### **REGULAR ARTICLES**



# Exploration of marine red seaweed as a dietary fish meal replacement and its potentiality on growth, hematological, biochemical, and enzyme activity in freshwater fish *Labeo rohita*

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#### **Abstract**

The present study investigated the dietary fishmeal replacement by marine red seaweed (Halymenia dilatata) meal (RSM) on growth performance, feed utilization, chemical body composition, hematological constituents, digestive, antioxidant, and metabolic enzymes in freshwater fish Labeo rohita (Rohu) fingerlings. The fish were fed with RSM-free control diet (RSM0) and four experimental diets, which replaced fish meal (FM) with varying levels of RSM (25%, 50%, 75%, and 100%, represented as RSM25, RSM50, RSM75, and RSM100 respectively). After a 60-day feeding trial, the survival rate (SR), growth performance (length gain, weight gain, and specific growth rate), protein efficiency ratio, chemical body composition (protein, lipid, and ash), and digestive enzymes (amylase and protease) were significantly increased (P < 0.05) in the fish fed with RSM50 diet containing 39% protein level. The sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) revealed a higher staining intensity of muscle proteins in fish fed with the RSM50 diet. However, the hematological constituents (hemoglobin, hematocrit, red blood cell, white blood cell, mean corpuscular volume, mean corpuscular hemoglobin, and mean corpuscular hemoglobin concentration), antioxidant enzyme - superoxide dismutase, and metabolic enzymes (glutamic oxaloacetic transaminase and glutamic pyruvic transaminase) were not significantly altered in RSM50 diet when compared to control. In contrast, hematological constituents were decreased (P < 0.05), and antioxidant and metabolic enzymes were increased in rolu fed with RSM75 and RSM100 (P < 0.05). Furthermore, these findings suggest that RSM might be adopted at a pace of 37% (estimated polynomial second-order regression) and is found to be beneficial for freshwater fish L. rohita diets that enhance growth and immune responses. The current study recommended substituting (50%) of marine red seaweed (Halymenia dilatata) for fish meal significantly improves the growth performance, chemical body composition, and digestive enzymes of L. rohita and this could be a valuable natural replacement for fishmeal to reduce the production cost of aquatic feed.

Keywords Fish nutrition · Weight gain · Feed efficiency · Polynomial regression · Protein profile

### Introduction

Aquaculture contributes half of the total global biodiversity, and India shares 7.58% of global production next to China (Boyd et al. 2020). According to the

National Fisheries Development Board (NFDB), aquaculture is one of the food-producing industries in India that is growing rapidly. Fish are an important part of the aquatic fauna and have a significant potential for long-term economic growth (Pounds et al. 2022; Iitembu

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et al. 2022; Muñiz et al. 2022). Increased aquaculture production is required to meet the growing population's animal protein requirements in order to alleviate poverty and malnutrition and also ensure a sustainable livelihood (Madsen et al. 2022; Akegbejo-Samsons, 2022). Globally, the production of freshwater fish Labeo rohita (rohu) was over 1.67 million tonnes in 2014 (FAO 2018). In Asia, the Indian major carp L. rohita contributes 35% of total aquaculture production, due to its economic significance, accelerated growth yield, controlled reproduction, and customer demand (Harikrishnan et al. 2021). However, recently, there is a demand for fish meal (FM), which is used as a protein source in commercial fish feeds (Naylor et al. 2009; FAO 2014; Maiolo et al. 2020). Therefore, research has already commenced finding alternative ingredients to replace FM in formulated feeds (Samaddar et al. 2015; Panase et al. 2018; Hassaan et al. 2019; Oliva-Teles et al. 2022).

Nowadays, seaweeds have been closely associated with human, animal, and fish life and have been used as a source of food, feed, and medicine (Demirel et al. 2009; Jimenez-Escrig et al. 2011; Rebours et al. 2014; García-Vaquero and Hayes 2016; Terriente-Palacios and Castellari, 2022). The edible seaweeds contain moderate concentrations of proteins, lipids, high levels of fiber, vitamins, minerals, omega-3 fatty acids, and antioxidant compounds like alkaloids, steroids, and terpenoids (Murata and Nakazoe 2001; Holdt and Kraan 2011; Rajapakse and Kim 2011; Afonso et al. 2021; Morais et al. 2020). The red seaweed Porphyra spp. and Undaria spp. have a higher protein when compared with brown (Laminaria spp. and Undaria spp.) and green (Ulva sp.) seaweeds (Murata and Nakazoe 2001; Marsham et al. 2007; Peñalver et al. 2020). In order to assess the potential of seaweed used as a nutritional aid, research is necessary to perform aquaculture nutrition—related studies, which are infrequent in fish (García-Ortega et al. 2016; Roleda and Hurd 2019). Previous reports suggest that the dietary inclusion of seaweed as the replacement for fishmeal, Porphyra purpurea (red seaweed) in the diet of Chelon labrosus (thicklip grey mullet) (Davies et al. 1997); Oreochromis niloticus (Nile tilapia) were fed with *Ulva rigida* (green seaweed) (Azaza et al. 2008); Incorporation of Porphyra dioica (red seaweed) to Oncorhynchus mykiss (Rainbow trout) (Soler-Vila et al. 2009); and *Pagrus major* (Red seabream) were fed with Ascophyllum nodosum (brown seaweed) (Nakagawa et al. 1997); whereas, Mustafa et al. (1995) had shown incorporation of all three seaweeds (Ascophyllum nodosum, Porphyra yezoensis, and Ulva pertusa) in the diet of Pagrus major (Red seabream). With these evidences, seaweeds are highly recommended to be included in the fish feed that enhance the growth, physiological activity, and disease resistance of cultured fish.

Halymenia, a marine red seaweed (Family: Halymeniaceae), is found in various parts of the world (Guiry and Guiry 2020). Among them, the species *Halymenia* dilatata (Jainab et al. 2019), Halymenia maculata (Fantonalgo. 2018), Halymenia durvillaei (Fenoradosoa et al. 2009; Boominathan et al. 2022), Halymenia formosa (Hurtado et al. 2020), Halymenia floresia (Malairaj et al. 2016; das Neves Amorim et al. 2011), and Halymenia palmata (Deepak et al. 2019) are better sources of proteins, fatty acids, minerals (I, Cu, Ni, Cd, and Zn), pigments (carotene, chlorophyll a, chlorophyll d, lutein, phycocyanin, r-phycoerythrin, and zeaxanthin), and polysaccharides (sugars, carrageenan, floridean starch, funoran, furcellarin, and galactan) (Aldon 1998; Shah et al. 2022). They also act as antimicrobial, antioxidant, food additives, nematicide, pesticide, antiandrogenic, and flavors involving in anti-inflammatory activities (Balasubramanian et al. 2021; Fredrick Raja. 2022; Angulo et al. 2020). The present research is aimed to investigate fish meal replacement with a marine red seaweed Halymenia dilatata in the diet of Labeo rohita. The growth performance, digestive enzymes, chemical body composition, hematological parameters, antioxidant and metabolic enzymes were evaluated in this study. In addition, polynomial second-order (quadratic) regression model was adopted to estimate the optimal replacement of fishmeal with H. dilatata.

### **Materials and methods**

# Collection of marine red seaweed and feed ingredients

The marine red seaweed, *Halymenia dilatata*, was collected from Mandapam coastal area in the Gulf of Mannar (Lat. 09° 17.417′N; Long. 079° 08.558′E), Ramanathapuram district, Tamil Nadu, India. The seaweed was rinsed in pure water, allowed to air-dry then milled into a fine powder and kept in sterile, airtight containers. Feed ingredients such as fish meal (FM), groundnut oil cake (GOC), soybean meal (SBM), wheat bran (WB), rice bran (RB), and tapioca flour (TF) were purchased from the local market, and these ingredients were also powdered. The vitamins and minerals premixture was purchased from the medical shop at Tiruchirappalli, Tamil Nadu, India. All the feed ingredients were analyzed for proximate composition by the method of Castell and Tiews (1980), as given in AOAC (1995).



### **Diet preparation**

Diet formulation was done to fulfill the basic requirement of protein (30–40%) and lipid (5–16%) for L. rohita fingerlings (Satpathy et al. 2003). The feed ingredients, protein (FM, GOC, and SBM), and carbohydrate (WB, RB, and TF) sources were combined, steam-cooked for 15 min at 60 °C, and then dried at 28 °C. The vitamin and mineral premixture was added to the experimental diets along with sunflower oil (Table 1). RSM-free control diet (RSM0) and four experimental diets, including varying levels of RSM (RSM25 = 25%, RSM50 = 50%, RSM75 = 75%, and RSM100 = 100%), were formulated. The dough was prepared with 10% hot water and pelletized separately. An air blower in a drying cabinet maintained at 38 °C was used to air-dry the experimental diets until the moisture content was around 10%. Followed by air-drying, the diets were chopped up into pellets sized (< 1 mm) and kept at - 20 °C until use. Proximate composition analyses of these diets were also performed, according to AOAC (1995). The results of the proximate composition of FM replaced with RSM included diets did not show much difference comparing to basic nutritional requirements of *L. rohita* fingerlings (Table 1).

**Table 1** Formulation and proximate composition (g kg<sup>-1</sup>) of fish meal replacement by red seaweed (H. dilatata) meal incorporated diets

Ingredients (g kg <sup>-1</sup> )	Experimental diets								
	RSM0	RSM25	RSM50	RSM75	RSM100				
Fish meal	200	150	100	50	0				
Red seaweed meal	0	50	100	150	200				
Groundnut oil cake	200	200	200	200	200				
Soybean meal	200	200	200	200	200				
Rice bran	120	120	120	120	120				
Wheat bran	120	120	120	120	120				
Tapioca flour	120	120	120	120	120				
Sunflower oil	20	20	20	20	20				
Vitamin and mineral mix <sup>a</sup>	20	20	20	20	20				
Proximate composition	n (% of a	lry matter	basis)						
Moisture	8.58	8.53	8.49	8.35	8.41				
Crude protein	41.67	40.32	38.55	37.09	36.16				
Crude fiber	3.58	3.34	3.27	3.20	3.08				
Ether extract	6.93	6.75	6.31	5.96	5.74				
Ash	9.56	9.37	9.45	9.18	9.29				
NFE	29.68	31.69	33.93	36.22	37.32				
Gross energy (kJ $g^{-1}$ )	43.31	43.80	43.52	43.78	43.61				

NFE, nitrogen-free extract

<sup>a</sup>Becosules capsules: thiamine mononitrate (USP), 50 mg; riboflavin (USP), 25 mg; pyridoxine HCI (USP), 10 mg; cyanocobalamin (USP), 15 mcg; niacinamide (USP), 100 mg; calcium pantothenate (USP), 25 mg; folic acid (USP), 1 mg; ascorbic acid (BP), 150 mg

## Collection and acclimatization of experimental fish

The freshwater fish, L. rohita fingerlings, were collected from Nathan fish farm in Thanjavur, Tamil Nadu, India. They were transported to the laboratory in well-oxygenated polythene bags, and they were stored in cement tanks with the following dimensions: height, 1.04 m; radius, 0.96 m, and volume,  $\approx 3000$  L. They were allowed to acclimatize to the laboratory conditions for 2 weeks. During acclimatization, the fish were fed with commercial feed ad libitum. To keep the fish in a healthy condition. the water was frequently changed (by 50%) each day, and the tanks were constantly aerated. The physicochemical properties were sustained as temperature,  $29.14 \pm 0.51$  °C; pH,  $7.20 \pm 0.30$ ; total dissolved solids,  $0.102 \pm 0.10$  g L<sup>-1</sup>; and dissolved oxygen,  $6.15 \pm 0.40$  mg L<sup>-1</sup>, throughout the experimental period (APHA 1998).

# Feeding experiment

While maintaining and handling fish throughout the experiment, all applicable criteria for the care and use of animals provided by the Committee for the Purpose of Control and Supervision of Experiments on Animals (CPCSEA), Government of India, have been carefully followed. Twenty L. rohita fingerlings  $(3.35 \pm 0.06 \text{ g}; 6.47 \pm 0.06 \text{ cm})$  $(\text{mean} \pm S.D)$  for each diet in triplicates were maintained in plastic troughs (height, 27 cm; radius, 21.5 cm) with 20 L of water. The experimental groups were fed with RSM0, RSM25, RSM50, RSM75, and RSM100 experimental diets. The feed quantity was adjusted every 10 days based on the 10% bodyweight of fingerlings. Throughout the 60-day experimental period, the daily feed was split in half and fed twice a day (at 09:00 and 18:00); moderate aeration was continually administered to maintain the ideal oxygen level (>  $6.00 \text{ mg L}^{-1}$ ).

### **Growth parameters**

After the feeding trial, survival rate (SR), growth parameters [final length (FL), length gain (LG), final weight (FW), weight gain (WG), specific growth rate (SGR)], and feed utilization [feed conversion ratio (FCR) and protein efficiency ratio (PER)] were determined according to García-Ortega et al. (2016) as follows:

Survival rate (%)

 $= 100 \times (\text{final number of fish / initial number of fish})$ 

Length gain (cm) = FL - IL



where FL is the final length of individual fish (cm) and IL is the initial length of individual fish (cm)

Weight gain 
$$(g) = FW - IW$$

where FW is the final weight of individual fish (g) and IW is the initial weight of individual fish (g)

Specific growth rate  $(\% \text{ day } -^1) = 100 \times (log_{10} \text{ FW } - log_{10} \text{IW}) / \text{ days}$ 

Feed conversion ratio (g)

= individual feed intake (g)/ individual weight gain (g)

Protein efficiency ratio (g) = (FW - IW)/ individual feed intake  $\times$  crude protein g  $-^1$  of diet

# Analysis of the proximate composition

Fish muscle tissues were analyzed in triplicates for proximate composition (moisture, protein, lipid, and ash) using the following standard methods. Protein was determined by the method of Lowry (1951) using trichloroacetic acid (TCA) precipitation. Lipid was assessed by the method of Barnes and Blackstock (1973) using chloroform—methanol extraction (Folch et al. 1957). The content of moisture and ash was determined, according to AOAC (1995).

# Analysis of protein profile/sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE)

Fish muscle tissues were homogenized in phosphate buffer (pH 7.4) solution and centrifuged at 14,000 rpm at 4 °C for 15 min. The soluble protein content was determined according to the method of Bradford (1976). For the determination of the molecular weight of proteins, 10% SDS-PAGE (Laemmli 1970) was performed in a vertical gel electrophoresis system (Bio-Rad, Hercules, CA, USA). Thermo Fisher Scientific (Massachusetts, USA) 10–180-kDa protein molecular markers were used as a standard marker. The gel was stained by Coomassie blue (0.025%). Molecular weights of the polypeptide bands were determined by comparing the relative mobility of protein bands to the standard protein marker.

## **Analysis of hematological parameters**

The cardiac puncture was done, and the blood samples were collected in a heparinized (Beparine<sup>R</sup> heparin sodium, IP 1000 IU mL<sup>-1</sup>, an anticoagulant) medical grade disposable readymade (26-gauge needle) syringe from the experimental fish after 60 days experimental trial. Then, the blood was transferred into heparinized plastic vials. The blood was analyzed for hematological parameters [hemoglobin (Hb), hematocrit (Hct), red blood cells (RBC), white blood cells

(WBC), mean cell volume (MCV), mean cell hemoglobin (MCH), and mean cell hemoglobin concentration (MCHC)] using Mindray Auto Hematology Analyzer BC 2800 (Mindray Bio-Medical Electronics Co., Ltd, Shenzhen, China).

### Assay of digestive enzymes

The digestive tract of fish was carefully dissected out and homogenized [1:10 (w/v %) wet mass: volume ice-cold double distilled water] by a glass homogenizer and centrifuged at 9300 g for 20 min at 4  $^{\circ}$ C. The suspension of the supernatant was used as an enzyme source.

Amylase activity was determined according to the starch-hydrolysis method of Bernfeld (1955). Maltose was used as standard, and 1 unit of amylase activity (U) was determined as the amount of enzyme to produce 1 mg of maltose  $\min^{-1} \operatorname{mg}^{-1}$  protein at 25 °C.

Protease activity was assayed by the casein-hydrolysis method of Furne et al. (2005), L-tyrosine was used as standard, and 1 unit of protease enzyme activity (U) was determined as the amount of enzyme desired to liberate 1  $\mu$ g of tyrosine min<sup>-1</sup> mg<sup>-1</sup> protein at 37 °C.

## Assay of antioxidant enzyme (SOD) activity

Gills, liver, and muscle tissues of fish were individually homogenized (10% w/v) in ice-cold 50-mM Tris buffer (pH 7.4) and centrifuged at 9300 g for 20 min at 4 °C.

The supernatant was used to assay the SOD enzyme activity. Soluble tissue protein concentration was determined by the method of Lowry (1951). The SOD activity was measured using pyrogallol (10 mM) autoxidation in Tris buffer (50 mM, pH 7.0), as described by Marklund and Marklund (1974). The activity of SOD was expressed in U mg<sup>-1</sup> protein.

# Assay of metabolic enzymes (GOT) and (GPT) activities

Fish gills, liver, and muscle tissues were individually homogenized in a solution of 0.25-M sucrose and centrifuged at 9300 g for 20 min at 4 °C. The supernatant was obtained and used as a source of enzymes. The glutamic oxaloacetic transaminase (GOT) and glutamic pyruvic transaminase (GPT) were analyzed according to the method of Reitman and Frankel (1957) using the kit (Medox Biotech India PVT, Ltd).

For GOT analysis, the substrate solution, L-aspartic acid (500 μL; pH 7.4), was added to 100 μL of sample and incubated at 37 °C for 1 h. Further, 500 μL of 2,4-dinitrophenyl hydrazine was added and allowed to stand for 20 min at room temperature. The final mixture was added to 3 mL of freshly prepared 4-N sodium hydroxide solution. The colored solution was read at 505 nm using Synergy<sup>TM</sup> HTX Multi-Mode Microplate Reader (Biotek, Winooski, VT,



USA) within 15 min. Sodium pyruvate (160 U  $L^{-1}$ ) was used as a calibrator. The activity of GOT was expressed as U  $L^{-1}$ .

For GPT analysis, buffered L-alanine and 2-oxoglutarate substrate (500  $\mu L$ ; pH 7.4) was added to 100  $\mu L$  of sample and incubated at 37 °C for 20 min. Then, 500  $\mu L$  of 2,4-dinitrophenyl hydrazine was added and allowed to stand at 28 °C for 30 min followed by the addition of 3 mL of freshly prepared 4-N sodium hydroxide solution. The colored solution was read at 505 nm using a Synergy TM HTX Multi-Mode Microplate Reader (Biotek) within 15 min. Sodium pyruvate (170 U  $L^{-1}$ ) was used as a calibrator. The activity of GPT was expressed as U  $L^{-1}$ .

### **Data analysis**

The polynomial second-order (quadratic) regression analysis was used to evaluate the effect of the increasing levels of dietary RSM on growth performance, proximate composition, digestive enzyme activity, hematology, and immune response. One-way analysis of variance (ANOVA) was used to differentiate among the diets, and significant (P < 0.05) differences among diet means were ranked by Duncan's multiple range test (DMRT). The data were expressed as mean  $\pm$  SE (standard error of the mean).

#### Results

### Survival and growth parameters

The results of SR, growth performance (IL, FL, IW, FW, WG, SGR), and feed utilization (FCR and PER) were shown in Table 2. Significantly increased values (P < 0.05)

in SR, FW, WG, SGR, and PER values were observed in *L. rohita* fed with the diets containing FM replacement with RSM50 followed by RSM25 as compared with RSM0. Meanwhile, the SR, FW, WG, SGR, and PER for both diet RSM75 and RSM100 were significantly decreased (P < 0.05) in comparison to those of other diet groups. Additionally, the dietary FM replacement by RSM35, RSM34, and RSM37 level promoted the most significant amount of WG, SGR, and PER, as estimated by the polynomial regression with strong positive correlation ( $y = -0.000x^2 + 0.048x + 5.236$ ;  $y = -7E - 05x^2 + 0.004x + 0.671$ ;  $y = 0.000x^2 + 0.014x + 1.722$ , respectively) (Table 2; Fig. 1a, b, and d).

The FCR in *L. rohita* was insignificantly decreased (P > 0.05) in groups fed RSM incorporated diets, especially at the level of RSM50 followed by RSM25 as compared with the control group. The diet RSM50 showed the lowest FCR values among the experimental diets (Table 2). Additionally, the polynomial regression of dietary RSM level and FCR was the fit regression modeling with high goodness  $(y = 0.000x^2 - 0.008x + 1.392)$  (Table 2; Fig. 1c). The significant value of FCR was estimated at the RSM25 diet according to the regression curve.

### **Chemical body composition**

The protein, lipid, and ash contents were significantly elevated (P < 0.05) in *L. rohita* fed on dietary FM replacement by RSM50 and RSM25 diets when compared to other experimental diets, whereas in the moisture content, no significant (P > 0.05) change was detected among the diet groups, as shown in Table 3. However, the FM replacement by the RSM analysis showed that the dietary RSM level

Table 2 Survival, growth performance, and feed utilization of L. rohita fingerlings fed with experimental diets for 60 days

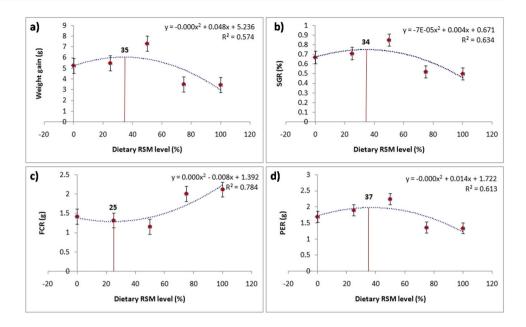
Parameters	Experimental of	liets			PSEM	Regression <sup>2</sup>		ANOVA	
	RSM0	RSM25	RSM50	RSM75	RSM100		Equation	$R^2$	P value
SR (%)	$80.00 \pm 5.00^{ab}$	$80.00 \pm 5.00^{ab}$	$86.66 \pm 2.88^a$	$71.66 \pm 5.77^{bc}$	$66.66 \pm 2.88^{c}$	5.12	$79 + 0.221x - 0.003x^2$	0.787	0.002
IL (cm)	$6.57 \pm 0.23^{a}$	$6.41 \pm 0.17^{a}$	$6.49 \pm 0.24^{a}$	$6.54 \pm 0.25^{a}$	$6.52 \pm 0.22^{a}$	0.07			
FL (cm)	$9.40 \pm 0.45^{bc}$	$9.66 \pm 0.50^{b}$	$10.90 \pm 0.41^{a}$	$8.77 \pm 0.35$ cd	$8.35 \pm 0.46^{d}$	0.81	$9.338 + 0.042x - 0.000x^2$	0.653	0.000
LG (cm)	$2.83 \pm 0.33^{bc}$	$3.25 \pm 0.52^{b}$	$4.41 \pm 0.35^{a}$	$2.23 \pm 0.22^{\text{ cd}}$	$1.83 \pm 0.57^{d}$	0.81	$2.802 + 0.044x - 0.000x^2$	0.670	0.000
IW (g)	$3.41 \pm 0.21^{a}$	$3.31 \pm 0.15^{a}$	$3.29 \pm 0.20^{a}$	$3.38 \pm 0.24^{a}$	$3.42 \pm 0.12^{a}$	0.03			
FW (g)	$8.67 \pm 0.72^{b}$	$8.80 \pm 0.85^{b}$	$10.60 \pm 0.78^{a}$	$6.90 \pm 0.73^{\circ}$	$6.87 \pm 0.82^{c}$	1.45	$8.636 + 0.044x - 0.000x^2$	0.563	0.001
WG (g)	$5.26 \pm 0.76^{b}$	$5.49 \pm 0.70^{b}$	$7.31 \pm 0.88^{a}$	$3.52 \pm 0.64^{\circ}$	$3.45 \pm 0.64^{c}$	1.47	$5.236 + 0.048x - 0.000x^2$	0.574	0.000
SGR (%)	$0.67 \pm 0.08^{b}$	$0.71 \pm 0.04^{b}$	$0.85 \pm 0.08^{a}$	$0.52 \pm 0.07^{c}$	$0.50 \pm 0.08^{c}$	0.12	$0.671 + 0.004x - E - 05x^2$	0.634	0.001
FCR (g)	$1.42 \pm 0.16^{b}$	$1.32 \pm 0.18^{b}$	$1.16 \pm 0.14^{b}$	$2.01 \pm 0.34^{a}$	$2.12 \pm 0.41^{a}$	0.28	$1.392 - 0.008x + 0.000x^2$	0.784	0.004
PER (g)	$1.70 \pm 0.20^{\rm bc}$	$1.90 \pm 0.27^{\rm ab}$	$2.25 \pm 0.29^{a}$	$1.36 \pm 0.21^{c}$	$1.34 \pm 0.26^{c}$	0.33	$1.722 + 0.014x - 0.000x^2$	0.613	0.006

The data were expressed as mean  $\pm$  standard error of the mean, n=3 triplicates per treatments

Different superscript letters within a row indicates statistically significant (P<0.05) differences as evaluated by Duncan multiple range test PSEM, pooled standard error of the mean; RSM, red seaweed meal; SR, survival rate; FL, final length; LG, length gain; FW, final weight; WG, weight gain; SGR, specific growth rate; FCR, feed conversion ratio; PER, protein efficiency ratio



Fig. 1 Polynomial second-order (quadric) regression of weight gain (a), specific growth rate (SGR) (b), feed conversion ratio (FCR) (c), and protein efficiency ratio (PER) (d) in *L. rohita* fingerlings fed increasing inclusions of RSM for 60 days



fits with protein and lipid are polynomial regression with a strong positive correlation ( $R^2 = 0.679$  and 0.544, respectively) (Table 3).

### **Protein profile**

Polypeptide bands of molecular weights between 10 and 180 kDa were resolved in the muscle tissue of *L. rohita* fingerlings (Fig. 2). Seven Coomassie blue–stained protein bands (37, 30, 23, 17, 15, 13, and 10 kDa) were observed in FM replacement with RSM diets fed.

*L. rohita*, when compared with the standard molecular marker. Various polypeptide bands resolved in RSM25, RSM50, and RSM75 stained more intensely when compared with RSM0 and RSM100 diets. Mainly, 37, 23, 17, 15, 13, and 10 kDa polypeptide bands in diet RSM50-fed *L. rohita* were stained more intensely among the experimental groups.

# Hematology

The highest level of hematological indices (Hb, Hct, RBC, WBC, MCV, MCH, and MCHC) was recorded in *L. rohita* fed on RSM50, RSM25, and RSM0, and the lowest level was recorded in *L. rohita* fed on diet containing RSM75 followed by RSM100 diet (Table 4). The hematological indices of *L. rohita* were significantly altered by RSM75 and RSM100 diets (P > 0.05), as demonstrated in Table 4. In addition, the polynomial regression of high-goodness dietary substitution of FM with RSM levels and Hb, Hct, RBC, WBC, MCV, MCH, and MCHC was an effective regression model ( $R^2 = 0.734$ , 0.734, 0.709, 0.946, 0.887, 0.827, and 0.777 respectively; Table 4).

### **Digestive enzymes**

The amylase and protease activities were significantly enhanced (P < 0.05) in *L. rohita* fed with dietary FM replacement by

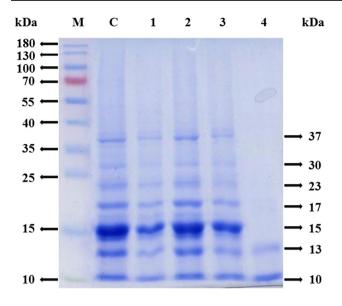
Table 3 Chemical body composition of L. rohita fed with experimental diets for 60 days

Parameters	Experimental diets						Regression		ANOVA	
	RSM0	RSM25	RSM50	RSM75	RSM100		Equation	$R^2$	P value	
Moisture (%)	78.79 ± 1.24 <sup>a</sup>	78.90 ± 1.12 <sup>a</sup>	$77.24 \pm 1.40^{a}$	79.36±0.98 <sup>a</sup>	81.94 ± 1.63 <sup>a</sup>	3.32	$79.56 - 0.186x + 0.002x^2$	0.572	0.934	
Protein (mg g <sup>-1</sup> wet wt.)	$125.17 \pm 8.07^{bc}$	$129.60 \pm 8.84^{ab}$	$139.18 \pm 9.37^{a}$	$117.55 \pm 2.92^{bc}$	$114.53 \pm 5.02^{\circ}$	7.88	$125.2 + 0.393x - 0.005x^2$	0.679	0.013	
Lipid (mg g <sup>-1</sup> wet wt.)	$56.86 \pm 4.45^{b}$	$58.43 \pm 2.96^{ab}$	$66.27 \pm 4.89^a$	$54.90 \pm 5.30^{b}$	$53.72 \pm 4.75^{b}$	4.72	$56.46 + 0.243x - 0.002x^2$	0.544	0.047	
Ash (%)	$2.01 \pm 0.06^{\rm bc}$	$2.08 \pm 0.06^{ab}$	$2.14 \pm 0.07^{a}$	$1.99 \pm 0.08^{bc}$	$1.94 \pm 0.04^{c}$	0.04	$2.013 + 0.004x - 5E - 05x^2$	0.800	0.023	

The data were expressed as mean  $\pm$  standard error of the mean, n=3 triplicates per treatments

Different superscript letters within a row indicate statistically significant (P<0.05) differences as evaluated by Duncan multiple range test PSEM, pooled standard error of the mean; RSM, red seaweed meal





**Fig. 2** Protein profile of the muscle tissue of *L. rohita* fingerlings fed with experimental diets for 60 days. Lane C, RSM0 diet–fed *L. rohita*; lane 1, RSM25 diet–fed *L. rohita*; lane 2, RSM50 diet–fed *L. rohita*; lane 3, RSM75 diet–fed *L. rohita*; lane 4, RSM100 diet–fed *L. rohita*; lane M, marker protein

RSM50 and RSM25 when compared to the other experimental diets (Table 5). Therefore, the polynomial regression of dietary RSM protease and amylase was the best regression model with high goodness ( $R^2$ =0.298 and 0.789) (Table 5).

# Antioxidant (SOD) and metabolic enzyme (GOT and GPT) activities

Insignificant values (P > 0.05) were observed in the SOD, GOT, and GPT enzyme levels in the gills, liver, and muscle of *L. rohita* fed on FM replacement by RSM50 and RSM25 diets when compared with that in the RSM0, RSM75, and RSM100 (Table 6). The antioxidant and metabolic enzymes

were increased significantly (P<0.05) in FM replaced with RSM75 and RSM100 (Table 6). Moreover, the polynomial regression of high-quality dietary replacement of FM with RSM levels and SOD, GOT, and GPT was an effective regression model (gills,  $R^2$ =0.943, 0.977, and 0.995; liver,  $R^2$ =0.957, 0.982, and 0.997; muscle:  $R^2$ =0.958, 0.938, and 0.923) (Table 6).

### Discussion

Marine seaweeds are increasingly used in biological applications, particularly in the pharmaceutical and feed formulation industries, because they contain a diverse range of bioactive and nutritional compounds and primary and secondary metabolites with potential therapeutic activity such as antimicrobial, anticancer, antiviral, anti-inflammatory, and immunomodulatory (Shah et al. 2022; Lomartire et al. 2021; Carpena et al. 2022; Ashkenazi et al. 2022). The marine seaweed-derived polysaccharides can be used for animal nutrition and improved animal intestinal integrity and enhanced immune responses (Leandro et al. 2019; Silva-Brito et al. 2022). Recently, many novel ingredients (insect meal, macroand microalgae, yeast) have been widely used as an alternative and essential protein diet for fish growth (Aragão et al. 2022). Pradhan et al. (2020) stated that plant-based alternatives to fishmeal supplementation increase the plankton population and also enhance fish growth. In this present study, the marine red seaweed (Halymenia dilatata) meal has been used as an alternative source for fishmeal in terms of growth performance, hematological, antioxidant, and digestive enzyme activity in the freshwater fish Labeo rohita.

The SR, LG, WG, SGR, and PER were significantly increased in *L. rohita* fed diets containing FM replacement with RSM50 diet (39% protein level) followed by RSM25 as compared with RSM0. The RSM properties like protein levels, vitamin and mineral composition, pigments, and

Table 4 Hematological constituents of L. rohita fed with experimental diets for 60 days

Parameters	Experimental diets						Regression		ANOVA
	RSM0	RSM25	RSM50	RSM75	RSM100		Equation	$R^2$	P value
Hb $(g dL^{-1})$	8.23 ± 0.61 <sup>a</sup>	$8.58 \pm 0.66^{a}$	8.95 ± 0.59 <sup>a</sup>	5.02 ± 0.37 <sup>b</sup>	5.17 ± 0.46 <sup>b</sup>	1.40	$8.454 + 0.015x - 0.000x^2$	0.734	0.000
Hct (%)	$31.07 \pm 1.24^{a}$	$31.92 \pm 1.02^{a}$	$32.66 \pm 1.05^a$	$20.11 \pm 1.09^{b}$	$21.02 \pm 1.02^{b}$	4.53	$31.85 + 0.022x - 0.001x^2$	0.734	0.000
RBC ( $\times 10^6  \mu L^{-1}$ )	$3.62 \pm 0.19^a$	$3.69 \pm 0.15^{a}$	$3.77 \pm 0.14^{a}$	$2.45 \pm 0.12^{\rm b}$	$2.61\pm0.08^{\mathrm{b}}$	0.48	$3.705 + 0.009x - 0.000x^2$	0.709	0.000
WBC ( $\times 10^{3} \mu L^{-1}$ )	$14.70 \pm 1.67^{a}$	$15.30 \pm 1.28^{\rm a}$	$15.66 \pm 1.83^a$	$9.70 \pm 1.51^{b}$	$6.10 \pm 1.59^{c}$	1.37	$14.74 + 0.007x - 0.001x^2$	0.946	0.000
MCV (fl)	$85.86 \pm 1.08^{a}$	$86.52 \pm 0.75^{a}$	$86.84 \pm 0.43^{a}$	$82.06 \pm 0.42^{b}$	$80.51 \pm 0.22^{b}$	1.36	$86.03 + 0.048x - 0.001x^2$	0.887	0.000
MCH (pg)	$22.71 \pm 0.49^a$	$23.22 \pm 0.84^{a}$	$23.72 \pm 0.68^{a}$	$20.47 \pm 0.51^{b}$	$19.78 \pm 0.65^{b}$	1.02	$22.82 + 0.035x - 0.000x^2$	0.827	0.000
$MCHC (g dL^{-1})$	$26.46 \pm 0.91^{ab}$	$26.85 \pm 1.21^{a}$	$27.38 \pm 0.92^{a}$	$24.94 \pm 0.48^{bc}$	$24.58 \pm 0.37^{c}$	0.81	$26.53 + 0.028x - 0.000x^2$	0.777	0.008

The data were expressed as mean  $\pm$  standard error of the mean, n=3 triplicates per treatments

Different superscript letters within a row indicate statistically significant (P < 0.05) differences as evaluated by Duncan multiple range test PSEM, pooled standard error of the mean; RSM, red seaweed meal; Hb, hemoglobin; Hct, hematocrit; RBC, red blood cell; WBC, white blood cell; MCV, mean corpuscular volume; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration



**Table 5** Digestive enzymes of *L. rohita* fingerlings fed with experimental diets for 60 days

Parameters	Experimental diets						Regression	n	
	RSM0	RSM25	RSM50	RSM75	RSM100		Equation $R^2$		P value
Amylase (IU mg <sup>-1</sup> protein)	$3.08 \pm 0.17^{b}$	$3.22 \pm 0.16^{b}$	$3.56 \pm 0.10^{a}$	$2.15 \pm 0.11^{d}$	$2.76 \pm 0.14^{c}$	0.63	$3.180 + 0.002x - 9E - 05x^2$	0.298	0.000
Protease (IU mg <sup>-1</sup> protein)	$0.21 \pm 0.06^{ab}$	$0.23 \pm 0.04^{ab}$	$0.27 \pm 0.05^{a}$	$0.20 \pm 0.04^{ab}$	$0.17 \pm 0.03^{b}$	0.02	$0.208 + 0.002x - 2E - 05x^2$	0.789	0.166

The data were expressed as mean  $\pm$  standard error of the mean, n=3 triplicates per treatments

Different superscript letters within a row indicate statistically significant (P < 0.05) differences as evaluated by Duncan multiple range test PSEM, pooled standard error of the mean; RSM, red seaweed meal

bioactive compounds enhanced fish growth (Saleh 2020). Our findings are in agreement with results obtained by Ergun (2009) who reported that lower-level inclusion of Ulva seaweed meal improves the growth of the fish through positive variations in the WG, PER, and FCR by the optimum level of lipids in the diets of seaweed meal. Marinho et al. (2013) investigated that up to 10% inclusion of Ulva meal doesn't affect the growth of the fish and it possessed higher PER and % of intake digestion which may be due to the presence of enormous amount of amino acids like methionine, phenylalanine, and histidine in red seaweed meal. Wassef (2001) reported that 20% dietary inclusion of Ulva seaweed meal was optimum for the growth of mullets because this seaweed has higher vitamin E content. Wan et al. (2016) suggested that 15% inclusion of Palmaria palmata (red seaweed) in Atlantic salmon enhances the growth and nutrient digestibility due to the absorption of nutrients and efficient digestion. The sulfated polysaccharides present in the *Ulva* sp. (seaweed meal) enhance the growth of the fish, hematology parameters, and immune responses of the Indian edible carp *Labeo rohita* (Harikrishnan et al. 2021). The results obtained by Thepot et al. (2022) indicate that red seaweed meal inclusion in the diet increases the growth of fish by reducing the hemolytic and respiratory burst activity as well as immune system simultaneously. Moreover, the alternatives for fishmeal from plant and animal sources possessed optimum dietary minerals that could enhance the growth of the aquatic organism (Ramasamy et al., 2021). Our data showed that poor survival, feed intake, and growth recorded in RSM75 and RSM100 diet-fed L. rohita might be deleteriously associated with the excess amount of RSM in the diet. This could be due to the decline in the amino acid levels of the diets consisting of fish meal replaced with algae/plant-based sources (Dileep et al. 2021). Higher

Table 6 Antioxidant (SOD) and metabolic (GOT and GPT) enzymes of L. rohita fingerlings fed with experimental diets for 60 days

Parameters	Experimental die	ts				PSEM	Regression		ANOVA
	RSM0	RSM25	RSM50	RSM75	RSM100		Equation	$R^2$	P value
Gills									
SOD (IU mg <sup>-1</sup> protein)	$17.36 \pm 1.48^{c}$	$17.45 \pm 1.31^{\circ}$	$17.40 \pm 1.42^{\circ}$	$24.09 \pm 1.70^{\rm b}$	$27.47 \pm 2.09^{a}$	1.59	$17.28 - 0.044x + 0.001x^2$	0.943	0.000
GOT (IU L <sup>-1</sup> )	$90.75 \pm 4.77^{b}$	$91.54 \pm 6.90^{b}$	$91.88 \pm 5.11^{b}$	$99.82 \pm 6.11^{ab}$	$106.34 \pm 6.43^{a}$	1.45	$90.89 - 0.06x + 0.002x^2$	0.977	0.035
GPT (IU L <sup>-1</sup> )	$127.40 \pm 11.31^{b}$	$129.99 \pm 10.66^{\rm b}$	$132.22 \pm 12.92^{ab}$	$138.69 \pm 11.28^{a}$	$145.77 \pm 13.33^{a}$	0.72	$127.6 + 0.030x + 0.001x^2$	0.995	0.386
Liver									
SOD (IU mg <sup>-1</sup> protein)	$23.33 \pm 1.70^{b}$	$23.77 \pm 2.17^{b}$	$23.91 \pm 1.75^{b}$	$27.60 \pm 2.17^{a}$	$29.63 \pm 1.45^{a}$	0.82	$23.32 - 0.011x + 0.000x^2$	0.957	0.007
GOT (IU L <sup>-1</sup> )	$115.81 \pm 4.90^{\circ}$	$116.66 \pm 5.11^{bc}$	$120.47 \pm 4.28^{\rm bc}$	$126.70 \pm 6.57^{\mathrm{ab}}$	$130.73 \pm 5.64^{a}$	1.20	$115.3 + 0.059x + 0.001x^2$	0.982	0.027
GPT (IU L <sup>-1</sup> )	$221.53 \pm 20.09^{\rm b}$	$225.68 \pm 14.62^{\rm b}$	$233.16 \pm 22.9^{b}$	$240.85 \pm 25.41^{a}$	$248.86 \pm 27.73^{a}$	0.72	$221.1 + 0.188x + 0.000x^2$	0.997	0.595
Muscle									
SOD (IU mg <sup>-1</sup> protein)	$14.06 \pm 1.27^{c}$	$14.87 \pm 2.09^{bc}$	$14.91 \pm 1.64^{bc}$	$17.92 \pm 1.81^{ab}$	$19.65 \pm 2.14^{a}$	0.68	$14.12 + 0.001x + 0.000x^2$	0.958	0.016
GOT (IU L <sup>-1</sup> )	$73.32 \pm 8.88^{\rm b}$	$74.51 \pm 6.81^{b}$	$72.32 \pm 3.92^{b}$	$82.47 \pm 4.69^{ab}$	$89.70 \pm 6.75^{a}$	2.62	$73.80 - 0.116x + 0.002x^2$	0.938	0.034
$GPT \; (IU \; L^{-1})$	$97.40 \pm 5.31^{b}$	$97.88 \pm 4.93^{b}$	$97.92 \pm 6.94^{\rm b}$	$108.14 \pm 9.40^{ab}$	$111.85 \pm 8.69^{a}$	2.67	$97.18 - 0.033x + 0.001x^2$	0.923	0.096

The data were expressed as mean  $\pm$  standard error of the mean, n=3 triplicates per treatments

Different superscript letters within a row indicate statistically significant (P < 0.05) differences as evaluated by Duncan multiple range test PSEM, pooled standard error of the mean; RSM, red seaweed meal; SOD, superoxide dismutase; GOT, glutamic oxaloacetic transaminase; GPT, glutamic pyruvic transaminase



inclusion levels of *U. rigida*, *C. barbata*, and *S. limacinum* incorporated diets had no significant effects on the growth and survival rate of O. niloticus and Epinephelus lanceolatus compared with fish fed with a control diet (Güroy et al. 2007; Azaza et al. 2008; García-Ortega et al. 2016). The decreased trend of FCR recorded up to RSM50 diet-fed fish indicates the fact that the formulated diets were superior in quality. Nonetheless, feed conversion efficiency (FCE) was similar in O. mykiss and Sciaenops ocellatus, fed with 50% inclusion of seaweed (U. lactuca, Enteromorpha linza, S. limacinum, and Arthrospira sp.) incorporated diets (Yildirim et al. 2009; Perez-Velazquez et al. 2018). Moreover, in this study, dietary FM replacement by RSM35, RSM34, and RSM37 inclusion promotes the most significant level of WG, SGR, and PER, as estimated by the polynomial regression equation. However, the RSM level that recorded the lowest FCR was RSM25. Several other studies indicated that the seaweed (Gracilaria bursa-pastoris, U. rigida, Gracilaria cornea, and Gracilaria arcuata) meals have essential amino acids, fatty acids, and other required compounds required for the growth and survival of European sea bass D. labrax and O. niloticus (Valente et al. 2006; Younis et al. 2018). According to Vazirzadeh et al. (2022), dietary inclusion of seaweeds (Gracilariopsis persica, Hypnea flagelliformis, and Sargassum boveanum) in O. mykiss had no adverse effects on fish growth and flesh quality. The seaweeds were used as a 10-15% replacement for soymeal and wheat flour in the fish diet. The study of Sáez et al. (2020) stated that low level (5%) inclusion of macroalgae *Ulva ohnoi* in diets of *Solea senegalensis* (Senegalese sole) improved the fillet quality.

The experimental fish fed with RSM50 in the diet had significantly increased the protein, lipid, and ash content when compared with control and other experimental diets suggesting that the optimum levels of dietary RSM compromised the nutritional quality of the experimental diets. López et al. (2012) demonstrated that FM replacement with 75% of Arthrospira meal and 25% of soybean meal mixture had significant results on the growth, feed utilization efficiency, and chemical composition of juvenile O. mykiss. Furthermore, in the present study, polypeptide bands at 10, 13, 15, 17, 23, and 37 kDa shown in SDS-PAGE were more intense in the muscle tissues of FM replaced with RSM50 diet–fed L. rohita. Red seaweed meal inclusion diets had influenced the protein synthesis due to the formation of ARA/EPA (arachidonic acid/eicosapentaenoic acid) ratio in the eicosanoid metabolism (Osmond et al. 2021). The protein bands were indicating the interactions of TOR (target of rapamycin) signalling which is actively involving in the protein synthesis, and it is influenced by the amino acids present in the RSM included diets (Qin et al. 2022). This recommends that the seaweeds can contribute to the absorption of dietary carbohydrates and protein as energy sources and stimulate the assimilation of nutrients in the fish body (Olvera-Novoa et al. 1998, Kut-Güroy et al. 2007; Yildirim et al. 2009; Walker and Berlinsky 2011; Xuan et al. 2013; Peixoto et al. 2016; Younis et al. 2018; Liao et al. 2022). Carbohydrate levels were gradually increasing in diets from control to RSM100 because carbohydrate sources interchange the monomers involved in membrane transport modes through the molecule's metabolism thus increasing the absorption (de Souza et al., 2021). However, Stadtlander et al. (2013) pointed out that during isocaloric (FM replacement with Porphyra yezoensis Ueda meal) diets, Oreochromis niloticus exhibited growth depression in response to excessive/ moderate levels of dietary protein. Similarly, in this study, FM replacement with RSM75 and RSM100 (lower protein levels) diets depress the growth performance and chemical composition of L. rohita.

The use of active immunostimulant ingredients, seaweed meals, extracts, and isolated compounds was suitable for fish cultivation (Peixoto et al. 2016; Lafarga et al. 2020). Seaweed-based immunostimulants have the potential to improve the function of the fish head kidney directly (Thanigaivel et al. 2015; Araujo et al. 2016; Thepot et al. 2021) and also modulate the gut microbiota that enhances immunity indirectly (Guiry and Guiry. 2020). This may influence the hematological (Hb, Hct, RBC, and WBC) status of fish as observed in different studies, because these are the monitoring tool for the fish health status and oxidative stress-related responses (Kalla et al. 2008; Wan et al. 2016; Sotoudeh and Jafari 2017; Hassaan et al. 2019). Generally, algae are rich in phytic acid; it has a tendency to bind with cations as well as proteins and cause modifications in the blood parameters (Musa et al., 2021). The blood chemistry (glutamyl oxaloacetic transaminase, total bilirubin, glucose, blood urea nitrogen, and glutamic pyruvate transaminase) values were not significant when fed with Eucheuma denticulatum included diet-fed Panaeolus olivaceus (Ragaza et al. 2015). Results of hematological indices (Hb, Hct, RBC, WBC, MCV, MCH, and MCHC) in the current study were in general within the normal range when fed with FM replacement with RSM50 incorporated diet. However, hematological constituents were significantly decreased when dietary FM was replaced by RSM75 and RSM100. These were in agreement with the results obtained by Madibana et al. (2017), who reported that the blood parameters (Hct, neutrophil, monocyte, thrombocytes, eosinophil, and basophil) were a functional tool for assessing the enzyme ratio in the blood which plays a crucial role in cell damage to fish fed with seaweed Ulva sp. meal.

The changes in digestive enzyme activity in fishes indicate physiological responses to different nutritional (carbohydrates, lipids, and proteins) constituents of

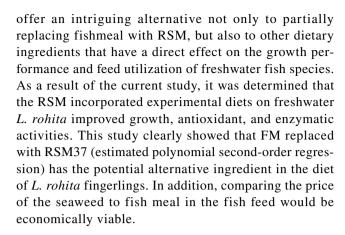


experimental diets (Hidalgo et al. 1999; Fernandez et al. 2001; Hani et al. 2018). Amylase and protease enzyme levels were based on the effects of the diets fed by the fish related to the feed utilization (Pradhan et al. 2014). Our findings have shown that higher enzyme activities (amylase and protease) in FM replacement with RSM50 diet-fed fish contain around 39% of protein and 34% of carbohydrate levels. Higher digestive enzyme activity facilitates increased digestion by the diet, and gut microbiota coordinates the fish for digestion and absorption of nutrients (Kamunde et al. 2019). These results indicated that the incorporated RSM acted as a carbohydrate and protein source, and it was well utilized by L. rohita. Similarly, previous studies showed increased digestive enzymes (protease, amylase, and lipase) in fishes, O. mykiss and O. niloticus, fed with 30% seaweed meals of Phytolacca dioica, Sargassum muticum, Spirodela polyrrhiza, Gracilaria vermiculophylla, and Ulva spp. (Fasakin et al. 2001; Pereira et al. 2012; Silva et al. 2015; Tharaka et al. 2020).

In this study, the activities of antioxidant (SOD) and metabolic (GOT and GPT) enzymes were insignificant in gills, liver, and muscle tissues of fish fed with FM replacement up to RSM50 diet. Meanwhile, significant alterations were observed in antioxidants and metabolic enzyme activities in RSM75 and RSM100 incorporated diet-fed fish due to the activation of the antioxidant pathway by the red seaweed meal, which is an excellent source of antioxidant properties (Kiadaliri et al. 2020). Similarly, increased activities of two liver-bound enzymes, alkaline phosphatase (ALP) and alanine aminotransferase (ALT), were also observed in fish fed with *Ulva* meal-rich diets indicating increased antioxidants levels by seaweed components (Madibana et al. 2017). These enzymes are used to maintain the nutritional status of the blood vascular system and liver functions. Variations in these enzyme levels led to plasma membrane damage (Hassaan et al. 2019). Dietary antioxidants like red seaweed meal prevent oxidative stress in fish (Batista et al. 2020). The present study results are similar to that of the previous studies that demonstrate dietary red seaweed supplementation improves antioxidant capacity indirectly by modifying the activities of antioxidant defense mechanisms (Kim and Lee 2008; Luo et al. 2012; Queiroz et al. 2014; Thanigaivel et al. 2015).

# **Conclusion**

The results of the present study indicate the fact that RSM incorporation in aquafeeds (39% protein level) could be an essential tool to increase the production of freshwater fish *L. rohita* fingerlings. These findings



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Author contribution Dinesh Babu Manikandan: Formal analysis, investigation, resources, methodology, validation, visualization, and writing—original draft. Srinivasan Veeran: Formal analysis, investigation, resources, methodology, data curation, validation, visualization, writing—original draft, and writing—review and editing. Subburaj Seenivasan: Formal analysis, investigation, methodology, validation, visualization, and writing—original draft. Arun Sridhar: Investigation, methodology, data curation, and visualization. Manikandan Arumugam: Formal analysis, investigation, methodology, and visualization. Zhou Yangen: Formal analysis, validation, and writing—review and editing. Thirumurugan Ramasamy: Conceptualization, formal analysis, investigation, methodology, project administration, supervision, writing—original draft, and writing—review and editing.

**Data availability** All data generated or analyzed during this study are included in this published article.

Code availability Not applicable.

### **Declarations**

**Ethics approval** All applicable international, national, and/or institutional guidelines for the care and use of animals were followed by the authors.

Consent to participate Not applicable.

Consent for publication Not applicable.

**Conflict of interest** The authors declare no competing interests.



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