are shown in **Table 6**. No significant differences were found in the whole-body composition among juvenile pompano *T. ovatus* fed with different amounts of dietary SPP ($P > 0.05$).

Serum Antioxidant and Immune Enzyme Activities

As SPP level increased, AKP activity showed an increasing trend followed by a decreasing trend (**Table 7**; $P < 0.05$). AKP activity in the S10 group was significantly higher than that in the control group ($P < 0.05$). ACP activity showed an increasing tendency followed by a decreasing trend, but the difference among the groups was not significant ($P > 0.05$). LYZ activity increased and then decreased, showing a trend change in line with AKP activity. LYZ activity in the S10 group was significantly higher than that in the control group ($P < 0.05$). According to the second-order

polynomial analysis model of LYZ activity, the SPP replacement level of juvenile pompano *T. ovatus* was 11.82%, corresponding to SPP replacement level (**Figure 1**).

T-SOD activity in the S5, S10, and S20 groups was not significantly different from that of the control group, or the high substitution group (S25) had significantly lower T-SOD activity than that in the control group. The GSH-Px activity showed an increase followed by a decrease. GSH-Px activity in the S10 and S20 groups was significantly higher than in the FM group ($P < 0.05$). As SPP increased, MDA content showed a gradually decreasing trend, and the groups in S10, S20, and S25 were significantly lower than that in the FM group ($P < 0.05$). No significant difference in T-AOC activity was found among the groups ($P > 0.05$), but T-AOC activity was higher in the substitution groups than in the FM group.

Serum Biochemical Indices

TP content in the S5, S10, and S20 groups were significantly higher than that in the FM group (**Table 8**, $P < 0.05$), but difference in TP content between the S25 and FM groups was not significant ($P > 0.05$). As SPP increased, TG levels gradually decreased, and the FM groups had the highest TG levels ($P < 0.05$). T-CHO and TG levels in the experimental group gradually decreased and were significantly lower than those in the FM group ($P < 0.05$). No significant differences in glucose content were found among the groups ($P > 0.05$).

Liver and Intestinal Morphology

Observation of liver tissue (**Figure 2**) showed that serious vacuolation occurred in the FM group, but the cell boundaries of FM, S5, and S10 groups were obvious, and the rate of intracellular vacuolization decreased. The nuclei of hepatocytes in the S20 group began to gradually lyse or disappear, and the nuclei of hepatocytes increased in spacing and showed blurred outlines after cell disintegration.

We stained the liver with oil red to observe fat deposition in the liver (the fat was stained red, **Figure 3**). Fat deposition tended to decrease with increasing SPP level. The fat droplets in the FM group were large and dense, whereas the fat droplets were significantly smaller and the deposition sites were more dispersed in the experimental groups.

The hind-gut structure data of juvenile *T. ovatus* are provided (**Figures 4, 5**). As SPP level increased, villus length (VL) increased first and then decreased and was significantly higher in the S10 group than in the FM group ($P < 0.05$). The trend of muscle thickness (MT) was in line with that of VL. All substitution groups were significantly higher than the FM group ($P < 0.05$), and the S10 group was higher than all other groups.

Expression of Lipid Metabolism-Related Genes in the Liver

In the present study, the expression of lipid metabolism-related genes in the liver were analyzed (**Figure 6**). The hepatic carnitine palmitoyl transferase I (*cptI*) mRNA levels in the S5 and S10 groups were significantly higher than those in the other groups ($P < 0.05$). The fatty acid synthesis (*fsan*) mRNA

TABLE 5 | Growth performance and biometry for juvenile *Trachinotus ovatus* fed the experimental diet.

Parameters	Experimental diets				
	FM	S5	S10	S20	S25
IBW (g)	39.97 ± 0.21	39.94 ± 0.27	39.91 ± 0.08	39.75 ± 0.14	39.88 ± 0.18
WGR (%)	117.67 ± 3.45	118.10 ± 4.09	117.70 ± 3.24	111.82 ± 0.79	114.76 ± 1.04
SGR (%/d)	1.39 ± 0.03	1.39 ± 0.03	1.38 ± 0.04	1.34 ± 0.01	1.39 ± 0.04
FCR	2.43 ± 0.10	2.37 ± 0.10	2.39 ± 0.15	2.56 ± 0.05	2.53 ± 0.07
SR (%)	99.17 ± 1.44	100.00 ± 0.00	99.17 ± 1.44	98.33 ± 2.89	99.17 ± 1.44
VSI (%)	4.70 ± 0.29 ^b	4.21 ± 0.23 ^{ab}	4.06 ± 0.04 ^a	4.39 ± 0.12 ^{ab}	4.58 ± 0.18 ^b
HSI (%)	0.99 ± 0.12 ^b	0.98 ± 0.07 ^b	0.65 ± 0.03 ^a	0.54 ± 0.05 ^a	0.57 ± 0.03 ^a
CF (g/cm ³)	4.31 ± 0.04	4.35 ± 0.07	4.32 ± 0.14	4.55 ± 0.25	4.33 ± 0.06
DFI (%/days)	3.15 ± 0.07	3.14 ± 0.06	3.16 ± 0.10	3.23 ± 0.02	3.21 ± 0.04

Data are mean ± S.E.M. (n = 3). Values in the same row with different superscripts represent significant difference (P < 0.05).

IBW, initial mean body weight; WGR, weight gain rate; SGR, specific growth rate; FCR, feed coefficient rate; SR, survival rate; VSI, viscerosomatic index; HSI, hepatic somatic indices; CF, condition factor; DFI, daily feed intake.

TABLE 6 | The composition of whole body for juvenile *Trachinotus ovatus* fed the experimental diet (% dry matter).

Parameters	Experimental diets				
	FM	S5	S10	S20	S25
Moisture	68.91 ± 0.94	69.53 ± 0.56	69.28 ± 0.85	69.75 ± 0.34	67.37 ± 1.44
Crude protein	59.04 ± 0.64	60.16 ± 0.50	61.31 ± 0.39	61.06 ± 0.79	59.05 ± 1.77
Crude lipid	27.56 ± 0.47	27.15 ± 1.38	26.96 ± 2.92	25.79 ± 1.32	27.53 ± 2.98
Crude ash	13.32 ± 0.21	13.30 ± 0.44	12.69 ± 0.10	14.16 ± 0.34	12.85 ± 1.52

Data are mean ± S.E.M. (n = 3). Values in the same row with different superscripts represent significant difference (P < 0.05).

level was significantly down-regulated after the replacement of FM with SPP in compound feed (P < 0.05). The S25 group had significantly lower fatty acid-binding protein I (*fabp1*) and apolipoprotein B-100 (*apob100*) mRNA levels than the FM group (P < 0.05). No significant difference in sterol-regulatory element-binding protein-I (*srebp-1*) mRNA level was found between the FM and replacement groups (P > 0.05).

Expression of Intestinal Immunity-Related Genes

The expression of intestinal immunity-related genes of juvenile *T. ovatus* is shown (Figure 7). The expression of genes involved in intestinal immunity-related genes, including interleukin 10 (*il-10*) and complement 4 (*c4*), was significantly up-regulated after FM was replaced with SPP (P < 0.05). As SPP level increased, interleukin 1β (*il-1β*) mRNA level was significantly down-regulated (P < 0.05). No significant difference in nuclear factor kappa B (*nf-κb*) mRNA level was observed between the FM group and the replacement groups (P > 0.05).

DISCUSSION

Soya bean meal has a high crude protein content and is rich in essential amino acids and widely available. The presence of anti-nutritional factors limits the use of soybean meal compared with FM (Yildirim et al., 2009). Indeed, excessive levels of plant proteins in feed can inhibit animal growth (Floreto et al., 2000).

However, fermented soy peptides are rich in organic acids, bacterial active proteins, folic acid, and B vitamins and have increased palatability (Cao et al., 2007). In the present study, the replacement of FM with SPP resulted in no significant differences in the WGR and SGR of juvenile *T. ovatus* cultured in floating cages under experimental conditions. We recommend that it is feasible to replace FM with SPP in compound feeds, and the studies on which is consistent with *Epinephelus akaara* (Zhao S. Y. et al., 2016) and *Acipenser baerii* (Wang et al., 2010). This may be related to the nutritional profile of SPP, which contains nutrients, such as low-molecular-weight peptides, vitamins, and other special nutrients, making up for the shortcomings of many plant proteins (Puchalska et al., 2014). Moreover, small peptides are preferentially used as energy substrates for the structural and functional development of intestinal mucosal epithelial cells, effectively promoting the development of intestinal mucosal tissues (Wang et al., 2003). In the present study, the replacement of FM with SPP significantly promoted intestinal growth. Therefore, the high content of low molecular weight peptides (Puchalska et al., 2014) and intestinal growth are responsible for the unaffected growth rate for juvenile *T. ovatus*.

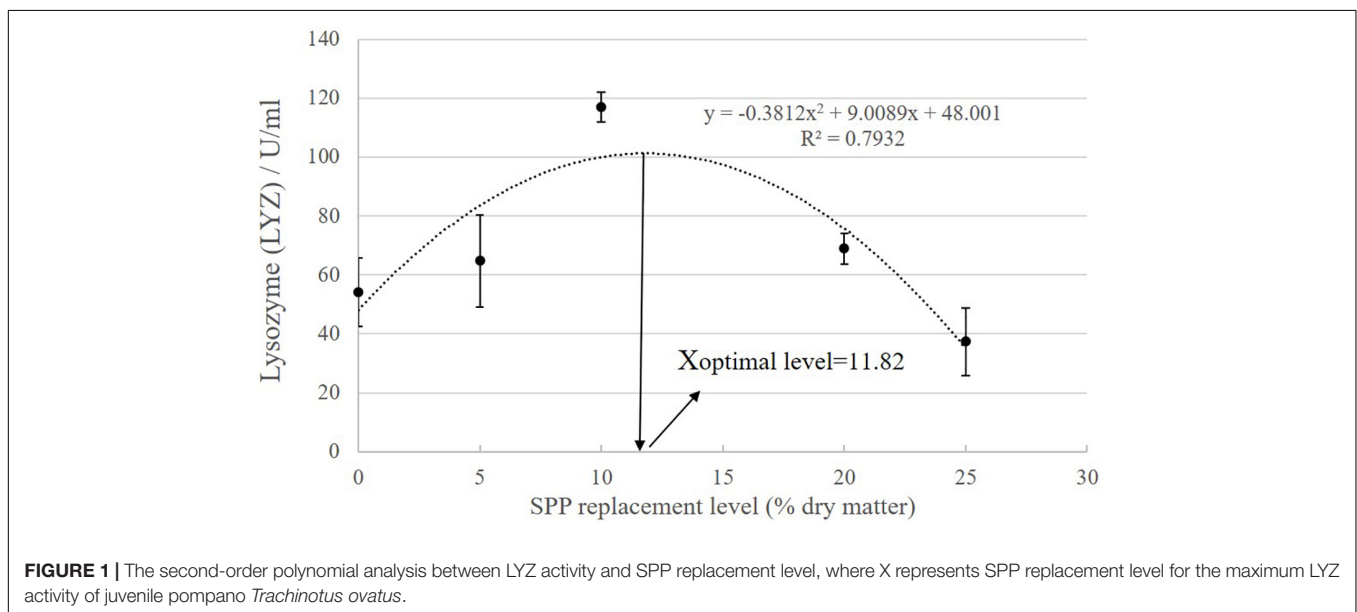
Body composition can directly reflect animal growth and indirectly reflect feed quality (Li et al., 2009). In the present study, no significant differences in moisture content and crude protein content were found among the groups. This study is in line with the studies on *Pelteobagrus fulvidraco* (Jing et al., 2021) and *Heterotis niloticus* (Monentcham et al., 2010). As SPP increased, the body fat content of fish decreased gradually, and all groups

TABLE 7 | Serum enzyme activity indices for juvenile *Trachinotus ovatus* fed the experimental diet.

Parameters	Experimental diets				
	FM	S5	S10	S20	S25
Serum immune ability					
AKP (U/L)	34.01 ± 9.19 ^a	41.40 ± 6.37 ^a	51.61 ± 6.54 ^b	41.08 ± 3.31 ^a	30.24 ± 1.96 ^a
ACP (U/L)	41.37 ± 0.71	41.83 ± 1.47	41.59 ± 3.19	40.41 ± 2.48	39.94 ± 0.71
LYZ (U/mL)	54.17 ± 11.53 ^{ab}	64.71 ± 15.55 ^{ab}	117.01 ± 4.97 ^c	68.92 ± 5.29 ^b	37.30 ± 11.47 ^a
Serum oxidation resistance					
T-SOD (U/ml prot)	35.41 ± 4.16 ^{bc}	42.81 ± 2.01 ^c	30.95 ± 6.85 ^{abc}	25.37 ± 3.62 ^{ab}	22.31 ± 5.57 ^a
GSH-Px (U/g prot)	25.26 ± 6.32 ^a	48.42 ± 3.65 ^{ab}	56.84 ± 10.94 ^b	56.84 ± 10.94 ^b	44.21 ± 12.63 ^{ab}
T-AOC (mM)	0.80 ± 0.01	0.82 ± 0.04	0.85 ± 0.02	0.81 ± 0.01	0.81 ± 0.01
MDA (nmol/mL)	6.36 ± 0.61 ^b	6.06 ± 0.30 ^{ab}	4.85 ± 0.30 ^a	4.85 ± 0.80 ^a	4.75 ± 0.35 ^a

Data are mean ± S.E.M. (n = 3). Values in the same row with different superscripts represent significant difference (P < 0.05).

AKP, alkaline phosphatase; ACP, acid phosphatase; LYZ, lysozyme; T-SOD, total superoxide dismutase; GSH-Px, glutathione peroxidase; T-AOC, total antioxidant capacity; MDA, malondialdehyde.



had lower fat content than the control group. In *Platichthys stellatus* (Jiang, 2013), small peptides could significantly reduce the amount of crude fat in the body. Moreover, Hou (2012) showed that SPP promoted fat and energy metabolism and inhibited fat deposition in the body. These effects are due to small peptides, which improve protein synthesis, reduce the deposition of free amino acids into fat (Boza et al., 2000), impede the absorption of fat and promote lipid metabolism (Christian, 2005). These results indicated that the replacement of FM with SPP can increase protein synthesis, promote the consumption of excess fat, and reduce fat deposition.

Condition factor, HSI, and VSI are important indices that reflect the body fat and lean status and growth status of fish (Aksnes et al., 2006a,b). In the present study, as SPP levels increased, the HSI and VSI significantly decreased, consistent with those of a hybrid grouper (*Epinephelus lanceolatus* ♂ × *Epinephelus fuscoguttatus* ♀; Jiang et al., 2015). This result may be related to the absorption mechanism of small

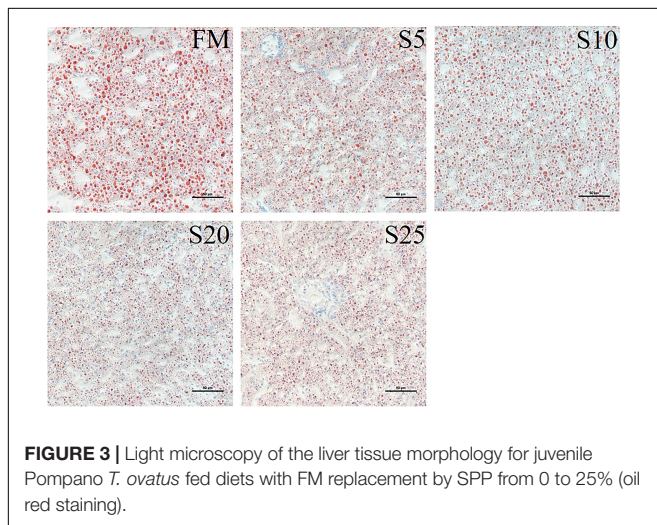
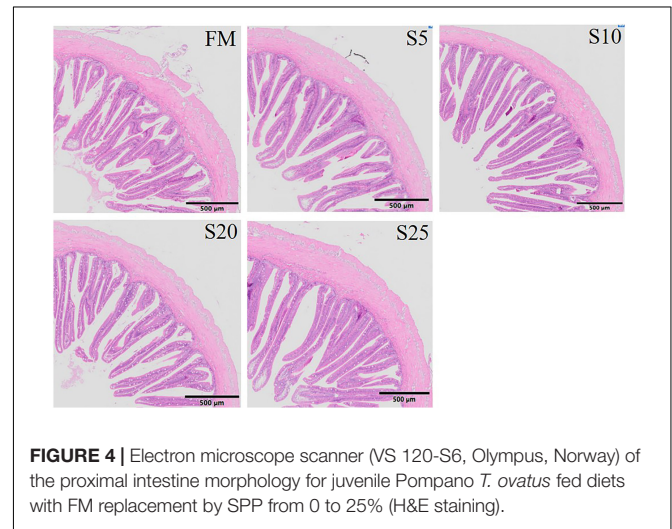
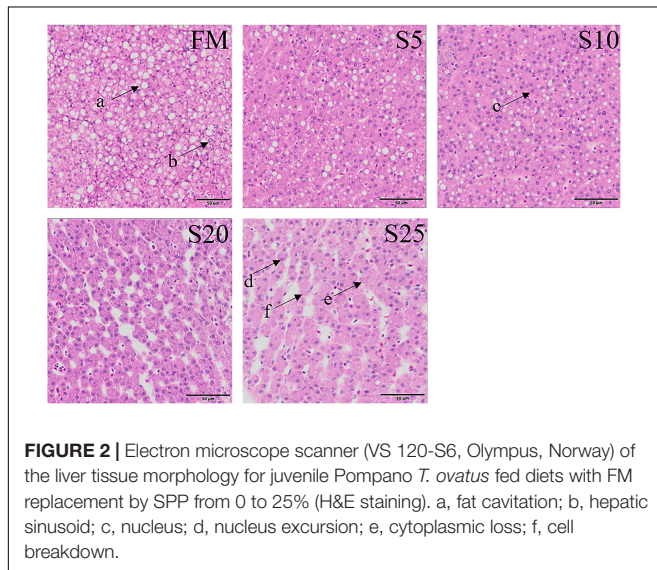
peptides that enter the bloodstream quickly and directly without degradation. Moreover, in *E. akaara*, reduction in HSI was associated with increased rates of small peptide and free amino acid transport and absorption and enhanced lipid metabolism (Zhao S. Y. et al., 2016). In the present study, a significant reduction in liver fat content was observed with oil red staining. Therefore, SPP may reduce liver fat accumulation and improve liver condition.

Animal blood is an important carrier of nutrients and metabolites, and elevated levels of TG and T-CHO in the blood can be detrimental to animal health (Kim et al., 2005). Increase or decrease in TG levels can be used as an indicator of fat metabolism and liver function (He et al., 2021). Nagasawa et al. (2003) reported that SPP significantly reduces triglyceride content and fatty acid synthase mRNA levels in adipose tissues, suggesting that soy isolate controlled gene expression in adipose tissues and effectively regulated adipocyte differentiation. In the present study, T-CHO and TG levels were significantly lower

TABLE 8 | Serum biochemical indices for juvenile *Trachinotus ovatus* fed the experimental diet.

Parameters	Experimental diets				
	FM	S5	S10	S20	S25
TP (g/L)	6.08 ± 1.80 ^a	28.64 ± 2.01 ^c	16.27 ± 1.74 ^b	12.37 ± 3.00 ^b	5.05 ± 1.60 ^a
TG (mmol/L)	2.02 ± 0.11 ^c	1.38 ± 0.14 ^b	1.24 ± 0.06 ^{ab}	1.09 ± 0.12 ^a	1.10 ± 0.10 ^{ab}
T-CHO (mmol/L)	23.78 ± 2.99 ^b	16.50 ± 0.31 ^a	16.35 ± 2.00 ^a	15.02 ± 1.78 ^a	12.65 ± 1.49 ^a
GLU (mmol/L)	4.36 ± 0.24	4.32 ± 0.31	3.95 ± 0.15	4.11 ± 0.34	3.78 ± 0.17

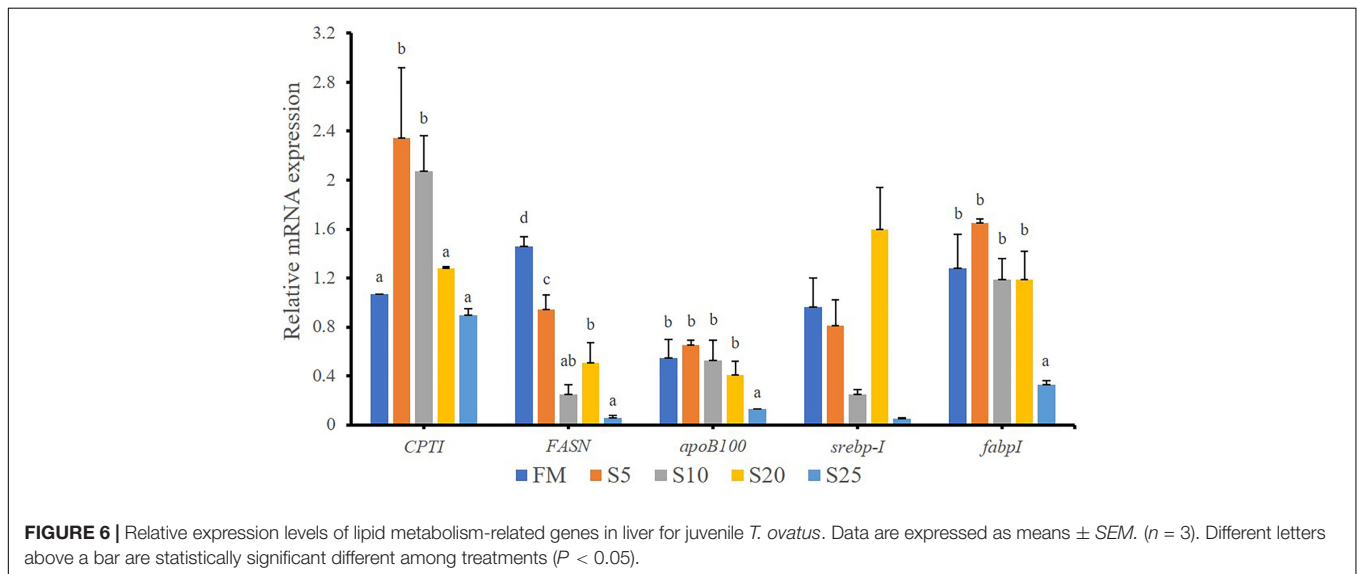
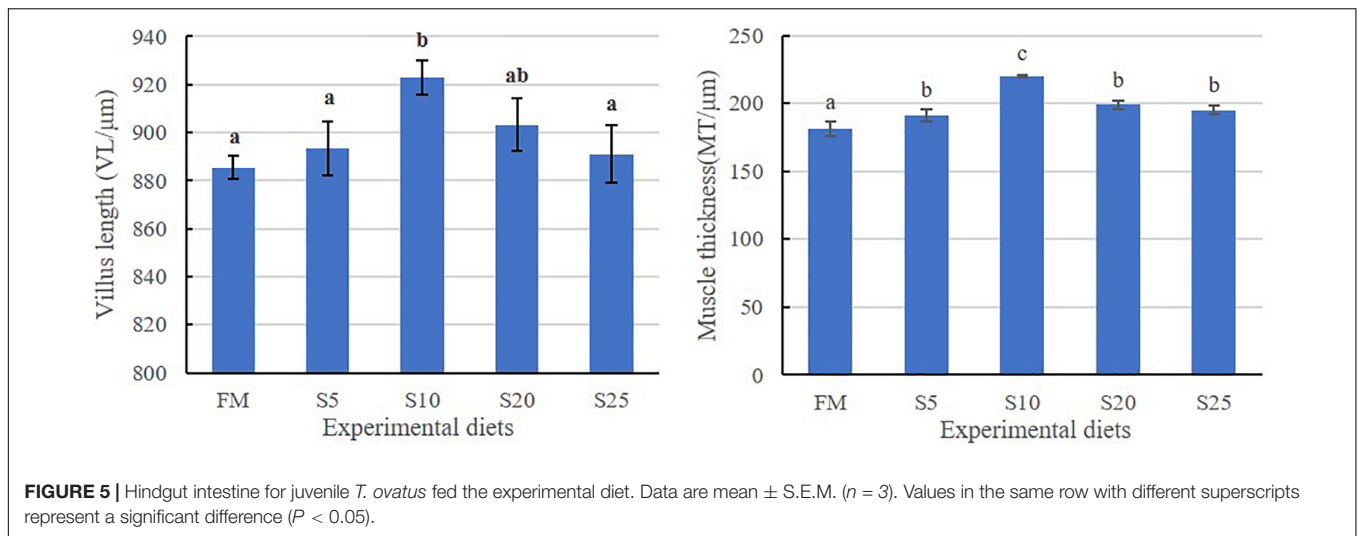
Data are mean ± S.E.M. (n = 3). Values in the same row with different superscripts represent significant difference ($P < 0.05$). TP, total protein; TG, triglyceride; T-CHO, total cholesterol; GLU, glucose.



than those in the control group (FM), indicating that SPP can effectively regulate TG and T-CHO metabolism. LPYPR (Leu-Pro-Tyr-Pro-Arg) and VK (Val-Lys) derived from glycine in soy are important components that lower cholesterol peptides and triglycerides, respectively (Inoue et al., 2011, 2015). SPP

are more beneficial to the balanced absorption of amino acids in fish than FM.

Fish fat metabolism mainly occurs in the liver, including synthesis, catabolism, and transport, and is regulated by a multitude of factors with synergistic activities (Hu, 2004; Weng et al., 2012). The *fasn* is the key enzyme for the *de novo* fatty acid biosynthesis, catalyzing the synthesis of malonyl-CoA and acetyl coenzyme A into long-chain saturated fatty acids (He et al., 2021). The *cpt1* is a key and rate-limiting enzyme that regulates the beta-oxidation of fatty acids, the main catabolic process in the body (Nilsson-Ehle et al., 1980). In the present study, the FM group had the highest liver *fasn* mRNA level, and *cpt1* expression in the S5 and S10 groups was significantly up-regulated. These findings were consistent with those of Nagasawa et al. (2003). In addition, fat deposition is related to fat transport rate. Increase in very-low-density lipoprotein (VLDL) secretion rate can reduce hepatic lipid deposition (Nagayoshi et al., 1995). Hussain et al. (2008) reported that *apob100* is an essential component in the assembly of VLDL particles. The *fabp1* can bind long-chain fatty acids for oxidation and storage through the cell membrane (Yan et al., 2015). In the present study, the S25 group had significantly lower in *apob100* and *fabp1* expression levels than that the other groups. The morphology of the liver cells in the FM, S5, and S10 groups was normal, and fat accumulation in the liver gradually decreased. These results were in line with those of the HSI and VSI analyses. With FM decreased, liver



cells were seriously damaged, the cells collapsed, and the nuclei disappeared. These effects were in line with the observation on *Oncorhynchus mykiss* (Feng et al., 2016), *Carassius auratus* (Shi et al., 2015), and *Lateolabrax japonicus* (Hu et al., 2013). We considered that excessive soy protein peptides can damage the liver, leading to dysfunction (Zhang et al., 2002). Therefore, our observations indicated that the replacement of 10% FM with SPP significantly promotes liver fat metabolism in grouper, reduces liver fat deposition, and improves liver condition.

The defense system can protect cells and cell membranes from oxidative damage and maintain normal physiological function (Bu et al., 2017). MDA is the oxidative end-product of free radicals acting on the peroxidation reaction of fat, and its content indirectly reflects the content of oxygen free radicals in the cells and the severity of free radical attacks (Cheng et al., 2017). In the present study, MDA content significantly decreased with increasing SPP supplementation. This result was in line with the results of Gyan et al. (2020). Deng et al. (2006)

reported that soybean peptides had strong DPPH (1,1-Diphenyl-2-picrylhydrazyl) radical-scavenging ability and most functional peptides with antioxidant properties were mainly concentrated in small peptides with molecular weights of lower than 5,000 Da. The molecular weights of SPP used in this experiment were mainly lower than 1,000 Da, providing a good molecular basis for antioxidant function. We suggest that SPP can inhibit free radicals and reduce the content of lipid peroxides during stress. GSH-Px is important antioxidant enzymes in fish (Cheng et al., 2017). GSH-Px catalyzes the reduction of glutathione and removes hydrogen peroxide and lipid peroxides produced during metabolism. In the present study, GSH-Px activity reached a maximum in the S10 and S20 groups significantly higher than that in the other groups. Similar results were *Cyprinus carpio* (Wang et al., 2014), *Ctenopharyngodon idella* (Zheng et al., 2012), and *P. fulvidraco* (Jing et al., 2021), indicating that SPP could increase antioxidant activity, and it is related to functional peptides with antioxidant properties in soy peptides

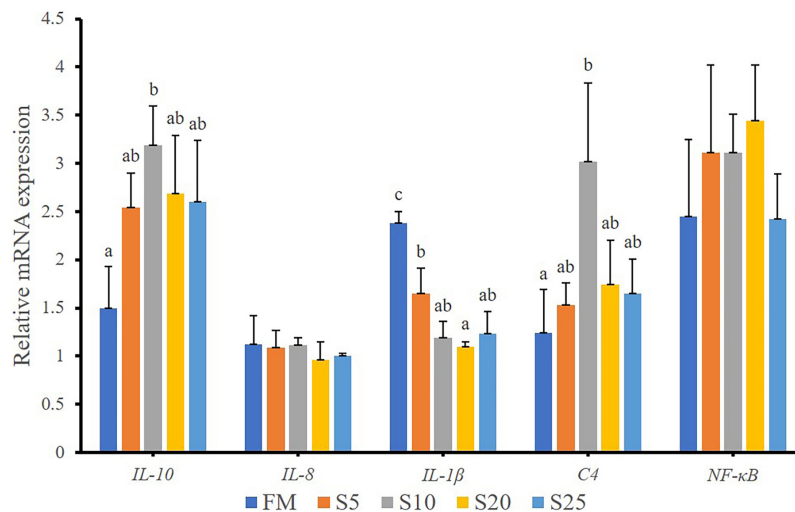


FIGURE 7 | Relative expression levels of immunity-related gene in intestine for juvenile *T. ovatus*. Data are expressed as means \pm SEM. Different letters above a bar are statistically significant different among treatments ($P < 0.05$).

(Deng et al., 2006). The above antioxidant indices indicate that soy protein peptides can increase the antioxidant activity for juvenile *T. ovatus*.

Active small peptides can participate in immune system regulation, can improve immunity, and exerts immunomodulatory effects, specifically stimulating macrophage phagocytosis and inhibiting lymphocyte proliferation (Kong et al., 2008). In serum, increases in LYZ, ACP, and AKP contribute to the immune system of aquatic organisms (Tseng et al., 2009). LYZ can hydrolyze mucopolysaccharides and catalyze the hydrolysis of glycosidic bonds in bacterial cell walls, thus causing bacterial cell walls to rupture (Xie et al., 2013). In the present study, serum LYZ activity significantly increased. The same effect was observed in *L. Vannamei*, indicating that the addition of small peptides significantly increases antimicrobial activity and lysozyme specific activity in sera (Lin et al., 2010). Based on the present results, the optimal SPP level that supported the most powerful immune function of juvenile *T. ovatus* was 11.82%, as predicted by the LYZ activity models. ACP and AKP catalyze phosphate monostearate hydrolysis and play important roles in the immune response of the body against pathogens (Liu et al., 2004). In the present study, serum AKP activity increased first and then decreased with as the amount of replacement SPP increased in *T. ovatus* diet. This result was in line with the results obtained in *Sardine* (Ben Khaled et al., 2012) and *P. fulvidraco* (Zhao Z. X. et al., 2016). In addition, soy peptides help eliminate anti-nutritional factors and act as immunomodulators, inducing defense genes involved in pathogenic attacks (Pearce et al., 2010).

As an important immune organ in the animal body, the intestine provides protection against external pathogens. As reported by Zhang et al. (2020), SPP can reduce inflammation and enhance immune function by regulating the expression of pro-inflammatory cytokines, such as *il-1β*, which in turn affects the expression and secretion levels of T cells. The *il-1β* plays an important role in inflammatory responses (Dinarello, 2000).

In the present study, the mRNA expression level of *il-1β* in the intestine was significantly down-regulated as SPP replacement level increased. Moreover, the anti-inflammatory factor *il-10* was significantly up-regulated in the S10 group. Complement is the main humoral component of the innate immune response and plays an essential role in the killing of microorganisms, phagocytosis, inflammatory response, immune complexes, and antibody production (Holland and Lambris, 2002). Fish can recognize and neutralize a variety of harmful microorganisms by specifically binding different forms of complement (Zarkadis et al., 2001). In the present study, the replacement of 10% FM with SPP significantly up-regulated gene expression of *c4*. The possible reasons are that small peptides are efficiently absorbed, and the amount of undigested protein in the intestine are reduced (Zheng et al., 2006). The exact mechanisms of the effects have not been explained and further studies are needed.

Among the internal organs of fish, the gut has the largest area of contact with the internal environment. The intestine is the main part of the fish that digests and absorbs nutrients (Zhou, 2012). The shape of the fold is positively correlated with nutrient absorption area (Wu et al., 2010). A decrease in the height of the intestinal fold means that the intestinal tract has a smaller absorptive area and absorb lower amounts of nutrients (Wu et al., 2014). The results of the present study showed that the replacement of FM with SPP can promote the growth of intestinal villi and muscle layer. Thus, the area of intestine absorption was enlarged. The present study observed a trend in line with the study of Jiang et al. (2010), indicating that increase in intestinal absorption area and muscle layer thickness can facilitate the absorption of SPP and improve feed utilization. In addition, the replacement of FM with a high proportion of SPP can inhibit villi growth. A high proportion of FM replacement reduces the protein metabolism of fish and damages the intestinal tissue structure (Wang et al., 2017). If the proportion of a plant-based protein used to replace FM is extremely high, the

intestinal tract of aquatic animals will be affected to some extent. This observation showed that under the present experimental conditions, the replacement of 10% FM with SPP could promote intestinal muscle thickening and wrinkled wall growth, thus improving intestinal absorption.

CONCLUSION

The replacement of FM with SPP can promote intestinal growth for nutrient absorption without affecting the growth performance of juvenile pompano *T. ovatus* under experimental conditions. In addition, SPP can reduce liver fat accumulation, improve liver condition, and enhance antioxidant capacity and immunity. Based on the second-order polynomial analysis model of LYZ activity, the optimal replacement SPP level for juvenile pompano *T. ovatus* is 11.82%.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation, to any qualified researcher.

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ETHICS STATEMENT

The animal study was reviewed and approved by the Guangdong Ocean University Institutional Animal Care and Use Committee.

AUTHOR CONTRIBUTIONS

HL and QY designed the experiments. HL carried out the experiments and drafted the manuscript. BT, MZ, ML, and SC were accountable for some aspects (such as ingredients and sites) of the work in ensuring that experiments can be carried out properly. QY and GR reviewed and revised the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: MZ and ML were employed by the company Yisheng (Yangjiang) Biotechnology Company Limited.

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