



Effect of microalga-based diet on oxidative stress enzymes of African catfish, *Clarias gariepinus*

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Abstract Here an indigenously isolated microalgal strain *Ascochloris* spp. cultivated in synthetic medium was evaluated as an aquaculture feed supplement. The daily dietary supplement includes microalgal feed (AF) and commercial diet feed (CF) (as control), respectively. These diets were fed separately to the juvenile *Clarias gariepinus* fishes ($n = 4$) under controlled conditions for an experimental period of 100 days. The protein, glycogen and lipid contents in the muscle extracts were found to be marginally higher in fishes that were fed with CF than AF diet. Similarly, CF fishes showed significantly higher glutathione-*s*-transferase, catalase, superoxide dismutase, and lipid peroxidase activities, except glutathione content. Zero mortality of the fishes with no significant difference in the overall body mass with the two dietary supplements strongly suggests that algal biomass could supplement the requisite nutrients for their metabolic activities. This preliminary investigation helps in exploring algal biomass as a potential alternative feed additive in the aquaculture industry.

Keywords Aquafeed · Aquaculture · Microalgae · Oxidative enzymes

Introduction

Aquaculture industry is one of the rapidly growing sectors recording an annual growth rate of 8.8% in past three decades. Fishmeal and fish oil are the most essential constituents of the aqua feed that contributes to the increasing costs and sustainability of aqua-industry. Commonly fishmeal is obtained by milling and drying of the whole fish and/or its remains, which is mostly governed by the seasonal catches of the fishes. Moreover, fish supply and climate changes are the two major hindering factors for fishmeal production and are threat for long-term sustainability. It was estimated that total marine catches is predicted to decline drastically to 5 million metric ton by the end of year 2020 (FAO 2012, 2018). Globally such alarming need enforces the aqua-industries for finding partial and complete fishmeal replacement. This have led to an extensive use of plant-based nutrient sources such as cereals, oilseed meals, pulses and protein concentrate meals (groundnut, peanut, cottonseed, sunflower seed, palm kernel, coconut, rapeseed, soybean protein concentrate, wheat gluten, corn

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gluten, etc.). Moreover, plant-diets must ensure accurate inclusion levels for metabolic synthesis of essential amino acids in these alternative diets supplemented fishes (Olsen and Hasan 2012). Recently, insects are being used as an excellent diet sources for fishes and are preferred as potential choices for fishmeal replacement. This includes locusts, common houseflies, mosquitoes, superworms, grasshoppers, termites, yellow mealworms, Asiatic rhinoceros beetles, domesticated silkworms and black soldier flies, respectively. However, chitin digestibility and bioaccumulation of insecticides in the fish body poses severe hurdles during these dietary inclusions (Henry et al. 2015).

Recently, microalgae have garnered interest in food and nutrition due to the presence of high amounts of proteins, vitamins, polyunsaturated fatty acids and antioxidants, respectively. Besides, microalgae are used in wide range of applications in renewable energy and nutraceutical sectors due to the generation of value-added products. Microalgae are fast-growing photosynthetic organisms which can produce a theoretical yield of approximately $77 \text{ g m}^{-2} \text{ day}^{-1}$ on dry biomass basis which accounts for $\sim 280 \text{ ton ha}^{-1} \text{ year}^{-1}$. About 5000 metric ton of algal biomass is processed annually in the US alone accounting for a turnover of 1.25 billion USD year (Khan et al. 2018). In contrast to the production and maintenance cost of fishmeal, microalgae require fewer resources and are extremely valuable. Arable land and freshwater are not a pre-requisite for microalgae cultivation as it posses the flexibility to grow in high salinity lands, wastewaters, industrial effluents and a wide variety of mediums rendering high algal yield. Apart from this, microalgal cultivation significantly reduces the greenhouse gas emissions (Park and Lee 2016). Besides, the conventional sources like soya, egg, rice, milk, meat and baker's yeast that are used in fish diets usually comprises up to 47% (w/w) protein content, whereas, microalgal species are known to contain very high protein contents up to 71% (w/w), respectively (Chacon-Lee and Gonzalez-Marino 2010). Along with this, the amino acid contents of almost all the microalgal species are in accordance with the levels present in other food proteins (Becker 2004).

Although to certain extent, there are several reports on the applicability of microalgal biomass as an aqua feed supplement. It is a well known fact that microalgae are regarded as an excellent source of antioxidants, and its consumption helps the aquatic organisms to cope up with the oxidative stress generated due to metabolic and other factors. Cahu et al. (1998) reported that feeding algae to *Dicentrarchus labrax* in larval stage significantly improved the digestive functions in the intestinal and pancreas. Similarly, effects on addition of sea weeds viz., *Spirulina* spp., *Chlorella* spp. and *Azolla* spp. as a partial fish meal replacement on *Macrobrachium rosenbergii* were studied and its non-toxicity on postlarval stages of fishes were reported (Radhakrishnan et al. 2014). Also, products derived from algae like rosamin and verdemin when added to the fish feed has significantly increased omega-3 long chain polyunsaturated fatty acid (PUFA) content in *Salmo salar* (Norambuena et al. 2015). Other than aquatic animals, cross-bred dairy goats supplemented with *Chlorella vulgaris* exhibited progressive formation of *trans*-11 C18:1 and monounsaturated fatty acids (MUFA) compared to the control diets (Tsiplakou et al. 2017). Furthermore, anti-proliferative effects of bioactive carotenoids extracted from *Chlorella ellipsoidea* and *Chlorella vulgaris* on human colon cancer cells were reported suggesting microalgae as a potential therapeutic agent (Cha et al. 2008).

Recently, we have identified an indigenously isolated microalgal strain, *Ascochloris* spp. from dairy effluent contaminated water samples collected from one of the largest dairy industry in India, AMUL (Anand Milk Union Limited), located in Anand district of Gujarat, India. Comprehensive studies on the microalgal growth properties, biomass and lipid productivities were reported earlier (Kumar et al. 2018). Since the acceptability and feasibility of any metabolic diet is determined by physical growth, biochemical and enzymatic indices of an organism, a preliminary assessment of such indices were performed on African catfish, *Clarias gariepinus*, to evaluate whether the *Ascochloris* spp. may be used as a partial replacement of conventional aquafeeds. *Clarias* spp. belongs to the family *Clariidae* and is a popular fish species for aquaculture business in Southeast Asian countries. Also, owing to its high growth rate, disease resistance, fecundity and stress handling, it has a high commercial significance worldwide (Muchlisin and Siti-Azizah 2010).

Materials and methods

All the chemicals and reagents used in the study were of analytical grade and procured from Sigma (St. Louis, MO, USA), Himedia (Himedia Laboratories Pvt Ltd., Mumbai, India) and Loba Chemie (Loba Chemie Pvt.



Ltd., Mumbai, India), respectively. The glass tanks used for maintaining fishes for the experimental period were procured locally.

Microalgae cultivation

All the experiments in the current study were performed using the microalgal strain, *Ascochloris* spp. as the algal feed. The microalga was grown in synthetic TAP (Tris–Acetate–Phosphate) medium under controlled conditions, i.e., light:dark cycle of 16:8 h at a constant temperature of 25 ± 1 °C with light intensity between ranging between 3366 and 3978 W/m² for a period of 5–17 days in a growth chamber (Supplementary Fig. 2c). The microalgal biomass was harvested using centrifugation at $10,000 \times g$ for 10 min and then dried at ≤ 50 °C until the moisture content reached $\leq 5\%$ (Supplementary Fig. 2d, e) (Kumar et al. 2018). In order to avoid minimal cell damage necessary precautions were taken during the entire process of microalgal cultivation to algal feed preparation. Further, microscopic analysis and chlorophyll content in the cell-free supernatant showed no cell disruption (Supplementary Fig. 2a).

In this study, we have analyzed the biochemical and enzymatic responses of *Clarias gariepinus* with two different dietary supplements, i.e., (a) algae fed (AF) and (b) commercial feed fed (CF), respectively. All the fish feeding experiments were carried out for a period of 100 days in the algal laboratory at Bioconversion technology division at Sardar Patel Renewable Energy Research Institute, Gujarat, India. The present study was undertaken with the approval from the institutional ethical committee members. The microalga, *Ascochloris* spp. used as an algal feed in the present study was isolated, morphologically and genetically identified and reported earlier by us (Kumar et al. 2018). Juvenile *Clarias gariepinus* fishes were procured from the local farm and acclimatized in the laboratory for 2 weeks before starting the feeding experiments. All the test fishes were selected based on their similar body weight (160–170 g) and 25–30 cm length, respectively. Fish feed experiments were conducted in 55 L glass water tanks (60 cm \times 30 cm \times 38 cm) with each tank comprising four fishes ($n = 4$). Each tank was provided with an aerator rod attached with a submersible air pump for uniform aeration throughout the tank. Fishes in tank (a) were fed with commercial feed, while tank (b) with alga feed, respectively.

Commercial feed (Make: Optimum aquarium food, Thailand), procured from local vendor, consist of 28% (w/w) crude protein, 3% (w/w) crude fat, 4% (w/w) crude fiber and 10% (w/w) moisture was used as control feed. While, dried algal biomass with moisture content $\pm 5\%$ was used as algal feed. No post-processing of the feed materials was involved. Fishes in both the tanks were fed twice a day with dried algal biomass and commercial feed, respectively, at 2% (w/w) of the fish body weight for a period of 100 days (Supplementary Data 1). The tanks were maintained at pH 8.0–8.5 and temperature 30–35 °C, respectively, by replacing the tank water with nearly 50% (v/v) of clean water every day. Body weight measurements of fishes were taken every fortnight. At the end of the experiment, the fishes were euthanized with Tricaine methane sulphonate (MS-222) followed by decapitation as described by Close et al. (1996). Muscle tissues were carefully excised under sterile conditions from the dorsal region using a sterile scalpel and approximately 1 gm each of sample were taken for pulverization in liquid nitrogen. All the samples were stored at -80 °C for further biochemical analysis.

Compositional analysis: the compositional analysis of the microalgal biomass was evaluated following the NREL laboratory analytical procedure (Laurens 2015). Ash content of the microalgal biomass was determined using dry oxidation method in a muffle furnace at 575 °C. Equation (1) was used for measuring percent ash content where DW_{sample} is the weight of the microalgal biomass on dry basis:

$$\text{Ash\%} = \frac{(\text{weight}_{\text{crucible+ash}} - \text{weight}_{\text{crucible}})}{DW_{\text{sample}}} \times 100. \quad (1)$$

Protein content (% w/w) in the microalgae biomass was determined using Kjeldahl method and was calculated using Eq. (2), where ‘%N’ is elemental nitrogen content and ‘N factor’ is a specific factor determined for algae (4.78).

$$\text{Protein\%} = \% N \times N_{\text{factor}}. \quad (2)$$



Lipid quantification (% *w/w*) was performed using Bligh and Dyer method (1959) where chloroform/methanol/water ratios were maintained at 1:2:0.8, respectively. Total carbohydrate content in the microalgal biomass was determined using two-step acid hydrolysis method where sulphuric acid breakdown the polysaccharides into monosaccharides and were quantified using HPLC (Schimadzu LC 2010C). The % carbohydrate was determined using Eq. (3):

$$\text{Carbohydrate\%} = \left(\frac{\text{total monosaccharides}}{\text{DW}_{\text{sample}}} \right) - \text{starch\%} \times 100. \quad (3)$$

Moisture content (%) in the microalgal biomass was calculated using oven-dry method, where 100 mg of wet alga was taken in a pre-weighed ceramic crucible and dried at 105 °C in a vacuum oven. Finally the % moisture was determined using Eq. (4).

$$\text{Moisture\%} = \frac{(\text{weight}_{\text{wetbiomass}} - \text{weight}_{\text{drybiomass}})}{\text{weight}_{\text{wetbiomass}}} \times 100. \quad (4)$$

Biochemical analysis: protein, glycogen and lipid determination

The tissue extractives from the fishes were obtained by finely crushing 0.5 g of muscle tissue in a specific extraction buffer that was described below for each individual. The tissue slurry was then sonicated for 20 s and centrifuged at 10,000 rpm \times g for 20 min to obtain clarified tissue extract. All the clarified extracts were stored at – 20 °C until further biochemical analysis. The muscle protein and carbohydrates were extracted in sodium phosphate buffer (50 mM, pH 7.0) containing 1 mM dithiothreitol, 1% (*v/v*) Triton-X 100, 1 mM ethylenediaminetetraacetic acid. The protein content was quantified spectrophotometrically using Bradford's method (Bradford 1976). Bovine serum albumin was used as external standard and the absorbance was recorded at 550 nm in microplate reader (Biotek, Synergy HT). Total glycogen content was determined using phenol–sulphuric acid method (Dubois et al. 1956). Dextrose was used as a standard reducing sugar and the absorbance was recorded at 490 nm. Lipid extraction from the muscle tissues was performed following Folch's method (Folch et al. 1957). Briefly, the muscle tissue was crushed in chloroform/methanol (2:1) and the two layers were separated by a separating funnel. The organic layer containing the lipids were collected and subjected to vacuum rotary evaporation method (Heildoph, Hei-vap). The lipid content was determined gravimetrically. Final results were expressed in mg g⁻¹ dry weight of the tissue.

Determination of oxidative stress enzymes

Glutathione (GSH) assay

For glutathione (GSH) determination, the muscle tissue was pulverized in 5% sulfosalicylic acid and the supernatant was separated by centrifugation at 10,000 \times g for 20 min. The GSH activity was estimated by enzymatic recycling method following the protocol described by Akerboom and Sies (1981). Briefly, 1 mL of potassium phosphate buffer (pH 7.0) containing reduced glutathione (0.5–2 nM), NADPH (50 μ L of 4 mg mL⁻¹), 5,5'-dithiobis(2-nitrobenzoic acid) (20 μ L of 1.5 mg mL⁻¹) was added to the muscle tissue extracts and the rate of reaction was recorded at 412 nm over a period of 5 min. Glutathione reductase (20 μ L of 6 U mL⁻¹) was used as control enzyme. The GSH activity was expressed as nmoles mg⁻¹ of protein of the sample.

Glutathione-s-transferase (GST) assay

0.5 g of the muscle tissue was crushed in 0.1 M sodium phosphate buffer, pH 6.0. The GST activity was determined using Habig's method (Habig et al. 1974) with minor modifications. The assay mixture consists of 100 mM 1-chloro, 2,4-dinitrobenzene, 200 mM L-glutathione (reduced) and 10 μ L of the tissue extract. A kinetic measurement for 5 min with an interval of 30 s was recorded at 340 nm ($\epsilon = 9.6$ mM) using UV



visible spectrophotometer (Shimadzu UV–Vis spectrophotometer UV-1700). GST activity was expressed as $\mu\text{mol mL}^{-1} \text{min}^{-1}$ and nmol mg^{-1} of protein sample.

Catalase assay

0.5 g of the muscle tissue was crushed in 50 mM sodium phosphate buffer (pH 7.0) consisting 1 mM EDTA. The catalase activity was determined by Beers and Sizer's method (1952). The assay mixture consists of 30 mM 30% H_2O_2 and 10 μL enzyme extract. Absorbance was immediately recorded at 240 nm for 2 min on UV visible spectrophotometer (Shimadzu UV–Vis 1700 spectrophotometer). The rate of decomposition of H_2O_2 is directly proportional to the catalase activity. One unit of catalase activity is defined as the amount of enzyme required to decompose 1 μM H_2O_2 . The enzyme activity was expressed as U mg^{-1} of protein.

Superoxide dismutase assay

0.5 g of the muscle tissue was crushed in 50 mM potassium phosphate buffer (pH 7.0) containing 0.25 M sucrose. The clear supernatant was collected by centrifugation at $10,000 \times g$ for 20 min and analyzed for superoxide dismutase (SOD) activity using xanthine/xanthine oxidase coupled reaction as described by McCord and Fridovich (1969). The kinetic measurements were recorded at 550 nm for 5 min. The SOD activity was expressed as U mg^{-1} of protein.

Lipid peroxidise assay

0.5 g of the muscle tissue was crushed in 20% trichloroacetic acid using Potter–Elvehjem homogenizer. The extract was then centrifuged at $10,000 \times g$ for 20 min at 4 °C to obtain clear supernatant. Lipid peroxidise activity was determined using a fluorometric assay method described by Ohkawa et al. (1979), where thio-barbituric acid (0.67%) was used as a substrate and 1,1,3,3-tetramethoxypropane as a standard. The chemical adduct was fluorometrically determined in a plate reader at $E_x/E_m = 532 \text{ nm}/553 \text{ nm}$, respectively. The lipid peroxidation levels were expressed in nmoles of malon dialdehyde per dry weight of the tissue.

Statistical analysis

All the experiments were conducted in triplicates and the data are presented as mean \pm SEM. One-way analysis of variance (ANOVA) was carried out using SigmaPlot ver. 14.0 software (SigmaPlot Inc, USA). The significant differences in the means between dietary treatments were evaluated by Tukey's multiple range test. Mean differences were considered significant at a p value of < 0.05 .

Results and discussion

Biochemical analysis

It was reported that inclusion of fresh and marine water-grown microalgae in the fish diet had significantly contributed in improving the healthy nature of the aqua species and also enhanced the immune responses towards oxidative stress (Mustafa et al. 1995; Velasquez et al. 2016). In this study, supplementing the algal biomass as aquafeed had not shown any significant differences in the overall body weight, as compared to the commercial feed, i.e., AF vs CF, respectively (Fig. 1), and is in line with other reported data. However, lower protein content ($9.03 \pm 0.08 \text{ mg g}^{-1}$) and lipid accumulation ($20.78 \pm 0.5 \text{ mg g}^{-1}$) was observed in AF, whereas in CF, significantly higher protein ($13.20 \pm 0.13 \text{ mg g}^{-1}$) and lipid ($24.02 \pm 0.06 \text{ mg g}^{-1}$) contents were recorded (Fig. 2a). One of the possible reasons could be due to the difference in the nature of feeder and the feed. The feeder, *Clarias gariepinus*, is a carnivorous fish, while the feed, i.e., microalgae, is of plant origin. Secondly, in general, it is known that lower digestibility of the dietary fibers is common in carnivorous fishes. The fibrous nature of the feed is mainly due to the carbohydrates of the microalga which accounted for lower feed intake, i.e., $60 \pm 7\%$ in AF as compared to $79 \pm 9\%$ in CF. Supporting our experimental results, it



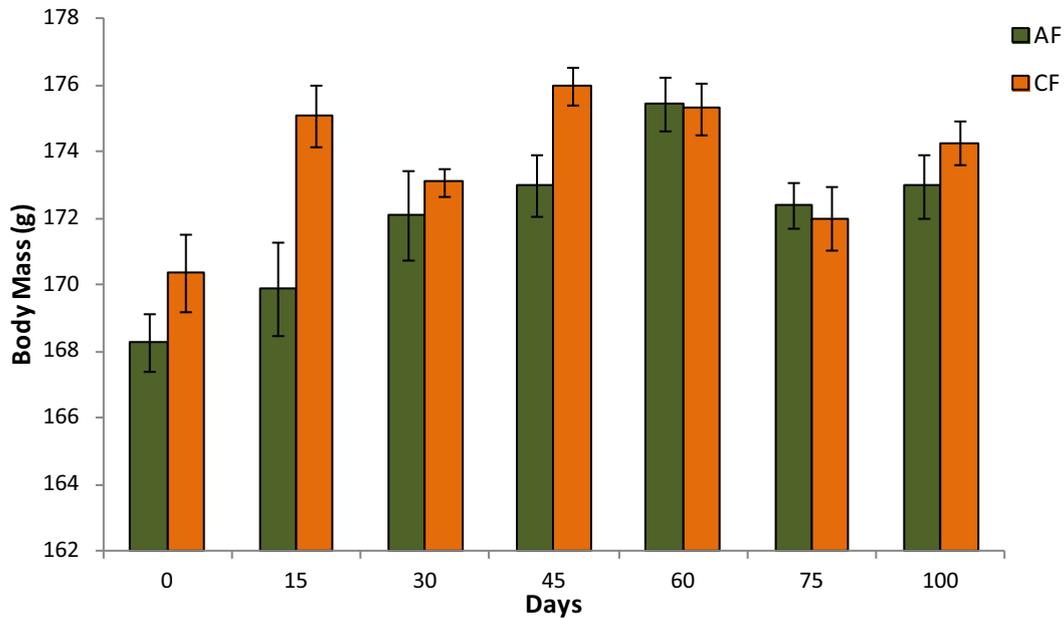


Fig. 1 Comparison of body mass of algae-fed fish (AF) and commercial feed-fed fish (CF) for the test period of 100 days. Body mass measurements were recorded every 15 days

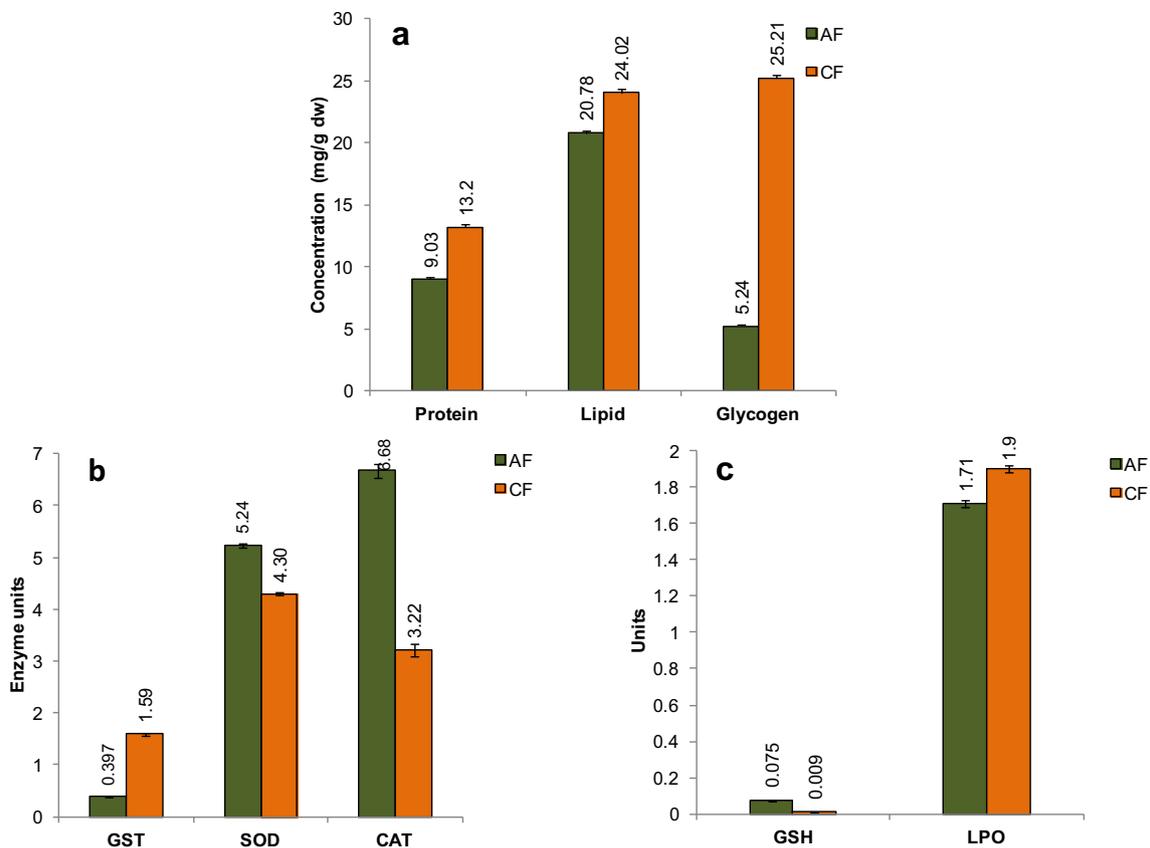


Fig. 2 Biochemical and enzymatic analysis of algae-fed fish AF and commercial feed-fed fish CF. **a** Protein, lipid and glycogen content, **b** enzyme activities, i.e., glutathione-*s*-transferase (GST), catalase (CAT), superoxide dismutase (SOD), **c** non-enzymatic antioxidant activity, i.e., lipid peroxidase (LPO) and glutathione (GSH). All the values are mean \pm SEM ($n = 4$, $p < 0.05$)



was reported that carnivorous fishes have shown poor metabolizing ability when fed with carbohydrate enriched diet (Shimeno et al. 1978). The complexity of algal polysaccharides and anti-nutritional factors viz. tannins, saponins, protease inhibitors, lectins and phytic acid that are found in microalgal strains might have also contributed to the poor growth indices in the fishes (Davies et al. 1997; Francis et al. 2001). Hence, marginally lower muscle protein contents were observed in AF, *Clarias gariepinus*, and correlated to their lower consumption of the algal diet. Our results were also in line with other reported data where lower protein and lipid contents in alga fed fishes were recorded, like in Nile tilapia, *Oreochromis niloticus*, Mozambique tilapia, *Oreochromis mossambicus* and flat fish, *Solea senegalensis* fed on *Hydrodictyon reticulatum*, *Spirulina maxima* and plant protein diet (Soyameal, soya protein concentrate, wheat gluten, etc.), respectively (Appler 1985; Olvera-Novoa et al. 1998; Valente et al. 2011). In another study, poor algal feed utilization, specific growth rate (SGR) and protein efficiency ratio (PER) was observed in when the feed was supplemented with *Ulva lactuca* and was reported recently by Abdel-Warith et al. (2016). Besides, rainbow trout, *Oncorhynchus mykiss* when fed with high carbohydrate diet exhibited stunted growth rate (Edwards et al. 1977). Our studies clearly suggests that microalga-based diet effect the physiological metabolism of *Clarias gariepinus*, especially the protein and lipid pathways, fatty acid biosynthesis, amino acid metabolism, intracellular protein transport and ribosome assembly (Panserat et al. 2008).

Enzymatic analysis

Reduced glutathione

Reduced glutathione (GSH) is one of the major non-enzymatic antioxidants. It plays a significant role in detoxification through chemical and enzymatic reactions (Ross 1988). The total glutathione contents in muscle extracts of AF and CF were 0.075 ± 0.001 nmoles mg^{-1} protein and 0.009 ± 0.001 nmoles mg^{-1} protein, respectively (Fig. 2c). Higher amounts of GSH in AF clearly indicates an effective anti-oxidative nature by *Clarias gariepinus*, whereas lower levels in CF suggests depleted levels of GSH arising primarily due to oxidative stress conditions, thereby conversion of the reduced GSH to oxidative glutathione (GSSG). GSSG mainly involves in secondary reactions like conjugation, reduction, and interaction with other non-enzymatic antioxidants (Forman et al. 2009). It is highly intriguing that supplementing *Ascochloris* spp. as an algal feed could drastically reduces the oxidative stress levels in the aquatic fishes as demonstrated in the current study. Similarly, other studies have reported that elevated GSH levels were observed in muscle and liver of *Sparus aurata* and *Salmo salar* fishes when fed with plant-protein diet (Sitjà-Bobadilla et al. 2005; Olsvik et al. 2011). This clearly suggests that plant-protein diets could be incorporated in the fish meals along with animal-based protein diets. In another study, it was observed that other than aquatic species, supplementation of *Chlorella* sp. and *Spirulina fusiformis* to mammalia (Swiss albino mice) showed elevated ratio of GSH/GSSG, reduced bacterial translocation and enhanced the defense mechanisms (Bedirli et al. 2009; Sharma et al. 2007).

Glutathione-s-transferase (GST)

GST is primarily responsible for detoxification of xenobiotic substances by conjugating it with endogenous reduced glutathione (GSH) (Jancova et al. 2010). GST activity in the muscle tissue extracts was significantly higher in the CF (1.59 ± 0.03 nmoles mg^{-1} protein) than in the AF (0.39 ± 0.02 nmoles mg^{-1} protein), respectively (Fig. 2b). As the microalgal cells were not found to be deformed or disrupted in the cultivation process, (Supplementary Fig. 2a), it was presumed that significant generation of any exogenous compounds were nullified during the microalgal growth and could have directly contributed in elevating the enzymatic response of GST in the AF. In contrary, commercial feed used in this investigation contained minute amounts of anti-oxidants (Astaxanthin), synbiotics, vitamins and minerals (Optimum aquarium food, Thailand). Detailed composition of the algal feed and the commercial feed is given in Table 1. But, it is not clearly known that the anti-oxidants incorporated in the commercial feed had any effect on reducing the generation of redox active substances during the metabolic growth of the fishes. Further molecular understanding on the anti-oxidants present in the *Ascochloris* spp. warrants the potential applicability of the microalga as an aqua feed. Others have reported that no significant difference in the GST levels was recorded in *Clarias gariepinus* and *Dicentrarchus labrax* when fed with fishmeal supplemented with microalgal species such as *Spirulina*

Table 1 Compositional analysis of feeds g% per kg dry matter

	Algal meal	Commercial meal
Moisture	8.56	10
Ash	5	13.56
Protein	20	28
Lipid	11	3
Carbohydrate	6	4
Antioxidants	ND	0.1
Vitamins	ND	2 (vitamin C and E)
Minerals premix	ND	2

ND not determined

platensis, *Chlorella vulgaris*, *Gracilaria* spp., *Ulva* spp., *Fucus* spp., respectively (Raji et al. 2018; Peixoto et al. 2016). However, incorporation of animal protein sources, i.e., cricket meal and housefly maggot meal in the diets of *Clarias gariepinus* and *Oreochromis niloticus* resulted in higher GST levels than the fishmeal diets alone (Ogunji et al. 2007; Taufek et al. 2016). Consequently, the origin of protein source cannot be solely contribute for direct enzymatic reactions of GST and thus further investigation is needed for determining the toxicity levels of fish feed constituents inducing enzymatic responses.

Superoxide dismutase (SOD) and catalase (CAT)

Superoxide dismutase (SOD) functions in dismutation of two molecules of superoxide anion (O_2^-) to molecular oxygen (O_2) and hydrogen peroxide (H_2O_2) (Fridovich 1995), whereas catalase (CAT) further helps in degradation of the generated hydrogen peroxide to water and molecular oxygen (Chance et al. 1952). Both these enzymes represent highest activity against reactive oxygen species and free radicals and reflect a rapid cellular metabolic rate. Kumar et al. (2014) reported that enzymatic and non-enzymatic antioxidant molecules present in microalgal strains scavenge free radicals and protect the muscle cells from oxidative stress when supplemented as fish feed. Higher catalase activity in AF ($6.68 \pm 0.13 \text{ U mL}^{-1}$) than in CF ($3.22 \pm 0.11 \text{ U mL}^{-1}$) clearly indicates the protective role of microalga *Ascochloris* spp. proving it as a viable alternative fish feed supplement along with the commercially available aquafeeds to a large extent. Similar observations were recorded with SOD, where microalga fed *Clarias gariepinus* effectively eradicated free radicals as evident from higher enzymatic response in the AF ($5.24 \pm 0.04 \text{ U mL}^{-1}$) than the CF ($4.30 \pm 0.02 \text{ U mL}^{-1}$), respectively (Fig. 2c). Similar experimental observations were reported by others where olive flounder, *Paralichthys olivaceus*, orange-spotted grouper *Epinephelus coioides* and *Cyprinus carpio* involving dietary *Chlorella vulgaris*, laminarin and *Chlorella* sp., respectively, exhibited similar responses (Rahimnejad et al. 2017; Yin et al. 2014; Stara et al. 2014). Elevated SOD and CAT activities are also reported in Sprague–Dawley rats fed with marine algae, *Pelvetia siliquosa* (Lee et al. 2003). Hence, it can be preliminarily concluded that the microalga fed fishes exhibit better defense mechanism against free radicals as compared to the commercial feed-fed fishes.

Lipid peroxidation

Free radicals are highly unstable molecules responsible for attacking carbon–carbon double bonds of intact lipid molecules, extracting hydrogen and forming peroxy radicals and hydroperoxides. Free radical mechanism is a self proliferating process and is majorly responsible for damaging cells (Yin et al. 2011), however the antioxidant responses nullify such effects of free radicals (Lesser 2006). The amount of malondialdehyde recorded in the CF and the AF were $1.90 \pm 0.01 \text{ nmoles g}^{-1}$ and $1.71 \pm 0.01 \text{ nmoles g}^{-1}$, respectively (Fig. 2c). Lipid peroxidation levels were marginally lower in the AF than in the CF, respectively. This clearly demonstrates that the microalgal antioxidant molecules could successfully mitigate the reactive oxygen species and protect the cells in the AF, whereas the commercial fishmeal had comparatively lower protective effects. This could be due to the degradation of bioactive molecules in commercial fish feed. Hence,



microalga-based diet had enhanced defense mechanisms than the commercial fish meal. Prevention of lipid peroxidation by microalgae was also observed in large yellow croaker, *Pseudosciaena crocea* fed with astaxanthin and *Haematococcus pluvialis* than the control diets without microalga supplementation (Li et al. 2014). Similarly, *Salmo salar* fishes fed with macroalga *Palmaria palmate* also demonstrated lower level of lipid peroxidation than the control diets excluding seaweed (Wan et al. 2016). Besides, supplementation of antioxidants (β -carotene, astaxanthin and lutein) extracted from microalga *Spirulina platensis*, *Haematococcus pluvialis* and *Botryococcus braunii*, respectively, to mammalian species (male Wistar rats) could significantly prevent lipid peroxidation levels by scavenging hydroxyl radicals and free radicals (Ranga Rao et al. 2010).

Statistical analysis for all the experimental parameters performed was found with *p* values much less than 0.05 (level of significance) (Supplementary data 2) which is a strong evidence against null hypothesis. Therefore the null hypothesis is rejected proving that there is a significant difference between the algal feed and commercial feed on various parameters of the specimen under study.

Conclusion

Based on the biochemical and enzymatic responses of GSH, GST, SOD, CAT and LPO with *Ascochloris* spp., it clearly suggests that the microalga could be a prospective candidate in aquafeed formulations. However, other than the aquafeed, the acceptability of microalga diet by the fishes also depends on several collective factors such as palatability, absorption of nutrients, antioxidant potential, voluntary feed intake by test animals, age and stress physiology of animals, respectively. This study presents a preliminary assessment of the microalgal strain, *Ascochloris* spp. as an aquafeed supplement. Further investigation determining the optimum constitution of the alga biomass viz., powdery and granular, in the fish diets formulations aided with binders and other animal protein constituents will reveal the applicability of the microalgal species as a potential aquafeed supplement. Also, cellular and molecular investigations further warrant the physiological and metabolic adaptation of the fresh-water and marine-water fishes towards establishing microalgae as a potential supplement of the commercial aquafeeds.

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Author contributions AKK is a principal scientist and he has designed, supervised the research work and performed critical refinement of manuscript. SS is an associate scientist responsible for large-scale outdoor microalgae cultivation and dissection and sampling of experimental fishes. DD is a research fellow and has performed the fish feeding, enzymatic experiments, data analysis. GD is a research fellow responsible for large-scale outdoor microalgae cultivation in photo bioreactors, raceway pond and harvesting algal biomass. ES and AP are scientific technical assistants and have performed the microalgae growth, enrichment and culturing experiments, species identification and lipid profiling. AK, DD and SS have written the manuscript. All the authors have read and approved the final manuscript.

Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

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