

1 **EnzoMeal as a fish meal replacer: effects on growth, digestive and**
2 **oxidative status of yellow perch (*Perca flavescens*)**
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1 **Abstract**

2 This experiment was conducted to develop a tandem chemical and enzymatic treatment of
3 soybean meal (SBM) to reduce the antinutritional factors (ANFs), and to produce EnzoMeal
4 (EZM) for fish feed. Using yellow perch (*Perca flavescens*) fingerlings, a 10-week
5 experiment was conducted to evaluate the nutritional quality of the EZM to compare with that
6 of fish meal (FM) and SBM. Fingerlings (297; 11.01 ± 0.19 g) were randomly distributed in
7 nine treatments with three replicates and fed iso-nitrogenous diets (crude protein 41%):
8 control (FM based protein); S₅₀ and S₁₀₀ (50% and 100% of FM protein replaced by SBM),
9 and EZM₅₀ and EZM₁₀₀ (50% and 100% of FM protein replaced by EZM). These diets were
10 also supplemented with an exogenous enzyme cocktail to reduce ANFs and named as S_{50+E},
11 S_{100+E}, EZM_{50+E}, and EZM_{100+E}, respectively. All diets were processed in a Brabender single
12 screw extruder, using a 1:1 compression ratio, with processing temperatures of approximately
13 40°C. The highest ($P < 0.05$) growth performance and nutrient utilization parameters (protein
14 efficiency ratio and protein productive value, PPV) were observed for the EZM_{50+E} group,
15 which were not statistically different to that for control and EZM₅₀ groups, and significantly
16 ($P < 0.05$) higher than all other groups. The least ($P < 0.05$) growth performance was
17 observed for the S₁₀₀, S_{100+E} and EZM₁₀₀ groups. On the other hand, the opposite trend was
18 observed for the feed conversion ratio. Digestive enzymes (amylase, lipase and protease)
19 activity was higher ($P < 0.05$) in the intestine than pyloric caeca of fish in all groups. The
20 highest protease activity (in intestine and pyloric caeca) was observed for the control group,
21 which is significantly similar to EZM₅₀, EZM_{50+E}, and S_{50+E} groups, and the lowest value was
22 observed for 100% replacement of FM protein by SBM and EZM fed groups. However,
23 inclusion of exogenous enzymes in feed showed positive effects in EZM_{50+E} compared to
24 EZM₅₀ for PPV and lipid productive value and amylase activity in the intestine. Activity of
25 protein metabolism enzymes i.e., alanine transaminase and aspartate aminotransferase in the

1 liver were the highest in the control, which was similar ($P < 0.05$) to the EZM_{50+E} and EZM₅₀
2 groups, whereas other groups exhibited lower activity. Antioxidant enzymes (catalase,
3 superoxide dismutase, and glutathione peroxidase) in the liver were the lowest in the control
4 group, which was significantly similar to the group fed the feed with 50% replacement of FM
5 protein, whereas complete (100%) replacement of FM protein exhibited the highest activity.
6 Conclusively, performance of the EZM₅₀ and EZM_{50+E} groups were similar to the FM fed
7 group, and better than SBM fed groups, and EnzoMeal has considerable potential as
8 alternatives to fishmeal in aquafeed.

9

10 **Keywords:** EnzoMeal, antinutritional factors, exogenous enzymes cocktail, growth, protein
11 metabolism, antioxidant enzymes, yellow perch.

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1 **Introduction**

2 Aquaculture is the fastest-growing sector of food production. Traditionally, fish meal (FM)
3 has been the main source of dietary protein for farmed fish. However, the increasing demand
4 in contrast to limited supply for this finite protein source has caused a significant increase in
5 FM prices in the last decade (Kader *et al.* 2010; Kumar *et al.* 2010a; Kumar *et al.* 2010b;
6 Kumar *et al.* 2011a; Kumar *et al.* 2011b; Kumar *et al.* 2012a). Therefore, it is a matter of
7 urgency that alternative protein sources for fish diets be found to support global aquaculture
8 development (Kader *et al.* 2010; Kumar *et al.* 2011a; Kumar *et al.* 2011b). Several alternative
9 proteins have been tested for fish feed, among them soybean appeared to be the best FM
10 replacer for fish diet. It has been reported that 30–50% FM could be replaced by soybean
11 meal (SBM). Soybean meals have a high content of available protein with a well-balanced
12 amino acid profile, a constant composition, a reasonable price, and there is a steady supply
13 available; however, their methionine level is low, and they also contain approximately 30%
14 of indigestible carbohydrates including non-starch polysaccharides (NSP), and several
15 compounds or anti-nutritional factors (ANFs; protease inhibitors, lectin, and phytate) that
16 may disturb the digestive process (Hernandez *et al.* 2007; Kumar *et al.* 2011a; Kumar *et al.*
17 2011b; Kumar *et al.* 2012a; Olli *et al.* 1994; Storebakken *et al.* 2000), as many of them have
18 been reported to hinder digestion and absorption of nutrients and restrict their potential as
19 replacements of FM in fish diets (Bureau *et al.* 1998; Hernandez *et al.* 2007; Klein *et al.*
20 1998; Kumar *et al.* 2011a; Kumar *et al.* 2011b; Kumar *et al.* 2012a; McGoogan & Gatlin
21 1997; Olli & Krogdahl 1994; Rumsey *et al.* 1994).

22 Digestion and absorption of nutrients depends on the activity of the digestive
23 enzymes, in particular those located in the intestine, which are responsible for the final stages
24 of breaking down and assimilation of the food (Klein *et al.* 1998; Kumar *et al.* 2011a; Kumar
25 *et al.* 2011b; Kumar *et al.* 2012a; Silva *et al.* 2010). In salmonids, higher inclusion levels of

1 solvent-extracted SBM led to a marked reduction in the activities of such enzymes in
2 enterocytes of the distal intestine (Bakke-McKellep *et al.* 2000; Krogdahl *et al.* 1995;
3 Krogdahl *et al.* 2003; Kumar *et al.* 2011a; Kumar *et al.* 2011b; Kumar *et al.* 2012a; Silva *et*
4 *al.* 2010). This suggests that measuring the activities of intestinal enzymes may represent a
5 sensitive tool to study the effects of differently-processed SBMs and cocktail exogenous
6 enzymes on nutrient bioavailability and to ascertain tolerability to certain soy-ANFs in
7 various fish species.

8 In addition, nutrition plays an important role in animal welfare by maintaining this
9 fragile oxidative balance, either by supplying nutrients that enhance the antioxidant system or
10 avoiding those that would induce an increase of reactive oxygen species (ROS) production by
11 different physiological pathways. The oxidative stress in aquatic organisms is more profound
12 during nutritional deficiency, elevated temperature, hypoxia and exposure to xenobiotics
13 (Avanzo *et al.* 2002; Hwang & Lin 2002; Kolkovski *et al.* 2000; Radhakrishnan *et al.* 2014;
14 Romeo *et al.* 2000). However, either an increase in ROS production above the level that can
15 be removed by antioxidant defenses, or a decrease in the capacity of the antioxidant defenses,
16 could result in oxidative damage to key molecules, including DNA, protein, and lipids (lipid
17 peroxidation) (Halliwell & Gutteridge 1999).

18 While the predominant concern about the effects of various alternative plant proteins
19 is on fish growth and feed efficiency, relatively few studies have monitored the dietary
20 influence on the biochemical index of fish, such as changes in protein metabolism enzyme
21 activities (Krogdahl *et al.* 2003; Kumar *et al.* 2010b; Kumar *et al.* 2011a; Kumar *et al.* 2011b;
22 Kumar *et al.* 2012a), hepatic metabolism (Vilhelmsson *et al.* 2004) and oxidative status,
23 which have also been used to provide an indication of disturbance by specific ingredient to
24 the metabolic function and nutrient utilization by fish (Kumar *et al.* 2011a; Kumar *et al.*
25 2011b; Kumar *et al.* 2012a; Lin & Luo 2011).

1 Yellow perch (*Perca flavescens*) is a major species of the Great Lakes region,
2 particularly in the north central region of the USA, and is a consumer favorite due to low fat
3 content and good taste (Fallahi *et al.* 2012; González *et al.* 2006). However, commercial
4 production of yellow perch has not rapidly developed yet, partially due to limited information
5 on nutritional requirements. A commercially available specialized diet has not been
6 developed for yellow perch. Therefore, it has made it essential to develop suitable complete
7 and supplemental diets for use in yellow perch. It may well be that fish with more
8 omnivorous feeding habits, like yellow perch, could make more efficient use of high levels of
9 dietary SBM.

10 To our knowledge, no studies have been reported on the use of enzyme-treated SBMs
11 as a FM substitute in yellow perch diets. The aims of this study were to 1) test a modified
12 soybean meal diet (EnzoMeal) against a regular soybean meal diet and a standard fish meal
13 diet using yellow perch, and determine the benefits and deficiencies of the EnzoMeal as a fish
14 food ingredient; and 2) evaluate the possibility of replacing different levels of FM protein by
15 SBM and EZM in the practical diet of yellow perch.

16

17 **Materials and methods**

18 This study and all experimental procedures involving animals were performed according to
19 the protocol approved by the Ohio State University Institutional Animal Care and Use
20 Committee.

21 *Preparation of EnzoMeal and diet formulations*

22 The EZM was obtained by heating SBM (NUTRASOY® 7B, obtained from Archer Daniels
23 Midland, Decatur, IL, USA) in 3% hydrogen peroxide solution using a proprietary catalyst.
24 The product was isolated and further treated with an enzyme cocktail.

1 Prior to feed formulation, the proximate composition of SBM, whole wheat meal, EZM, soy
2 protein isolate, and FM were determined. Nine isonitrogenous and isolipidic diets were
3 formulated. Experimental diets containing crude protein 41%, crude lipid 11%, vitamin
4 premix 2%, mineral premix 2%, and Titanium oxide (TiO₂) 0.2% were prepared. TiO₂ was
5 added for digestibility measurement. Methionine was supplemented at the rate of 0.3% in
6 100% FM protein replacement diets. Experimental diets are as follows: control (FM based
7 protein); S₅₀ and S₁₀₀ (50% and 100% of FM protein replaced by SBM), and EZM₅₀ and
8 EZM₁₀₀ (50% and 100% of FM protein replaced by EZM), and SBM and EZM based diets
9 were also supplemented with an exogenous enzyme cocktail (phytase, NSPase, and
10 carbohydrase) and named as S_{50+E}, S_{100+E}, EZM_{50+E} and EZM_{100+E} respectively (Table 1).

11

12 *Enzyme cocktail supplementation*

13 Enzymes (cocktail) and buffer were mixed only with SBM and EZM and kept at room
14 temperature for three hours to hydrolyze the carbohydrates and phytate. Thereafter, these
15 soybean products (hydrolyzed SBM and EZM) were mixed along with other fish feed
16 ingredients for fish diet preparation.

17 *2.1.2. Feed extrusion:* All ingredients were ground with a Wiley mill (Model 4, Thomas
18 Scientific, Swedesboro, NJ, USA) to an average particle size of approximately 500 µm. All
19 components were then combined and mixed for 15 min (Kushlan Products, Inc., Goldendale,
20 WA). After all ingredients were thoroughly homogenized, each blend was adjusted to a
21 desired pre-extrusion moisture content of ~40% by adding adequate amounts of water, then
22 mixed again for 15 min. The extrusion processing of each blend was performed using a single
23 screw extruder (Brabender Plasti-Corder, Model PL 2000, South Hackensack, NJ, USA),
24 which had a compression ratio of 3:1, a screw length-to-diameter ratio of 20:1, and a barrel
25 length of 317.5 mm. The die assembly was conical, and tapered from an initial diameter of

1 6.0 mm to a diameter of 3.0 mm at the discharge opening. The length of the die was 27.0 mm,
2 which resulted in a die length-to-diameter ratio of 9.0. A 7.5 HP (5.5 kW) motor powered the
3 extruder. The screw speed was set to 100 rpm during extrusion, and the temperature of all
4 extrusion zones (feed, transition, and die) were set to 40°C for the duration of processing.
5 After extrusion, the pelletized feed blends were dried in a laboratory oven (Thelco Precision,
6 Jovan, Winchester, VA) at 50°C for 24 h. After drying, the diets were broken by hand, sieved
7 into proper pellet size, and were stored at -15°C. Enzymes may have been active during the
8 feed drying process, thus hydrolyzing substrates in the feed ingredients.

9

10 *Experimental system and animals*

11 A mixed strain of yellow perch fingerlings (about 10-12 g) was produced in the breeding
12 program at the Ohio Center for Aquaculture Research and Development at The Ohio State
13 University South Centers, Piketon, OH, USA. These fingerlings (N = 297) were randomly
14 distributed in nine groups with three replicates, each having 11 fish (11.01 ± 0.19 g) in a
15 round tank (55 L capacity). At the start of the experiment, six fish were preserved at -20°C
16 for analysis of the initial body composition. After fish were acclimatized for two weeks in the
17 experimental tanks, fish were trained for different experimental diets for 1 week. The all
18 tanks were supplied with water from a flow-through system. The system was subjected to a
19 photoperiod of 12 h light: 12 h darkness. Water quality was monitored throughout the
20 experiment. All the water parameters were in the optimum range (temperature 17.4 – 20.1°C,
21 pH 7.0 – 7.5, and dissolved oxygen 6.7 – 8.4 mg l⁻¹). Water flow was adjusted to the same
22 level for all tanks. During the experimental period, a ration totaling 3% body weight daily
23 was hand-fed to the fish in three equal proportions at 9:00, 13:00, and 17:00 h. Fish were
24 weighed individually at the beginning of the experiment and at every other week intervals
25 during the experimental period to adjust the feeding level for the subsequent week. On the

1 designated weigh days, the fish were weighed in the morning, just prior to the 9:00 h feeding,
2 having not had feed for 16 hours prior.

3 The experiment lasted for 10 weeks. At the end of the experiment, three fish per
4 group were anaesthetized with tricaine methanesulfonate (MS222; 250 mg/L). Anaesthetized
5 fish were carefully dissected to isolate the intestine, pyloric caeca, and liver. These samples
6 were stored at -80°C for determination of activities of digestive, protein metabolism and
7 antioxidant enzymes. Four additional fish per group were killed by a blow to the head with a
8 metal rod and then stored at -20°C for chemical composition analysis. Prior to determination
9 of the proximate composition, the fish were autoclaved at 121°C for 20 min, thoroughly
10 homogenized using an Ultra-Turrax T25, frozen overnight and freeze-dried.

11

12 *Proximate analysis of feed ingredients, experimental diets, and whole body of fish*

13 All samples were ground to a fine powder by using a grinder prior to analyses. The proximate
14 composition of the feed ingredients, experimental feeds, and of the fish carcasses was
15 determined according to the official methods (AOAC, 1990), i.e., for moisture (oven-drying
16 at 105°C overnight), crude protein by CN analyser (N X 6.25), fat [by extraction according to
17 the method described by Smedes (1999) as modified by Schlechtriem *et al.* (2003) and ash
18 (oven incineration at 480°C overnight).

19

20 *Growth and nutrient utilization parameters*

21 Growth performance and diet nutrient utilization were assessed in terms of:

22 — Body mass gain percentage (BMG, %) = [(final body mass - initial body mass) /
23 initial body mass] X 100

24 — Specific growth rate (SGR, g/day) = [(ln final body mass in g) - (ln initial body mass
25 in g) / number of trial days] X 100

1 — Metabolic growth rate (MGR, $\text{g kg}^{0.8} \text{ day}^{-1}$) = (Body mass gain in g) / [{"(initial body
2 mass in g / 1000)^{0.8} + (final body mass in g / 1000)^{0.8}} / 2] / number of trial days

3 — Feed conversion ratio (FCR) = dry feed fed (g)/body mass gain (g)

4 — Protein efficiency ratio (PER) = body mass gain (g)/crude protein fed (g)

5 — Protein productive value (PPV, %) = [(final fish body protein in g - initial fish body
6 protein in g) / total protein consumed in g] X 100

7 — Lipid productive value (LPV, %) = [(final fish body lipid in g - initial fish body lipid
8 in g) / total crude lipid consumed in g] X 100

9

10 *Organ indices*

11 *Relative intestinal length (RIL), hepatosomatic index (HSI) and intestinal somatic indexes*
12 *(ISI) are calculated as indicated below:*

13 $\text{RIL} = \text{Intestine length (mm)} / \text{body mass (g)}$

14 $\text{HSI} = \text{Liver mass (g)} \times 100 / \text{body mass (g)}$ and

15 $\text{ISI} = \text{Intestinal mass (g)} \times 100 / \text{body mass (g)}$.

16

17 *Enzyme assays*

18 *Digestive enzyme assay*

19 Amylase activity was measured in the pyloric caeca and intestine by using an amylase assay
20 kit (Kit no. # MAK009 SIGMA, Aldrich, USA). Amylase activity was determined using a
21 coupled enzymatic assay, which resulted in a colorimetric product, proportional to the
22 amount of substrate, ethylidene-pNP-G7, cleaved by the amylase. One unit is the amount of
23 amylase that cleaves ethylidene-pNP-G7 to generate 1.0 μmole of p-nitrophenol per minute at
24 25°C. RayBiotech's Protease Activity Assay Kit (CODE: 68AT-Protease-S100, Ray Bio,
25 USA) was used to determine the protease activity in pyloric caeca and intestine of fish. One

1 unit is defined as the amount of protease that cleaves the substrate, to yield an amount of
2 fluorescence equivalent to 1.0 nmol of unquenched fluorescein isothiocyanate labeled casein
3 per minute at 25°C. Lipase activity was determined using a coupled enzyme reaction, which
4 resulted in a colorimetric product proportional to the enzymatic activity present in pyloric
5 caeca and intestine of fish. Lipase Activity was measured using an assay Kit (MAK046
6 Sigma Aldrich, USA). One unit of lipase is the amount of enzyme that will generate 1.0
7 μ mole of glycerol from triglycerides per minute at 37°C.

8

9 *Protein metabolism enzyme*

10 BioVision's Aspartate aminotransferase (AST) assay Kit (Catalog# K753-100, Bio Vision,
11 USA) was used to determine the AST activity in liver of fish. The principle is amino group is
12 transferred from aspartate to α -ketoglutarate. The products of this reversible transamination
13 reaction are oxaloacetate and glutamate. The glutamate is detected in a reaction that
14 concomitantly converts a nearly colorless probe to color. One unit of AST is the amount of
15 enzyme that will generate 1.0 mmole of glutamate per minute at pH 8.0 at 37°C. Alanine
16 transaminase (ALT) activity was quantified by using an ALT activity Assay Kit (Cayman
17 Chemical Item Number 700260, USA). Measurement of the ALT activity was carried out by
18 monitoring the rate of NADH oxidation in a coupled reaction system employing lactate
19 dehydrogenase (LDH). The oxidation of NADH to NAD^+ was accompanied by a decrease in
20 absorbance. One unit is defined as the amount of enzyme that will cause the oxidation of 1.0
21 μ mol of NADH to NAD^+ per minute at 37°C.

22

23 *Antioxidant enzymes*

24 A superoxide dismutase (SOD, inhibition rate %) assay kit (Catalog # 19160 SIGMA, Sigma
25 Aldrich, USA) was used to assess the SOD enzyme activity in the liver of fish. SOD was

1 assayed by utilizing Dojindo's highly water-soluble tetrazolium salt, WST-1 (2-(4-
2 Iodophenyl) - 3-(4-nitrophenyl)-5-(2,4-disulfophenyl) - 2H-tetrazolium, monosodium salt)
3 that produces a water-soluble formazan dye upon reduction with a superoxide anion.
4 *Catalase* enzyme activity was determined with a catalase assay kit (Cayman Chemical Item
5 Number 707002, USA). The determination is based on the reaction of the enzyme with
6 methanol in the presence of an optimal concentration of H₂O₂. The formaldehyde produced
7 was measured spectrophotometrically with 4-amino-3-hydrazino-5-mercapto-1,2,4-triazole
8 (Purpald) as the chromogen. Purpald specifically forms a bicyclic heterocycle with
9 aldehydes, which, upon oxidation, changes from colorless to a purple color. One unit is
10 defined as the amount of enzyme that will cause the formation of 1.0 nmol of formaldehyde
11 per minute at 25°C. A glutathione peroxidase (GPx) colorimetric assay kit (Catalog # K762-
12 100, Bio Vision, USA) was used to determine the GPx activity in the liver. One unit is
13 defined as the amount of enzyme that will cause the oxidation of 1.0 μmol of NADPH to
14 NADP⁺ under the assay kit condition per minute at 25°C.

15 *Statistical analysis*

16 All data were subjected to a one-way analysis of variance ANOVA and the significance of
17 the differences between means was tested using Tukey's HSD (Honestly Significant
18 Difference) test (P < 0.05). The software used was SAS, Version 9.3. Values are expressed as
19 means ± standard deviation.

20

21

22 **Results**

23 *Proximate composition of feed ingredient and experimental diets*

24 Total crude protein content in the EZM was 58%, and SBM contains 47.5% crude protein
25 (Table 2), which revealed that crude protein amount increased almost 22%, whereas ANFs

1 especially oligosaccharides, NSPs, and phytate content decreased (>50%) in EZM compared
2 to SBM (data not shown). Heat labile antinutrients such as lectin and trypsin inhibitor were
3 absent in EZM because we have used the roasted SBM.

4 The composition of experimental diets is shown in Table 1. Proximate composition of
5 feed ingredients and experimental diets are shown in Table 2. Experimental diets contained
6 about 41% crude protein and 15% crude lipid and were isonitrogenous and isolipidic.

7

8 *Chemical composition of whole body of fish and organ indices*

9 The effects of different inclusion levels of EZM and SBM with and without supplementations
10 of exogenous cocktail enzymes on the body composition of yellow perch are presented in
11 Table 3. The highest ($P < 0.05$) inclusion of EZM and SBM (100% FM protein replacement
12 groups) regardless the inclusion of exogenous enzymes, exhibited higher ($P < 0.05$) moisture
13 content than that of the control and other groups; whereas crude protein and lipid content
14 exhibited an opposite trend. The highest ($P < 0.05$) moisture content was observed in the S_{100}
15 group, which was significantly similar ($P > 0.05$) to the 100% replacement groups (S_{100+E} ,
16 EZM_{100} and EZM_{100+E}), whereas the lowest value was found in the control group. Whole
17 body protein content was the highest ($P < 0.05$) in the EZM_{50+E} group, which is statistically
18 similar to EZM_{50} , S_{50+E} , and control groups; whereas the lowest value was observed for 100%
19 replacement groups (S_{100} , EZM_{100} , and S_{100+E}). The highest ($P < 0.05$) crude lipid content was
20 observed in EZM_{50} , which was similar to S_{50} and the control groups, and these values were
21 statistically higher than other groups. Dietary soybean protein and exogenous enzyme
22 supplementation did not significantly ($P > 0.05$) affect the ash content in the whole body of
23 the fish.

24 Hepato somatic index, GSI, CI, and RIL of yellow perch fed different experimental
25 diets are shown in Table 3. The hepatosomatic indexes were higher in EZM fed groups

1 (except EZM_{100+E}) than regular SBM fed groups. The relative intestinal length values of
2 plant-fed groups were significantly ($P < 0.05$) higher than the control group. EZM and SBM
3 did not influence ($P > 0.05$) the GSI of fish among the groups.

4

5 *Fish behavior, feed intake, growth and nutrient utilization*

6 Based on the visual observation during the feeding period, palatability or acceptability of feed
7 was good and the behaviour of fish was normal, except in the 100% FM replacement groups,
8 wherein these groups exhibited lower palatability than other groups.

9 The effects of the two processed soybean meals (EZM and SBM) with and without
10 the exogenous enzyme cocktail for growth performance, feed intake and nutrient utilization
11 are summarized in Tables 3 and 4, and Figures 1A and 2. The highest ($P < 0.05$) growth
12 performance (FBM, weight gain, SGR, and MGR) were observed for the EZM_{50+E} group,
13 which were not statistically different to that for the control and EZM₅₀ groups, and
14 significantly ($P < 0.05$) higher than all other groups. The least ($P < 0.05$) growth performance
15 was observed for the S₁₀₀, S_{100+E}, and EZM₁₀₀ groups. On the other hand, the opposite trend
16 was observed for the FCR. Feed intake was positively correlated to the growth performance
17 (tables 3 and 4). The highest ($P < 0.05$) PER, PPV, and LPV values were observed for the
18 S_{50+E} group which is statistically similar to the control and EZM₅₀ groups. The lowest values
19 for these parameters were observed in the 100% FM replacement groups. Overall, EZM fed
20 groups exhibited higher ($P < 0.05$) growth performance and feed utilization than SBM fed
21 groups; whereas dietary supplementation of exogenous enzymes both in EZM or SBM did
22 not significantly affect the growth performance and feed utilization of yellow perch except
23 PPV and LPV in the EZM_{50+E} group.

24

25 *Enzymes*

1 Digestive enzyme activities in pyloric caeca and the intestine of yellow perch are presented in
2 Figure 3A, 3B and 3C. Digestive enzymes (amylase, lipase, and protease) activity was higher
3 ($P < 0.05$) in the intestine than in pyloric caeca of fish in all groups. Amylase activity in
4 pyloric caeca and intestine was the highest ($P < 0.05$) in EZM_{50+E} whereas other groups
5 exhibited lower ($P < 0.05$) value. Inclusion of exogenous enzymes in feed showed positive
6 effects ($P < 0.05$) in EZM_{50+E} compared to EZM_{50} for the amylase activity in the intestine. The
7 highest protease activity (in intestine and pyloric) was observed for the control group, which
8 is significantly similar to the EZM_{50} , EZM_{50+E} , and S_{50+E} groups, and the lowest ($P < 0.05$)
9 value was observed for 100% replacement of FM protein by SBM and EZM fed groups.
10 Lipase activity in the intestine was the highest ($P < 0.05$) in EZM_{50+E} , which was significantly
11 similar ($P > 0.05$) to the control, EZM_{50} , and 50% replacement groups (FM replaced by SBM),
12 whereas 100% FM replacement groups exhibited lower activities of lipase in the intestine.
13 Amylase activity in the pyloric caeca did not differ significantly ($P > 0.05$) among the groups.

14 Dietary inclusion of EZM and SBM significantly ($P < 0.05$) changes the activities of
15 protein metabolism enzymes (ALT and AST) in liver of yellow perch (Figure 4A). Activities
16 of protein metabolism enzymes i.e., ALT and AST in liver were the highest in control, which
17 was similar ($P < 0.05$) to the EZM_{50+E} and EZM_{50} groups whereas other groups exhibited
18 lower activity. Activities of protein metabolism enzymes were positive correlated with
19 growth performance and feed utilization. Alanine transaminase and AST activities in liver
20 exhibited positive response (not significantly) with dietary inclusion of the exogenous
21 enzyme cocktail in EZM and SBM based diets.

22 Antioxidant enzyme (catalase, SOD, and GPx) activities in liver were significantly
23 altered by different experimental diets (Figures 1B and 4B). Antioxidant enzymes such as
24 catalase, SOD, and GPx in the liver were the lowest ($P < 0.05$) in the control group, which
25 was significantly similar to the 50% replacement of FM protein fed groups, whereas groups

1 having complete (100%) replacement of FM protein exhibited the highest activity. Dietary
2 supplementation of exogenous enzyme cocktail in SBM and EZM based diets did not
3 influence ($P > 0.05$) the antioxidant enzyme activity in fish.

4

5 **Discussion**

6 The use of plant protein products in aquaculture diets is generally limited by their low levels
7 of digestible protein and/or energy (Kumar et al., 2011a; Kumar et al., 2011b). The seed
8 meals of soybeans, lupins, pea, and Jatropha represent some of those plant products that
9 contain a considerable amount of proteins, and efforts have been made to further enhance
10 their protein levels by processing technologies (Glencross *et al.* 2005; Kumar *et al.* 2012a;
11 Makkar & Becker 2009; Overland *et al.* 2009). In the present study we have developed a
12 method to produce soybean meal containing a high amount of protein (58%) with no ANFs
13 (oligosaccharides, NSP, and phytate), and tested efficacy in replacing fish meal in yellow
14 perch. To our knowledge, no studies have been reported on the use of enzyme-treated SBM
15 as a FM substitute in yellow perch diets.

16

17 *Feed intake, growth and nutrient utilization*

18 The results of the present study clearly show that the nutritive value of complete diets for
19 yellow perch may vary when differently processed soybean meals (roasted SBM and EZM
20 along with or without the inclusion of exogenous enzymes) are used to replace varying
21 amounts of FM protein. In this experiment relative to control (fish meal based diet),
22 palatability was not adversely affected by type of SBM (regular SBM or EZM) inclusion in
23 the diet; however, the level of SBM inclusion adversely affected the feed intake. In general,
24 feed intake has an inverse relationship to higher FM replacement levels with plant proteins in

1 omnivorous and carnivorous fish (Kader *et al.* 2011; Kader *et al.* 2012; Uyan *et al.* 2006). In
2 the present experiment, feed intake was lower in the 100% FM replacement groups compared
3 to the 50% replacement groups. However, there were no significant effects of
4 supplementation of exogenous enzymes in either EZM or SBM based diets on feed intake and
5 growth performance. In this study, there was a decreasing trend in feed intake with increasing
6 dietary SBM and EZM level, similar to the findings of some previous studies (Espe *et al.*
7 2006; Kaushik *et al.* 2004). Kaushik *et al.* (2004) used a regression model, while Espe *et al.*
8 (2006) used a total replacement of FM, adding different ingredients to increase voluntary feed
9 intake. The former reported that feeding diets containing only 0.5% FM to European seabass
10 did not affect feed intake and reduce performance on high inclusion of plant ingredients,
11 while in the latter study only one of the groups reduced feed intake, but all reduced growth as
12 compared to the FM control. These results indicate that the effects on the growth response
13 might directly be attributed to the partly even significantly lower feed intake of fish fed the
14 plant protein diets (Espe *et al.* 2006). Hence, in our study, reduced feed intake (consequently
15 a lower intake of essential nutrients and digestible energy) could be the main reason for
16 reduced growth performance of yellow perch as the dietary SBM and EZM level increased.

17 Reduced nutrient/energy bioavailability and growth in response to complete feeds
18 containing high levels of conventionally-processed SBM and EZM, relative to FM-based
19 diets, has been observed in almost all carnivorous and omnivorous fish species, including
20 yellow perch investigated to date, and various causes and mechanisms have been claimed as
21 possible reasons. Of these, oligosaccharides and NSPs in soy preparations have been
22 suggested to lead to reduced bioavailability of all nutrients and energy through mechanisms
23 involving a binding action with bile salts and/or by an obstructing action on digestive
24 enzymes coupled with changes in digesta viscosity and transit rate (Francis *et al.* 2001;
25 Tibaldi *et al.* 2006). This is supported by findings that soy-protein derivatives obtained

1 through processes, which ensure partial (50%) removal of non-starch carbohydrate fractions,
2 like EZM or the enzyme-treated SBMs here tested, resulted in higher nutritive value relative
3 to regular SBM when used at 50 and 100% replacement levels (Bureau *et al.* 1998;
4 Hernandez *et al.* 2007; Kumar *et al.* 2010a; Kumar *et al.* 2010b; Kumar *et al.* 2011a; Kumar
5 *et al.* 2011b; McGoogan & Gatlin 1997; Olli & Krogdahl 1994; Rumsey *et al.* 1994).
6 Deficiencies in minerals or essential amino acids may also have impacted the ability of perch
7 to utilize soy products at levels greater than 500 g/kg (Brown *et al.* 1997; Kasper *et al.* 2007;
8 Ketola 1975). In the present study, the use of the EZM with zero oligosaccharides to
9 substitute up to 50% FM protein in the diet resulted in nutrient utilization (FCR, PER, PPV
10 and LPV) values and growth performance, which did not differ from those measured in fish
11 fed the control diet (Refstie *et al.* 1999). Our results are in concurrence with many studies
12 (Denstadli *et al.* 2007; Glencross *et al.* 2005; Kaushik *et al.* 2004; Mambrini *et al.* 1999;
13 Overland *et al.* 2009; Vielma *et al.* 2000), which have shown that soy, lupin, and pea protein
14 isolates and soy protein concentrate can replace 50-75% of FM protein in rainbow trout and
15 Atlantic salmon diets without impairing the growth performance and nutrient utilization.
16 Médale *et al.* (1998) observed that total replacement of FM by soy protein concentrate led to
17 a significant decrease in feed intake and resulted in poor growth partially due to methionine
18 deficiency in the soy protein concentrate based diet.

19 Soybean meal fed groups exhibited lower growth performance compared to EZM fed
20 groups. Hence, there seems to be some evidence that a higher intake of soy non-starch
21 carbohydrates lowers the nutritive value of yellow perch diets, but the extent to which this
22 depressive effect is attributable to soy oligosaccharides or NSPs could not be established in
23 this experiment.

24 The weight gain data indicate that SBM/EZM inclusion had no positive impact on
25 production when combined with supplemental exogenous cocktail enzymes (NSPase, phytase

1 and carbohydrase) but there is increasing trend (not significantly) towards growth and
2 nutrient utilization of feed. Our results concur with other reports suggesting no adverse
3 effects when these enzymes are added to feeds (Allan *et al.* 1998; Kocher *et al.* 2003; Stone
4 *et al.* 2003). However, research with other species has shown a positive impact when a
5 supplemental exogenous cocktail enzyme has been included in aquafeeds wherein a
6 multienzyme (amylase, protease, β - gluconase, β -glucosidase, cellulose and combination of
7 these enzymes) in feed significantly improved weight gain, FCR, and other nutritional
8 parameters (Baas & Thacker 1996; Castañón *et al.* 1997; Cowieson *et al.* 2006; Denstadli *et*
9 *al.* 2011). The mechanism by which exogenous multienzymes enhance nutrient digestion and
10 utilization in plant proteins have not been identified, but disruption of cell wall integrity and
11 the breakdown of the highly viscous NSP and carbohydrate were thought to be the major
12 factors involved (Sinha *et al.* 2011). The supplemental exogenous enzymes had been shown
13 to decrease digesta viscosity, feed and protein utilization, and improve digestibilities of
14 amino acids, protein, and lipid in fish (Denstadli *et al.* 2011).

15

16 *Biochemical composition*

17 Efficient protein synthesis requires sufficient availability of all essential amino acids
18 (Dabrowski & Guderly 2002; Kumar *et al.* 2011a; Kumar *et al.* 2011b). Unbalanced amino
19 acid concentrations in a diet resulted in increased protein degradation (Langar *et al.* 1993;
20 von der Decken & Lied 1993), and thereby increased protein turnover (Martin *et al.* 2003).
21 The plant protein (SBM) based diets lower nitrogen retention in salmon and trout because
22 these diets have less digestible energy and an amino acid profile that is suboptimal for muscle
23 growth (Cheng *et al.* 2003; Kumar *et al.* 2011a; Kumar *et al.* 2011b; Pack *et al.* 1995; Refstie
24 *et al.* 2000). A similar pattern was found in the present study. Crude protein content of whole
25 yellow perch was lower in groups fed soy-based protein feeds with 100% FM replacement

1 than in the control and in the groups fed 50% FM protein, regardless the inclusion of
2 exogenous enzymes. However, interestingly, crude protein content in the whole body was
3 higher in the partially (50% replacement) soy based protein fed groups than in the 100%
4 replacement groups. Similarly, Barrows *et al.* (2008) and Cheng *et al.* (2003) also found that
5 the body protein content increased significantly when SBM replaced FM in trout diet. This
6 indicates that the combination of 50% FM and 50% soy based protein diets contain optimum
7 digestible energy and balanced amino acid profile optimal for yellow perch muscle growth.

8 In the present study, exogenous enzyme supplementation did not enhance nutrient
9 utilization and nutrient deposition in yellow perch. Our results are in contrast with other
10 studies, which showed that inclusion of exogenous enzymes in soy based diets increased
11 protein synthesis in fish muscle (Ai *et al.* 2007; Danicke *et al.* 2003).

12 In this study, moisture content exhibited an inverse relationship with lipid content. A
13 similar trend has been reported in a study by Hasan *et al.* (1997), wherein FM protein was
14 replaced by plant protein such as mustard, sesame, linseed, copra, and groundnut oil cakes.
15 Our study also showed that complete (100%) replacement of FM protein by soy protein
16 exhibited lower lipid content in fish than the control group. This suggests that fish were
17 utilizing the dietary lipid for energy purposes and the contribution/utilization of energy from
18 the nitrogen-free extract was minimal. In addition, the change in the efficiency of lipid
19 digestion and absorption from the diet may have contributed to the reduction in whole body
20 total fat level. Partial (50%) replacement of FM protein by soy protein showed higher lipid
21 content in fish than that of 100% FM replacement groups. Similarly, Hasan *et al.* (1997) and
22 Mazurkiewicz (2009) observed that partial FM protein replaced by plant protein in fish diet
23 exhibited higher lipid deposition. There is evidence that partial replacement of FM by plant
24 protein sources such as corn gluten meal and soy protein concentrates increases hepatic
25 lipogenic enzyme activities in seabass that leads to higher whole body lipid amounts (Dias

1 1999; Kaushik *et al.* 2004). Another possible reason for higher lipid retention could be a
2 higher supply of some of the dispensable amino acids such as glutamic acid in excess by the
3 plant protein fed diets (Barrows *et al.* 2008; Kumar *et al.* 2011a; Kumar *et al.* 2011b). These
4 authors also indicated the involvement of possible metabolic or endocrine mechanisms in
5 eliciting such differences in whole body lipid deposition (Kumar *et al.* 2011a; Kumar *et al.*
6 2011b). In the present study, a higher value of HSI in the partial FM protein replacement
7 groups suggests higher lipid deposition in liver. Hepatosomatic index values of above 1, as
8 observed here, are common in yellow perch.

9

10 *Digestive enzyme activities and relative intestinal length (RIL)*

11 Digestion and absorption of nutrients depends on the activity of the digestive enzymes, in
12 particular those located in the pyloric caeca and intestine, which are responsible for the
13 breaking down and assimilation of the feed (Klein *et al.* 1998). The study of digestive
14 enzymes in fish shows a wide range of potential benefits, suggesting that determining of the
15 properties, function and optimal conditions for nutrients (carbohydrate, protein and lipid)
16 hydrolysis of digestive enzymes would provide a more accurate measurement for
17 carbohydrate, protein and lipid digestibility (Klein *et al.* 1998; Silva *et al.* 2010). Therefore,
18 digestive enzyme activities are considered as predictors of potential feed utilization and
19 growth differences in fish (Lin & Luo 2011).

20 The digestive enzymatic activity measured in the pyloric caeca was significantly
21 lower than that in the intestinal digestive tract of yellow perch (Figure 2) because high
22 enzymatic activity in the intestine plays an important role in absorption of the digested
23 nutrients. Bowyer *et al.* (2013) observed the similar results that the lower digestive enzyme
24 activity in the pyloric caeca than the distal intestine, it appears as a symptom of reduced
25 functionality of the distal intestine and, thus, reduced capacity for reabsorption of digestive

1 secretions in salmon fed SBM based diets (Bowyer *et al.* 2013; Krogdahl *et al.* 2003).
2 However, our results are in contrast with other studies (Harpaz & Uni 1999; Harpaz *et al.*
3 2005a; Harpaz *et al.* 2005b; Krogdahl *et al.* 1999), wherein they have observed more of the
4 high enzymatic activity in the pyloric caeca than in the intestine.

5 In the present study, higher (>50% FM protein replacement) inclusion levels of soy
6 protein (SBM and EZM) led to a marked reduction in the activities of digestive enzymes
7 (amylase, protease, and lipase; except lipase activity in pyloric caeca); and these enzymatic
8 activities were found to decrease with lowering SGR and increasing FCR, which show that
9 digestive processes were affected in both the intestine and pyloric caeca with increasing
10 dietary soy protein (SBM and EZM) levels. Current results suggest that enzyme capacity of
11 fish could be improved using nutrients (low dietary inclusion levels of soy protein, 50% FM
12 protein replacement) that stimulate enzyme secretion. Similar results were observed in
13 Atlantic cod and Atlantic salmon (Bakke-McKellep *et al.* 2000; Bowyer *et al.* 2013; Bureau
14 *et al.* 1998; Hidalgo *et al.* 1999; Krogdahl *et al.* 1995; Krogdahl *et al.* 2003; Lemieux *et al.*
15 1999). However, high percentages of soy protein (100% replacement of FM by SBM and
16 EZM) led to decreased digestive enzyme activity, and major reasons could be the presence of
17 heat stable antinutrients (phytate, oligosaccharides, and NSPs) in soy based diets. Phytate is
18 known to inhibit activities of digestive enzymes such as pepsin, trypsin, and alpha-amylase
19 (Alarcon *et al.* 1999; Kumar *et al.* 2010a; Kumar *et al.* 2011a; Kumar *et al.* 2011b; Robaina
20 *et al.* 1995), or to form complexes with minerals (Sugiura *et al.* 1999; Teskeredzic *et al.*
21 1995) and proteins (Moyano *et al.* 1999), thereby modifying digestion processes and
22 impairing intestinal absorption. High content of oligosaccharides and NSP in SBM (15%
23 NSP) and EZM (7% NSP) leads to a significant reduction in the activities of digestive
24 enzymes in the intestine and pyloric caeca (Baeverfjord & Krogdahl, 1996; Bakke-McKellep
25 *et al.* 2000; Bureau *et al.* 1998; Krogdahl *et al.* 1995; Krogdahl *et al.* 2003; Tibaldi *et al.*

1 2006) and even to diminishing of the carrier-mediated nutrient transport/absorption ability
2 (Nordrum *et al.* 2000; Tibaldi *et al.* 2006). The activity of the intestinal enzymes could
3 provide further insight on possible effects of differently modified SBMs on nutrient
4 bioavailability and on the sensitivity of yellow perch intestinal mucosa functions to certain
5 soy protein ANFs.

6 There was no significant effect of soy protein (SBM and EZM) levels on amylase
7 activity in pyloric caeca of yellow perch. Similar observation was reported by various
8 researchers (Lopez-Lopez *et al.* 2005; Lin & Luo *et al.* 2011) in other fish species. It is
9 suggested that animals have the capacity to adapt their digestive physiology in response to
10 changes in their nutrient requirements or dietary profile to some extent.

11 The exogenous enzymes (mixture of NSPase, phytase, and carbohydrase) had been
12 shown to decrease digesta viscosity and improve digestibilities of amino acids, protein, lipid,
13 and starch in domestic animals and fish (Ai *et al.* 2007; Bedford 1995; Classen 1996;
14 Cowieson *et al.* 2006; Sinha *et al.* 2011; Kumar *et al.* 2012b). Also, addition of those
15 enzymes will enhance the utilization of phytate, NSP, and oligosaccharides, which result in
16 higher utilization of protein. However, endogenous digestive enzymes, important for
17 digestion of nutrients, showed contradictory tendency; for example, secretion of pancreatic
18 juice and endogenous digestive enzyme activities increased with increasing viscosity due to
19 viscous NSP (Danicke *et al.* 2003; Ikegami *et al.* 1990). Supplementation of exogenous
20 enzymes significantly decreased the activities of protease, chymotrypsin, amylase, and lipase
21 in fish (Danicke *et al.* 2003; Li *et al.* 2004). However, in our study, supplementation of
22 exogenous enzymes (mixture of NSPase, phytase, and carbohydrase) did not change digestive
23 enzyme activity (except EZM_{50+E} group) in pyloric caeca and intestine. The exact mechanism
24 is not clearly understood in fish. More studies should be conducted to examine the

1 mechanism by which feed utilization could be improved with exogenous enzyme
2 supplementation.

3 Overall, our study suggests that measuring the activities of digestive enzymes may
4 represent a sensitive tool to study the effects of differently-processed SBMs on nutrient
5 bioavailability and to ascertain tolerability to soy-oligosaccharides in yellow perch. The
6 lower activity of digestive enzymes in soy protein (SBM and EZM) fed groups were
7 correlated with lower nutrient utilization parameters (PER, PPV, and LPV).

8 It is well known that carnivorous and omnivorous fish require longer time to digest
9 plant protein based diets compared to animal protein based diets (Buddington *et al.* 1997;
10 Kumar *et al.* 2011a; Kumar *et al.* 2011b). A direct relationship between the amount of dietary
11 plant protein and RIL has been reported earlier in fish (Kramer & Bryant 1995; Kumar *et al.*
12 2011a; Kumar *et al.* 2011b). In yellow perch, soy protein (SBM and EZM) based diets
13 exhibited higher RIL than the control group. RIL value increases as the soy protein (SBM and
14 EZM) inclusion increases in the fish diets (Overland *et al.* 2009). From a physiological view
15 point, a longer RIL would facilitate an increase in digestibility and retention time by
16 enhancing contact time of the digestive enzymes and the feed components, resulting in an
17 increase in their digestion and absorption (Kumar *et al.* 2011a; Kumar *et al.* 2011b).
18 Omnivorous fish such as yellow perch species showed compensation mechanisms, such as an
19 increase in RIL, and as a result increase in digestive activity, to achieve a digestive balance
20 and growth rates similar to those observed for the FM fed group.

21

22 *Protein metabolism enzymes*

23 Alanine aminotransferase and AST are two enzymes for protein metabolism, mostly active in
24 the liver, which are quantitatively important in transamination of amino acids. Alanine
25 aminotransferase and AST activities in liver were significantly different among the groups

1 but supplementation of exogenous enzymes in soy based diets did not change the
2 aminotransferase (ALT and AST) activity in fish. In the present study these enzyme activities
3 were positively correlated with growth performance and feed utilization. In the present study,
4 the activity of protein metabolism enzymes ALT and AST in liver was reduced with
5 increasing dietary SBM and EZM levels, which indicated the utilization of dietary protein
6 decreased and the liver was damaged to a certain extent (Lin *et al.* 2007; Luo 2011).
7 Aminotransferases, such as ALT and AST, catabolize amino acids and transfer amino groups
8 to alpha-keto acids (reversible catalysis) but when the available amino acids are deficient, the
9 keto acids may be reduced, thereby reducing the activity of ALT and AST (Lin and Luo,
10 2011). Decreasing liver AST and ALT activity and a tendency to lower liver ALT and AST
11 activity in fish given diets containing high level of soy protein compared to the low level of
12 soy protein based diets, indicate reduction in substrates available for transamination as a
13 consequence of reduced dietary levels of these amino acids (Lin & Luo 2011).

14

15 *Antioxidant enzymes*

16 Reactive oxygen species (ROS), which include hydroxyl radical, superoxide anion, hydrogen
17 peroxide, and singlet oxygen, are physiologically generated in a series of biochemical
18 reactions within cellular compartments and increase in physiological conditions that result in
19 oxidative stress (Dirks *et al.* 1982; Radhakrishnan *et al.* 2014). The increased levels of ROS
20 may lead to irreversible cell damage and eventually to cell death. Antioxidant enzymes play a
21 crucial role in the defense against oxidative cell damage, through catalyzing the breakdown
22 of superoxide anion to oxygen and hydrogen peroxide (McCord & Fridovich, 1988;
23 Radhakrishnan *et al.* 2014). In normal cells, there exists a delicate balance between the
24 prooxidant forces and antioxidant defenses known as redox balance. Previously some

1 researchers have reported that the overwhelming of antioxidant defenses of cells by
2 prooxidants leads to oxidative stress (Radhakrishnan *et al.* 2014).

3 The nutrient metabolism alterations induced by NSPs (from SBM and EZM) in fish
4 nutrition, like those observed in Nile tilapia, *Oreochromis niloticus*, and African catfish, in
5 which intestinal volatile fatty acid (VFA) production was increased by NSPs (Amirkolaie *et*
6 *al.* 2006; Enes *et al.* 2012; Leenhouders *et al.* 2007; Sinha *et al.* 2011), which may also
7 induce a modification in normal cellular oxidative balance. To minimize ROS related cell
8 damage, fish and other aerobic organisms possess antioxidant defense enzymatic
9 mechanisms, including the key enzymes with antioxidant activity (SOD, CAT and GPx)
10 (Enes *et al.* 2012; Halliwell & Gutteridge, 2007; Martinez-Alvarez *et al.* 2005).

11 In the present study, the activities of antioxidant enzymes (SOD, CAT and GPx) in
12 partial (50% FM replaced by soy protein) replacement groups did not show any significant
13 increase when compared with the control group; whereas 100% replacement groups exhibited
14 higher antioxidant enzyme activity than other groups. Antioxidant enzyme activities in the
15 liver increased as dietary soy protein increased, which suggests a rise in O_2^- generation
16 related to increased levels of NSP and oligosaccharides in soy based diets. It is known that
17 NSP (from SBM and EZM) tends to increase digesta viscosity and thus delay gastric
18 emptying rate in fish (Amirkolaie *et al.* 2005; Enes *et al.* 2012; Leenhouders *et al.* 2006;
19 Sinha *et al.* 2011; Storebakken 1985). This delay of digesta passage through the gut may
20 stimulate NSPs fermentation by intestine microbiota, thus leading to an increase of VFA
21 production. These VFA might have been mobilized for energy production, leading to a rise of
22 O_2^- levels and thus an increase of antioxidant enzyme activity (Choct & Kocher 2000; Enes
23 *et al.* 2012). Another possibility for the suggested rise of O_2^- levels would be an increase in
24 intestinal mucosal cell turnover and consequently, the need of energy supply for the process
25 in fish fed increasing levels of SBM (NSP) (Enes *et al.* 2012; Jin *et al.* 1999; Sinha *et al.*

1 2011). Similar results were observed in other studies wherein plant-containing diets increased
2 the hepatic glutathione redox status as well as antioxidant enzyme activities in fish (Enes et
3 al., 2012). This result indicated that high inclusion of soy protein (SBM and EZM) may exert
4 oxidative stress on yellow perch. In order to eliminate excess free radicals, the fish increased
5 antioxidant enzyme activities to protect the body under normal function (Enes *et al.* 2012;
6 Valko *et al.* 2007). In liver, this was evident by the increase of the GPx activity in a dose-
7 dependent manner with plant protein increment. This higher antioxidant activity could be due
8 to some of the plant constituents such as flavonoids present in soy based diets. Little
9 information about such mechanisms in fish was found in the literature, therefore further
10 research is warranted.

11 In the present study, dietary supplementation of exogenous enzymes did not
12 significantly change the activities of antioxidant enzymes (SOD, CAT, and GPx) in fish,
13 whereas, Zhu *et al.* (2014) found that supplementation of exogenous enzymes significantly
14 reduced SOD and CAT activities in fish. This suggests that exogenous enzymes could protect
15 the fish body from cell oxidative damage when fed mainly with plant source-protein. Another
16 reason is that the enhancement of aerobic metabolism due to dietary phytase supplementation
17 improves the gastrointestinal digestive enzymes (Nwanna, 2007; Zhu *et al.* 2014), and could
18 lead to the increased risk of oxidative stress (Martinez-Alvarez *et al.* 2005; Zhu *et al.* 2014).
19 However, the real mechanisms underlying the dietary exogenous enzymes actions remain
20 largely speculative and further research is required.

21 22 **Conclusions** 23

24 EnzoMeal (EZM) containing 580 g kg⁻¹ crude protein and no anti-nutritional factors, has been
25 evaluated as an alternative for fishmeal in the diet of yellow perch with significant success.
26 Higher growth performance and feed utilization was observed for 50% replacement of FM by

1 EZM fed groups compared to 100% replacement of FM by SBM and EZM fed groups. The
2 activity of protein metabolism enzymes AST and ALT was positively correlated with growth
3 performance and nutrient utilization parameters, whereas antioxidant enzymes (SOD, CAT,
4 and GPx) exhibited an opposite trend. The supplementation of exogenous enzymes (mixture
5 of NSPase, phytase, and carbohydrase) did not exhibit any positive effects on growth
6 performance, nutrient utilization, digestive physiology, protein metabolism, and oxidative
7 status in yellow perch. EnzoMeal with high protein and no ANFs has considerable potential
8 as alternatives to fishmeal in aquafeed. It can be used as one of the promising fishmeal
9 replacers in high-protein fish feeds and substitute 50% - 100% of FM protein without
10 sacrificing fish yield.

11

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18

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1 **Table 1** Composition of the experimental diets (g kg⁻¹ dry matter basis of ingredients) for
 2 yellow perch (*Perca flavescens*) fingerlings.

Experimental diets*	Control	S ₅₀	S _{50+E}	S ₁₀₀	S _{100+E}	EZM ₅₀	EZM _{50+E}	EZM ₁₀₀	EZM _{100+E}
Menhaden fish meal ^a	480.0	240.0	240.0	-	-	240.0	240.0	-	-
Wheat meal ^b	340.8	219.0	219.0	90.0	90.0	284.0	284.0	218.0	218.0
Soybean meal (SBM)	-	325.0	325.0	650.0	650.0	-	-	-	-
EnzoMeal (EZM)	-	-	-	-	-	269.0	269.0	538.0	538.0
Soy protein isolate ^c	-	16.0	16.0	28.0	28.0	7.0	7.0	12.0	12.0
Menhaden oil ^d	50.0	78.0	78.0	107.0	107.0	78.0	78.0	107.0	107.0
Soybean oil ^e	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0
Vitamin premix ^f	30.0	30.0	30.0	30.0	30.0	30.0	30.0	30.0	30.0
Mineral premix ^g	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0
TiO ₂ ^h	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0
Methionine ⁱ	-	-	-	3.0	3.0	-	-	3.0	3.0
Enzyme cocktail ^j	-	-	+	-	+	-	+	-	+
Total	1000	1000	1000	1000	1000	1000	1000	1000	1000

3 **Treatments***

4 **Control** - Fish meal (major source of protein)

5 **S₅₀** - 50% fish meal protein replaced by SBM

6 **S₅₀ +E** - 50% fish meal protein replaced by SBM + enzymes (cocktail)

7 **S₁₀₀** - 100% fish meal protein replaced by SBM

8 **S₁₀₀ +E** - 100% fish meal protein replaced by SBM + enzymes (cocktail)

9 **EZM₅₀** - 50% fish meal protein replaced by EZM

10 **EZM₅₀ +E** - 50% fish meal protein replaced by EZM + enzymes (cocktail)

11 **EZM₁₀₀** - 100% fish meal protein replaced by EZM

12 **EZM₁₀₀ +E** - 100% fish meal protein replaced by EZM + enzymes (cocktail)

13

14 ^aIPC 740; International Proteins Corp., Minneapolis, Minnesota, USA.

15 ^bWhole wheat meal; Bob' s Red Mill Natural Foods, Inc. Milwaukie, OR, USA.

16 ^cSolae, LLC, St. Louis, USA.

17 ^dVirginia Prime; Omega Protein, Inc., Reedville, Virginia, USA.

18 ^eProduct number OF1870E; Consumers Supply Distributing, Sioux City, Iowa, USA.

19 ^fTest Diet; Land O'Lakes Purina Feed, Richmond, Indiana. Pantothenic acid, 4,601 mg/kg;
 20 pyridoxine, 823 mg/kg; riboflavin, 3,000 mg/kg; niacin, 5,000 mg/kg; folic acid, 1,800 mg/kg;
 21 thiamin hydrochloride, 4,503 mg/kg; biotin, 500 mg/kg; vitamin B12, 200 µg/kg; choline, 50,001
 22 mg/kg, menadione, 1,040 mg/kg; vitamin A, 96.1 international units (IU)/g; vitamin D3, 120 IU/g;
 23 vitamin E, 3,000 IU/kg; and ascorbic acid (L-ascorbyl-2-polyphosphate), 17,500 mg/kg.

24 ^gTest Diet; Land O'Lakes Purina Feed. Calcium, 8.00%; phosphorus, 8.00%; potassium, 5.00%;
 25 magnesium, 1.33%; sodium, 3.48%; chloride, 2.80%; fluorine, 144 mg/kg; iron, 1,600 mg/kg; zinc,
 26 1,091 mg/kg; manganese, 276 mg/kg; copper, 126.2 mg/kg; cobalt, 248.1 mg/kg; iodine, 114.68
 27 mg/kg; chromium, 8.0 mg/kg; molybdenum, 5.61 mg/kg; and selenium, 5.02 mg/kg.

28 ^h and ⁱ Sigma Aldrich, Saint Louis, MO, USA.

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30

1 **Table 2** Proximate compositions of feed ingredients and experimental diets (% dry matter
 2 basis).

Feed ingredients	Dry matter	Crude protein	Crude lipid	Nitrogen free extract	Ash
Fish meal	92.45	67.05	10.77	4.45	10.18
Soybean meal	91.24	47.45	2.45	28.58	12.76
EnzoMeal	91.05	58	1.87	24.90	6.28
Soy protein isolate	94.05	91.03	0.74	0.77	4.05
Whole wheat meal	90.19	11.85	2.15	74.53	1.76
Experimental diets*					
Control	95.09	40.78	14.48	27.41	12.42
S₅₀	95.89	41.94	14.56	30.49	8.90
S₅₀+E	94.84	40.48	15.29	30.47	8.60
S₁₀₀	94.41	41.01	14.34	33.23	5.83
S₁₀₀+E	95.91	42.16	14.52	33.52	5.71
EZM₅₀	94.6	42.30	14.9	29.23	8.17
EZM₅₀+E	95.27	41.52	15.35	29.71	8.19
EZM₁₀₀	94.5	41.91	15.09	31.94	4.56
EZM₁₀₀ +E	94.73	42.03	15.27	33.23	4.21

3 * See footnotes to Table 1.

Table 3

Chemical composition of whole body (at the start and at the end of the experiment, g kg⁻¹ wet basis ± SD), relative intestinal length (RIL, mm g⁻¹), hepato somatic index (HSI) and feed intake of yellow perch (*Perca flavescens*) fingerlings of different experimental groups.

	Moisture	Crude protein	Crude lipid	Ash	Calcium	Phosphorus	Magnesium	Potassium
Initial fish	73.79 ± 0.55	14.55 ± 0.58	3.60 ± 0.34	4.69 ± 0.88	1.69 ± 0.01	2.16 ± 0.01	0.02 ± 0.00	0.21 ± 0.00
Treatment*	Moisture	Crude protein	Crude lipid	Ash	RIL	HSI	GSI	Feed intake (g)
Control	67.96 ± 1.19 ^b	16.32 ± 0.71 ^{ab}	9.02 ± 0.35 ^{ab}	3.86 ± 1.52	0.49 ± 0.03 ^c	1.79 ± 0.37 ^{ab}	2.49 ± 0.05	36.96 ± 0.86 ^a
S₅₀	68.03 ± 1.38 ^b	16.95 ± 0.86 ^{ab}	7.27 ± 3.16 ^{ab}	4.79 ± 0.57	0.64 ± 0.02 ^a	1.32 ± 0.41 ^c	2.07 ± 0.22	36.45 ± 0.77 ^a
S₅₀+E	70.86 ± 2.80 ^{ab}	17.77 ± 1.05 ^{ab}	6.20 ± 1.26 ^b	5.09 ± 1.68	0.66 ± 0.01 ^a	1.61 ± 0.41 ^b	2.43 ± 0.64	35.62 ± 2.93 ^a
S₁₀₀	73.63 ± 1.19 ^a	14.67 ± 0.89 ^b	5.15 ± 0.91 ^c	3.67 ± 0.71	0.61 ± 0.05 ^{ab}	1.41 ± 0.21 ^b	2.75 ± 0.28	18.32 ± 0.85 ^b
S₁₀₀+E	73.37 ± 2.76 ^{ab}	15.16 ± 2.24 ^b	7.64 ± 1.34 ^b	4.92 ± 0.41	0.64 ± 0.04 ^a	1.54 ± 0.05 ^b	2.84 ± 0.29	18.98 ± 0.78 ^b
EZM₅₀	69.15 ± 1.29 ^{ab}	17.97 ± 0.53 ^a	9.65 ± 0.26 ^a	4.28 ± 1.02	0.55 ± 0.05 ^b	1.91 ± 0.06 ^a	2.20 ± 0.24	37.34 ± 1.06 ^a
EZM₅₀+E	69.56 ± 0.84 ^{ab}	18.04 ± 0.84 ^a	5.87 ± 0.25 ^b	4.41 ± 0.47	0.59 ± 0.04 ^b	2.13 ± 0.06 ^a	2.25 ± 0.32	39.09 ± 0.77 ^a
EZM₁₀₀	72.86 ± 1.97 ^{ab}	14.59 ± 1.52 ^b	4.30 ± 0.19 ^c	5.62 ± 0.91	0.61 ± 0.02 ^{ab}	2.05 ± 0.63 ^a	2.64 ± 0.77	19.46 ± 0.62 ^b
EZM₁₀₀+E	71.21 ± 2.73 ^{ab}	14.42 ± 1.16 ^b	4.96 ± 0.52 ^c	3.96 ± 0.79	0.60 ± 0.06 ^{ab}	1.29 ± 0.04 ^c	2.40 ± 0.24	21.67 ± 2.13 ^b

* See footnotes to Table 1

Values are mean (n = 3) ± standard deviation. Mean values in the same column with different superscript differ significantly (P < 0.05).

Table 4

Growth performance of Yellow perch (*Perca flavescens*) fingerlings fed with the experimental diets for ten weeks.

Treatment*	Initial weight (g)	Final weight (g)	Weight gain (g)	Body mass gain (%)	Metabolic growth rate (gkg ^{0.8} day ⁻¹)
Control	10.96 ± 0.06	39.96 ± 2.42 ^{ab}	29.00 ± 2.41 ^{ab}	264.65 ± 21.88 ^{ab}	8.03 ± 0.38 ^{ab}
S₅₀	11.04 ± 0.07	33.13 ± 2.30 ^{cd}	22.09 ± 2.25 ^{cd}	199.98 ± 19.43 ^{cd}	6.80 ± 0.42 ^c
S₅₀+E	10.91 ± 0.05	36.90 ± 4.88 ^{bc}	25.99 ± 4.65 ^{bc}	237.11 ± 74.87 ^{bc}	7.42 ± 1.49 ^{bc}
S₁₀₀	10.98 ± 0.07	22.04 ± 1.65 ^e	11.06 ± 1.59 ^e	100.71 ± 13.95 ^e	4.24 ± 0.45 ^e
S₁₀₀+E	10.99 ± 0.11	22.51 ± 2.54 ^e	11.52 ± 2.38 ^e	104.85 ± 23.82 ^e	4.35 ± 0.76 ^e
EZM₅₀	11.16 ± 0.14	40.02 ± 0.63 ^{ab}	28.85 ± 0.66 ^{ab}	258.50 ± 7.28 ^{ab}	7.96 ± 0.12 ^{ab}
EZM₅₀+E	11.06 ± 0.13	45.38 ± 1.70 ^a	34.32 ± 1.78 ^a	310.28 ± 18.54 ^a	8.79 ± 0.27 ^a
EZM₁₀₀	10.71 ± 0.14	24.91 ± 0.85 ^e	14.20 ± 1.18 ^e	133.02 ± 15.78 ^{de}	5.16 ± 0.38 ^{de}
EZM₁₀₀+E	11.18 ± 0.12	27.89 ± 1.71 ^{de}	16.72 ± 1.61 ^{de}	149.54 ± 13.48 ^d	5.64 ± 0.35 ^e

* See footnotes to Table 1.

Values are mean (n = 3) ± standard deviation. Mean values in the same column with different superscript differ significantly (P < 0.05).

Figure legends

Fig. 1. (A) Specific growth rate (%/day) and (B) antioxidant enzyme (Glutathione peroxidase) activity in liver of yellow perch (*Perca flavescens*).

Values are mean (n = 3) ± standard deviation. Mean values with different superscript differ significantly (P < 0.05).

Fig. 2. Nutrient utilization parameters (A) feed conversion ratio and protein efficiency ratio, (B) protein productive value and lipid productive value of yellow perch (*Perca flavescens*).

Values are mean (n = 3) ± standard deviation. Mean values with different superscript differ significantly (P < 0.05).

Fig. 3. Digestive enzymes (A) amylase (B) protease and (C) lipase activity in pyloric caeca and intestine of yellow perch (*Perca flavescens*).

Values are mean (n = 3) ± standard deviation. Mean values with different superscript differ significantly (P < 0.05).

Fig. 4. (A) Protein metabolism enzymes (alanine amino transferase, ALT and aspartate transferase, AST) activity in liver, (B) antioxidant enzymes (superoxide dismutase and catalase) activity in liver of yellow perch (*Perca flavescens*).

Values are mean (n = 3) ± standard deviation. Mean values with different superscript differ significantly (P < 0.05).