

Effects of dietary tea polyphenols on growth, biochemical and antioxidant responses, fatty acid composition and expression of lipid metabolism related genes of large yellow croaker (*Larimichthys crocea*)

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Abstract

The study was conducted to investigate the effects of dietary tea polyphenols (TP) on growth performance, biochemical and antioxidants responses, fatty acid composition, and lipid metabolism-related gene expressions of large yellow croaker (*Larimichthys crocea*). Four diets were formulated with different levels of TP (0.00%, 0.01%, 0.02% and 0.05%). Results showed that growth performance of *L. crocea* were not different among dietary treatments. Compared with the control group, fish in 0.02% TP group had lower body and hepatic lipid content and lower total cholesterol content. The minimum content of triglycerides and low-density lipoprotein-cholesterol were found in 0.05% TP group. Hepatic n-6 PUFA and n-3 PUFA were significantly higher in TP supplementation groups. Malondialdehyde content was lower in TP supplementation groups, and superoxide dismutase activity was higher in 0.01% TP group than the control group. The mRNA expressions of carnitine palmitoyltransferase1, acyl-CoA oxidase and peroxisome proliferators-activated receptor α were up-regulated in 0.01% and 0.02% TP groups, while lipoprotein lipase expression was down-regulated in TP supplementation groups than the control group. Results suggested that 0.01%–0.02% TP supplementation could reduce the deposition of liver lipid of *L. crocea* caused by high-lipid diet, which might be due to the increase in lipid oxidation related gene expressions.

KEYWORDS

antioxidant capacity, fatty acid composition, gene expression, large yellow croaker, tea polyphenols

1 | INTRODUCTION

High lipid diets (HLD) has been increasingly used for cost-effective farming and protein sparing effect in aquaculture in recent years, indicating that HLD may be more helpful for improving protein utilization and maximizing nitrogen retention in fish (Boujard et al., 2004). However, HLD often lead to abnormal lipid accumulation in

the liver with the low disease resistance and hepatocyte injury of farmed fish (Ai et al., 2004; Du et al., 2008), which is harmful to the health and production of fish (Cabello, 2006). More researches are needed to explore a way to reverse the negative effects of HLD.

Tea polyphenols (TP), the main constituent of tea, have been reported to have various pharmacological and biological functions,

such as antioxidant (Benzie & Szeto, 1999), antibacterial and antiviral (Hattori, Kusumoto, Namba, Ishigami & Hara, 1990; Nakayama et al., 1993), anti-obesity (Murase, Nagasawa, Suzuki, Hase & Tokimitsu, 2002), and hypocholesterolemic activities (Ikeda et al., 2003). Besides, studies of lipid metabolism in animals and cells have found that TP reduced the contents of triacylglycerol, total cholesterol and low-density lipoprotein-cholesterol (Chan et al., 1999; Nanjo, Hara & Kikuchi, 1994), inhibited body and hepatic fat accumulation (Chaudhari & Hatwalne, 1977; Ishigaki, Tonooka, Matsumoto & Hara, 1991). Numerous studies have been carried out to investigate the medicinal properties and health benefits of TP on mammals. However, few reports have focused on the effects TP supplementation on the growth and lipid metabolism in fish.

Large yellow croaker (*Larimichthys crocea*), with its high level of production and delicious taste, has been widely cultured in south-east China. Furthermore, previous studies found that the lipid metabolism of the large yellow croaker is similar to other fish species and mammals, and could be a suitable research model for lipid metabolism in marine fish (Cai, Feng, Xiang, Mai & Ai, 2016; Yan et al., 2015; Zuo et al., 2012). Thus, this study was carried out to determine the effects of the supplementation of TP on growth, antioxidant capacity and lipid deposition in the liver of large yellow croaker fed HLD.

2 | MATERIALS AND METHODS

2.1 | Feed ingredients and diet formulation

Four isoproteic (43% crude protein) and isolipidic (18% crude lipid) diets were formulated to contain graded levels of TP (0.00%, 0.01%, 0.02% and 0.05%) (Table 1). Previous studies found that 12% dietary lipid content was an appropriate lipid content for large yellow croaker (Yi et al., 2014) and 18% dietary lipid content was a high lipid content for large yellow croaker (Wang, Yan, Xu, Ai & Mai, 2016; Yan et al., 2015). White fish meal, wheat gluten meal, casein, wheat meal and soybean meal were used as protein sources, whereas fish oil and soybean oil were used as the main lipid sources. Ingredients and nutrient composition of the four experimental diets are given in details (Table 1).

Dietary ingredients were ground into the fine powder through 320 μm mesh and weighed. All the ingredients were thoroughly mixed until they were homogenous in the mixer. After that, oil mixture and water were thoroughly mixed with all ingredients respectively. Pellets (4 \times 5 mm and 5 \times 5 mm) were made by automatic fish granulator (F-26, South China University of Technology), placed into an oven at 55°C for drying 12 hr, sealed in bags, and stored frozen (-20°C) in the feeding trial.

2.2 | Experimental procedure

Large yellow croaker was bought from the Fu Fa Aquatic Products, Ningde, China. Before the start of the experiment, the same batch

TABLE 1 Formulation and proximate analysis of the experimental diets (% dry weight)

Ingredient	Tea polyphenols level (% dry weight)			
	0	0.01	0.02	0.05
White fish meal	29.00	29.00	29.00	29.00
Wheat gluten meal	8.00	8.00	8.00	8.00
Casein	8.00	8.00	8.00	8.00
Wheat meal	15.50	15.49	15.48	15.45
Soybean meal	18.00	18.00	18.00	18.00
Fish oil	8.00	8.00	8.00	8.00
Soybean oil	6.00	6.00	6.00	6.00
Phospholipid	2.00	2.00	2.00	2.00
Mineral premix	2.00	2.00	2.00	2.00
Vitamin premix	2.00	2.00	2.00	2.00
Attractant	0.30	0.30	0.30	0.30
Sodium alginate	1.00	1.00	1.00	1.00
Ethoxyquin	0.10	0.10	0.10	0.10
Choline chloride	0.10	0.10	0.10	0.10
Tea polyphenols	0.00	0.01	0.02	0.05
(Total)	100.00	100.00	100.00	100.00
Proximate analysis ($n = 3$)				
Crude protein %	42.57	42.78	43.01	42.89
Crude Lipid %	17.78	17.58	17.32	17.76

All those ingredients were supplied by Great Seven Biotechnology Co., Ltd, China.

Mineral premix (mg or g kg⁻¹diet): CuSO₄·5H₂O, 10 mg; Na₂SeO₃ (1%), 25 mg; ZnSO₄·H₂O, 50 mg; CoCl₂·6H₂O (1%), 50 mg; MnSO₄·H₂O, 60 mg; FeSO₄·H₂O, 80 mg; Ca(IO₃)₂, 180 mg; MgSO₄·7H₂O, 1,200 mg; zeolite, 18.35 g.

Vitamin premix (mg or g kg⁻¹diet): vitamin D, 5 mg; vitamin K, 10 mg; vitamin B12, 10 mg; vitamin B6, 20 mg; folic acid, 20 mg; vitamin B1, 25 mg; vitamin A, 32 mg; vitamin B2, 45 mg; pantothenic acid, 60 mg; biotin, 60 mg; niacin acid, 200 mg; α -tocopherol, 240 mg; inositol, 800 mg; ascorbic acid, 2,000 mg; microcrystalline cellulose, 16.47 g.

Tea polyphenols: supplied by solar China; Concentration $\geq 95\%$,

Attractant: glycine and betaine.

of fish was reared in floating sea cages (2 \times 4 \times 2 m) and the control diet was fed for 2 weeks to adapt to the experimental conditions and diets.

At the beginning of the experiment, the fish were fasted for 24 hr and weighed. Sixty fish of similar sizes (15.88 \pm 0.12 g) were distributed into 12 sea cages (1 \times 1 \times 2 m), and each diet was randomly allocated to triplicate groups of fish. Fish were fed twice daily at 05:00 and 17:00 for 70 days. During the experimental period, water temperature ranged from 26.5 to 31.0°C, salinity was from 32‰ to 36‰ and dissolved oxygen was approximately 7 mg/L. At the end of the experiment, the fish fasted for 24 hr before harvest. Total number and body weight of fish in each cage were measured for analysis of weight gain (WG), survival rate (SR) and specific growth rate (SGR). The body weight and liver weight were measured for analysis of hepato-somatic index (HSI) and viscera-somatic index (VSI).

2.3 | Chemical analysis of diets and fish

Crude protein, crude lipid, moisture and ash content in the diet and whole body were determined following the procedures of the Association of Official Analytical Chemists (AOAC, 1995). Briefly, the moisture content was determined by drying to constant weight at 105°C, and crude protein content ($N \times 6.25$) was determined by the Kjeldahl method. Crude lipid content was determined by ether extraction using Soxhlet method. The fatty acid composition were analysed using the procedures described by Metcalfe, Schmitz and Pelka (1966) with some modification (Ai et al., 2008; Zuo et al., 2012). Fatty acid methyl esters were identified and quantified by HP6890 gas chromatograph (Agilent Technologies, Santa Clara, California, USA) with the capillary column (007-CW, Hewlett Packard, Palo Alto, CA, USA) and a flame ionization detector.

2.4 | Blood collection

Following the feeding trial, fish were fasted for 24 hr, then anaesthetized with MS222 (1:10,000) (Shanghai Reagent, China). Blood samples were obtained from the caudal vein with 1 ml syringes of 10 fish from each cage and allowed to clot in the cold for 6-8 hr. The clot was removed and residual blood cells were separated from the serum by centrifugation (3,500 r/min for 10 min). The serum stored at -80°C for later analysis.

2.5 | Plasma biochemical parameters

Aspartate transaminase (AST), alanine transaminase (ALT), total cholesterol (TC), triglycerides (TG), low-density lipoprotein-cholesterol (LDL-c) and high-density lipoprotein-cholesterol (HDL-c) are tested by the Mindry BS-180 automatic biochemical analyzer (Mindry China).

2.6 | Antioxidant index

Malondialdehyde (MDA), total antioxidant capacity (T-AOC) and Superoxide dismutase (SOD) were measured using a test kit (Nanjing Jiancheng Bioengineering Institute China). The reagent preparation, sample pretreatment and method of operation are in accordance with operating instructions.

2.7 | RNA extraction and real-time quantitative polymerase chain reaction (RT-qPCR)

Total RNA was extracted from liver using Trizol Reagent (Invitrogen, USA) and electrophoresed on a 1.2% denaturing agarose gel to test the integrity. Then, RNA reverse transcribed to cDNA by Prime Script-RT reagent Kit (Takara, Japan). RT-PCR was performed in a quantitative thermal cycler (CFX96™ Real-Time System, BIO-RAD, USA). The amplification was performed in a total volume of 25 µl, containing 1 µl of each primer, 1 µl cDNA product, 12.5 µl SYBR-Premix ExTaqII (Takara, Japan) and 9.5 µl RNA-free water. The real-time PCR program was as follows Zuo et al. (2012). The primer sequences of palmitoyltransferase1(CPT1), acyl-CoA oxidase (ACO), lipoprotein lipase (LPL) and peroxisome proliferators-activated receptor α (PPAR α) were from Cai et al. (2016) and Yan et al. (2015) and listed in Table 2. To calculate the expression of CPT 1, ACO, LPL and PPAR α , the comparative ct method ($2^{-\Delta\Delta t}$ method) was used as described by Yao et al. (2009).

3 | CALCULATIONS AND STATISTICAL ANALYSIS

$$\text{Weight gain (WG\%)} = 100 \times (\text{Wt} - \text{Wo}) / \text{Wo}$$

$$\text{Survival rate (SR\%)} = 100 \times \text{FN} / \text{IN}$$

$$\text{Specific growth rate (SGR\%)} = 100 \times (\text{Ln Wt} - \text{Ln Wo}) / t$$

$$\text{Hepato-somatic index (HSI\%)} = 100 \times \text{liver weight} / \text{body weight}$$

$$\text{Viscera-somatic index (VSI\%)} = 100 \times \text{visceral weight} / \text{body weight}$$

where FN is the final number and IN is the initial number of fish in each cage. Wt is the final weight, Wo is the initial weight of fish and t is the experiment period.

The results were presented as means \pm SEM (standard error of the mean). One-way ANOVA was used to analyse all the data. When there were significant differences ($p < .05$), the group means were further compared using Tukey's multiple range test. All statistical analyses were performed using SPSS 19.0 (SPSS, IL, USA).

4 | RESULTS

4.1 | Survival rate and growth

The survival rate was independent of the dietary treatments ($p > .05$). Although no significant differences were observed in

TABLE 2 Sequences of the PCR primers used in this study

Target gene	Forward (5'-3')	Reverse (5'-3')	Reference
CPT1	GCTGAGCCTGGTGAAGATGTTT	TCCATTGTTGTAATTGTTTACTGTCC	Yan et al. (2015)
ACO	AGTGCCAGATGATCTTGAAGC	CTGCCAGAGTAACCATTTCCT	Yan et al. (2015)
LPL	GAATTCAACGCGAAACACAG	ACGCTCATAGAGGGCAGACAC	Yan et al. (2015)
PPAR α	GTCAAGCAGATCCAGAACCC	TGGTCTTCCAGTGAGTATGAGCC	Cai et al. (2016)
β -actin	CTACGAGGGTTATGCCCTGCC	TGAAGGAGTAACCGCGCTCTGT	Yan et al. (2015)

CPT1, carnitine palmitoyltransferase1; ACO, acyl-CoA oxidase; LPL, lipoprotein lipase; PPAR α , peroxisome proliferators-activated receptor α .

TABLE 3 Growth response and survival of large yellow croaker fed diets with graded levels of tea polyphenols level (Means \pm SEM)

	Tea polyphenols level (% dry weight)			
	0	0.01	0.02	0.05
Initial body weight g	15.89 \pm 0.02	15.89 \pm 0.23	15.88 \pm 0.09	15.87 \pm 0.13
Final body weight g	36.71 \pm 3.05	37.96 \pm 2.45	37.53 \pm 0.37	36.80 \pm 0.50
Weight gain %	131.02 \pm 18.88	138.98 \pm 17.39	136.52 \pm 14.24	131.77 \pm 2.37
Specific growth rate %	1.21 \pm 0.07	1.26 \pm 0.05	1.24 \pm 0.04	1.20 \pm 0.02
Survival rate %	86.67 \pm 1.67	87.78 \pm 2.55	85.00 \pm 1.67	85.44 \pm 0.96
Hepato-somatic index %	2.96 \pm 0.31	2.92 \pm 0.39	3.03 \pm 0.12	2.93 \pm 0.26
Viscera-somatic index %	8.53 \pm 0.32	8.04 \pm 0.20	7.98 \pm 0.16	8.15 \pm 0.28

Data are expressed as means \pm SEM. Different letters in each row show significant differences among dietary treatments by Tukey's test ($p < .05$). SEM, standard error of means. ($n = 3$).

growth among dietary treatments ($p > .05$), growth had increased with dietary TP increasing from 0.01% to 0.02% and thereafter decreased (Table 3). There were no significant difference in HSI and VSI among dietary treatments ($p > .05$).

4.2 | Whole body composition

The lipid content in whole body and liver were decreased with dietary TP increasing from 0.01% to 0.02% and thereafter increased (Table 4). In whole body and liver, fish fed the diet with 0.02% TP had the lowest lipid ($p < .05$), while muscle lipid content was not significantly different among dietary treatments ($p > .05$). No significant differences were observed in crude protein, the moisture of whole body among dietary treatments ($p > .05$).

4.3 | Serum biochemical indexes

The content of TG, TC and LDL-c decreased whereas the HDL-c content increased with TP increasing from 0.01% to 0.05% (Figure 1). Fish fed the diet with 0.02% TP had significantly lower TC content compared with control group ($p < .05$), and also had lower content of TG and LDL-c compared with control group without significant differences ($p > .05$). The content of TG and LDL-c was significantly lower in fish fed the diet with 0.05% TP compared with control group ($p < .05$). On the contrary, the HDL-c content was significantly higher in 0.02% TP supplementation group compared with

control group ($p < .05$). Compared with the control group, the activity of AST and ALT were significantly lower in 0.02%TP group and 0.01% TP group respectively ($p < .05$).

4.4 | Fatty acid composition

The hepatic saturated fatty acids (SFA) gradually decreased whereas the n-6 PUFA and n-3 PUFA of the liver increased by dose-dependent with the dietary TP increasing from 0.01% to 0.05% (Table 5), but there were no significant differences in SFA, MUFA, n-3 PUFA, n-6 PUFA and other fatty acid in the muscle among treatment groups ($p > .05$) (Table 6). In the liver, the SFA and stearic acid were significantly lower in fish fed the diet with 0.01% TP compared with the control group ($p < .05$) and the lowest SFA appeared in 0.05% TP group ($p < .05$). The lowest C20:0 was observed in 0.05% TP group ($p < .05$). The C18:1n-9 was significantly lower in 0.02% TP group compared with control group ($p < .05$). The n-6 PUFA and C18:2n-6 were significantly higher in fish fed the diet with 0.02% and 0.05% TP compared with the control group ($p < .05$). The n-3 PUFA, C18:3n-3 and C20:3n-3 were significantly higher in 0.02% TP supplementation group compared with control group ($p < .05$). Although there was an upward trend of n-3 LC-PUFA, no significant difference was found among treatment groups ($p > .05$). No significant differences were observed in n-3 PUFA/n-6 PUFA, MUFA and other fatty acid among treatment groups ($p > .05$).

TABLE 4 Effects of tea polyphenols on body composition of large yellow croaker (Means \pm SEM)

Index	Tea polyphenols level (% dry weight)			
	0	0.01	0.02	0.05
Moisture (%)	72.22 \pm 1.62	72.11 \pm 0.79	71.46 \pm 0.31	71.84 \pm 0.64
Crude protein (%w.w.)	15.14 \pm 0.46	15.23 \pm 0.54	15.81 \pm 0.25	15.33 \pm 0.14
Crude lipid (% w.w.)	9.36 \pm 0.17 ^b	8.89 \pm 0.17 ^{ab}	8.55 \pm 0.32 ^a	8.80 \pm 0.35 ^{ab}
Liver lipid content (% w.w.)	39.91 \pm 0.78 ^c	38.42 \pm 1.32 ^c	33.90 \pm 0.30 ^a	35.87 \pm 0.33 ^b
Muscle lipid content (% w.w)	9.58 \pm 0.75	9.12 \pm 0.97	9.35 \pm 1.46	9.16 \pm 0.50

Data are expressed as means \pm SEM. Different letters in each row show significant differences among dietary treatments by Tukey's test ($p < .05$). SEM, standard error of means. ($n = 3$). w.w. means wet weight.

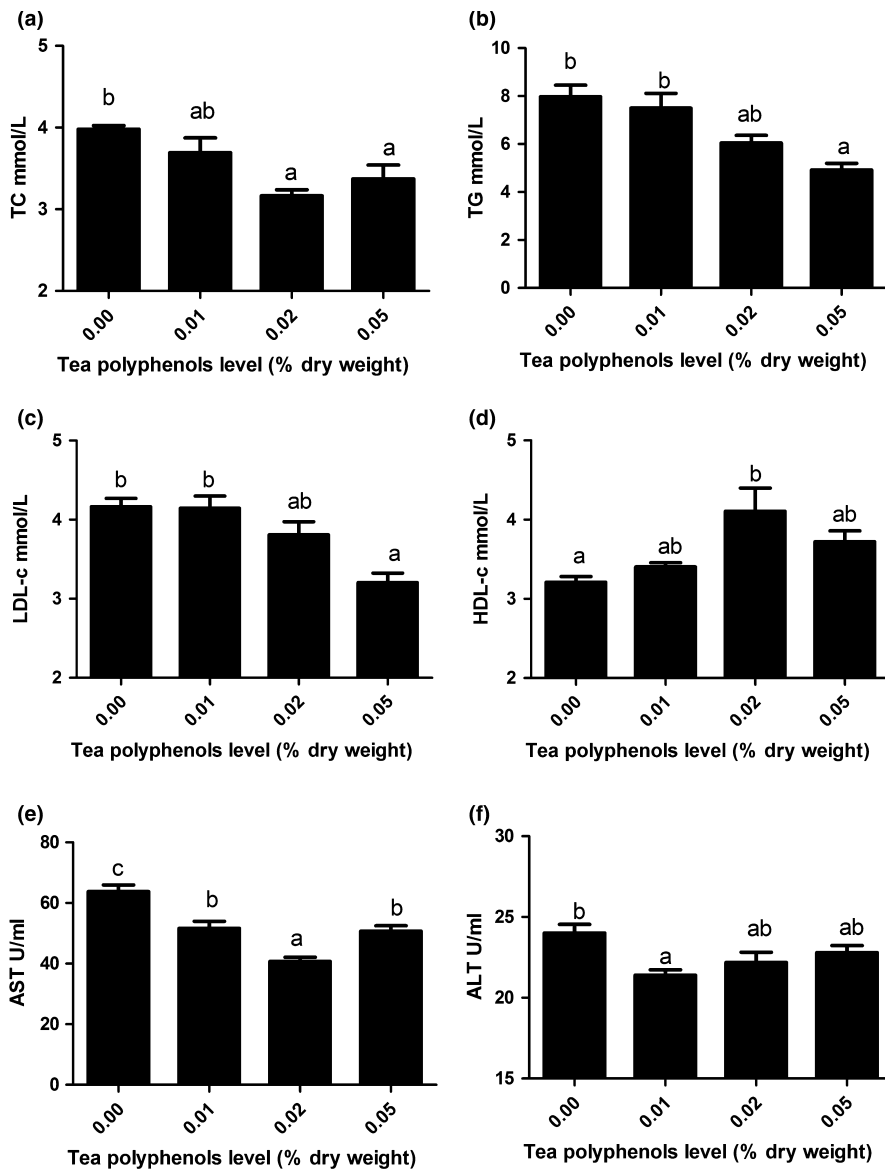


FIGURE 1 The serum biochemical indexes (a) TC. (b) TG. (c) LDL-c. (d) HDL-c. (e) AST. (f) ALT of large yellow croaker fed the experimental diets. Results are expressed as means + SEM ($n = 3$). Bars bearing with different letters are significantly different by Tukey's test ($p < .05$)

4.5 | Antioxidant indexes

Malondialdehyde content was significantly lower in TP supplementation groups compared with the control group ($p < .05$) (Figure 2a), and fish fed the diet with 0.02% TP had the lowest MDA ($p < .05$). The activity of SOD was higher in TP supplementation groups compared with the control group. Especially, fish fed the diet with 0.01% TP was significantly higher than the control group ($p < .05$) (Figure 2c). Also, fish fed the diet with 0.02% TP had significantly higher T-AOC compared with the control group ($p < .05$) (Figure 2b).

4.6 | Lipid metabolism related gene expression in the liver

The mRNA expression levels of CPT1, PPAR α and ACO significantly increased in fish fed diets with 0.01% and 0.02% TP ($p < .05$) and thereafter significantly dropped when dietary TP increased to 0.05%

compared with fish fed the diet with 0.02% TP ($p < .05$) (Figure 3a, b,c). The mRNA expressions of LPL in TP treatment groups were significantly lower than the control group ($p < .05$) (Figure 3d).

5 | DISCUSSION

In this study, although no significant differences were observed in growth performance of large yellow croaker among dietary treatments, growth had been increased with supplementation of TP increasing from 0.01% to 0.02%. Previous studies in mammals have found that TP or tea extracts can significantly reduce body weight (Bose et al., 2008; Tian et al., 2013; Wolfram, Wang & Thielecke, 2006). The differences were probably due to the different animal species used.

Results of this study found that 0.02% dietary TP significantly decreased lipid content of whole body and liver which is consistent

TABLE 5 Effects of tea polyphenols on fatty acid composition in the liver of large yellow croaker (Means \pm SEM) (% of total fatty acids)

Fatty acid (%)	Tea polyphenols level (% dry weight)			
	0	0.01	0.02	0.05
C14:0	2.97 \pm 0.24	3.02 \pm 0.10	3.12 \pm 0.19	3.32 \pm 0.15
C16:0	17.00 \pm 0.94	16.45 \pm 0.47	15.64 \pm 0.13	15.90 \pm 0.93
C18:0	6.22 \pm 0.60 ^c	5.30 \pm 0.17 ^b	4.87 \pm 0.30 ^b	3.82 \pm 0.39 ^a
C20:0	1.21 \pm 0.06 ^b	1.23 \pm 0.05 ^b	1.18 \pm 0.05 ^b	1.09 \pm 0.02 ^a
Σ SFA	27.40 \pm 0.58 ^c	26.01 \pm 0.35 ^b	24.82 \pm 0.42 ^a	24.14 \pm 0.69 ^a
C16:1n-7	7.82 \pm 0.78	7.90 \pm 0.39	7.60 \pm 0.21	8.60 \pm 0.59
C18:1n-9	20.32 \pm 0.83 ^b	19.55 \pm 0.65 ^{ab}	18.97 \pm 0.47 ^a	19.35 \pm 0.47 ^{ab}
Σ MUFA	28.14 \pm 1.34	27.45 \pm 0.68	26.57 \pm 0.44	27.95 \pm 1.04
C18:2n-6	23.64 \pm 1.36 ^a	24.14 \pm 0.27 ^{ab}	25.69 \pm 0.10 ^b	25.78 \pm 0.95 ^b
C20:4n-6	0.53 \pm 0.03	0.54 \pm 0.04	0.57 \pm 0.02	0.54 \pm 0.03
Σ n-6 PUFA	24.17 \pm 1.33 ^a	24.67 \pm 0.31 ^{ab}	26.26 \pm 0.08 ^b	26.32 \pm 0.99 ^b
C18:3n-3	2.27 \pm 0.09 ^a	2.35 \pm 0.08 ^a	2.48 \pm 0.02 ^b	2.56 \pm 0.07 ^b
C20:5n-3	2.89 \pm 0.12 ^a	3.03 \pm 0.22 ^{ab}	3.32 \pm 0.15 ^c	3.22 \pm 0.08 ^{bc}
C22:6n-3	2.34 \pm 0.03	2.31 \pm 0.20	2.45 \pm 0.16	2.24 \pm 0.28
Σ n-3 PUFA	7.50 \pm 0.20 ^a	7.68 \pm 0.48 ^{ab}	8.26 \pm 0.31 ^b	8.01 \pm 0.39 ^{ab}
n-3/n-6 PUFA	0.31 \pm 0.02	0.31 \pm 0.02	0.32 \pm 0.02	0.31 \pm 0.01
Σ n-3 LC-PUFA	5.23 \pm 0.13	5.34 \pm 0.41	5.78 \pm 0.29	5.45 \pm 0.35

Data are expressed as means \pm SEM. Different letters in each row show significant differences among dietary treatments by Tukey's test ($p < .05$). SEM, standard error of means. ($n = 3$).

SFA, saturated fatty acids; MUFA, mono-unsaturated fatty acids; n-6 PUFA, n-6 poly-unsaturated fatty acids; n-3 PUFA, n-3 poly-unsaturated fatty acids; LC-PUFA, long chain-polyunsaturated fatty acids.

TABLE 6 Effects of tea polyphenols on fatty acid composition in the muscle of large yellow croaker (Means \pm SEM) (% of total fatty acids)

Fatty acid (%)	Tea polyphenols level (% dry weight)			
	0	0.01	0.02	0.05
C14:0	3.88 \pm 0.42	3.81 \pm 0.23	3.81 \pm 0.50	3.59 \pm 0.26
C16:0	18.59 \pm 0.79	18.08 \pm 0.50	18.62 \pm 1.75	17.90 \pm 0.27
C18:0	4.64 \pm 0.51	4.81 \pm 0.02	5.06 \pm 0.14	4.81 \pm 0.33
C20:0	1.17 \pm 0.14	1.29 \pm 0.19	1.15 \pm 0.20	1.19 \pm 0.04
Σ SFA	28.28 \pm 1.03	27.99 \pm 0.70	28.63 \pm 2.27	27.49 \pm 0.19
C16:1n-7	6.62 \pm 0.49	6.51 \pm 0.24	6.68 \pm 0.69	6.17 \pm 0.28
C18:1n-9	16.93 \pm 0.26	17.17 \pm 0.33	17.45 \pm 0.23	17.04 \pm 0.40
Σ MUFA	23.55 \pm 0.74	23.69 \pm 0.52	24.13 \pm 0.84	23.21 \pm 0.22
C18:2n-6	22.68 \pm 1.59	22.56 \pm 1.29	22.82 \pm 1.23	23.00 \pm 0.26
C20:4n-6	0.70 \pm 0.13	0.77 \pm 0.88	0.77 \pm 0.03	0.73 \pm 0.03
Σ n-6 PUFA	23.38 \pm 1.46	23.33 \pm 1.33	23.59 \pm 1.22	23.73 \pm 0.28
C18:3n-3	2.30 \pm 0.23	2.27 \pm 0.08	2.23 \pm 0.11	2.33 \pm 0.14
C20:5n-3	4.63 \pm 0.32	4.41 \pm 0.09	4.25 \pm 0.21	4.46 \pm 0.16
C22:6n-3	4.62 \pm 0.42	4.24 \pm 0.36	4.28 \pm 0.07	4.17 \pm 0.12
Σ n-3 PUFA	11.55 \pm 0.72	10.92 \pm 0.37	10.75 \pm 0.33	10.96 \pm 0.18
n-3/n-6 PUFA	0.50 \pm 0.02	0.47 \pm 0.03	0.46 \pm 0.03	0.46 \pm 0.01

Data are expressed as means \pm SEM. Different letters in each row show significant differences among dietary treatments by Tukey's test ($p < .05$). SEM, standard error of means. ($n = 3$).

SFA, saturated fatty acids; MUFA, mono-unsaturated fatty acids; n-6 PUFA, n-6 poly-unsaturated fatty acids; n-3 PUFA, n-3 poly-unsaturated fatty acids.

with studies in mammals that TP has the function of lipid-lowering and anti-obesity effects (Bursill & Roach, 2007; Tian et al., 2013; Wolfram et al., 2006). Correspondingly, the content of plasma TC,

TG and LDL-c were decreased in TP supplementation groups, and 0.02%–0.05% TP supplementation can significantly increase plasma HDL-c content. Furthermore, in this study, TP supplementation

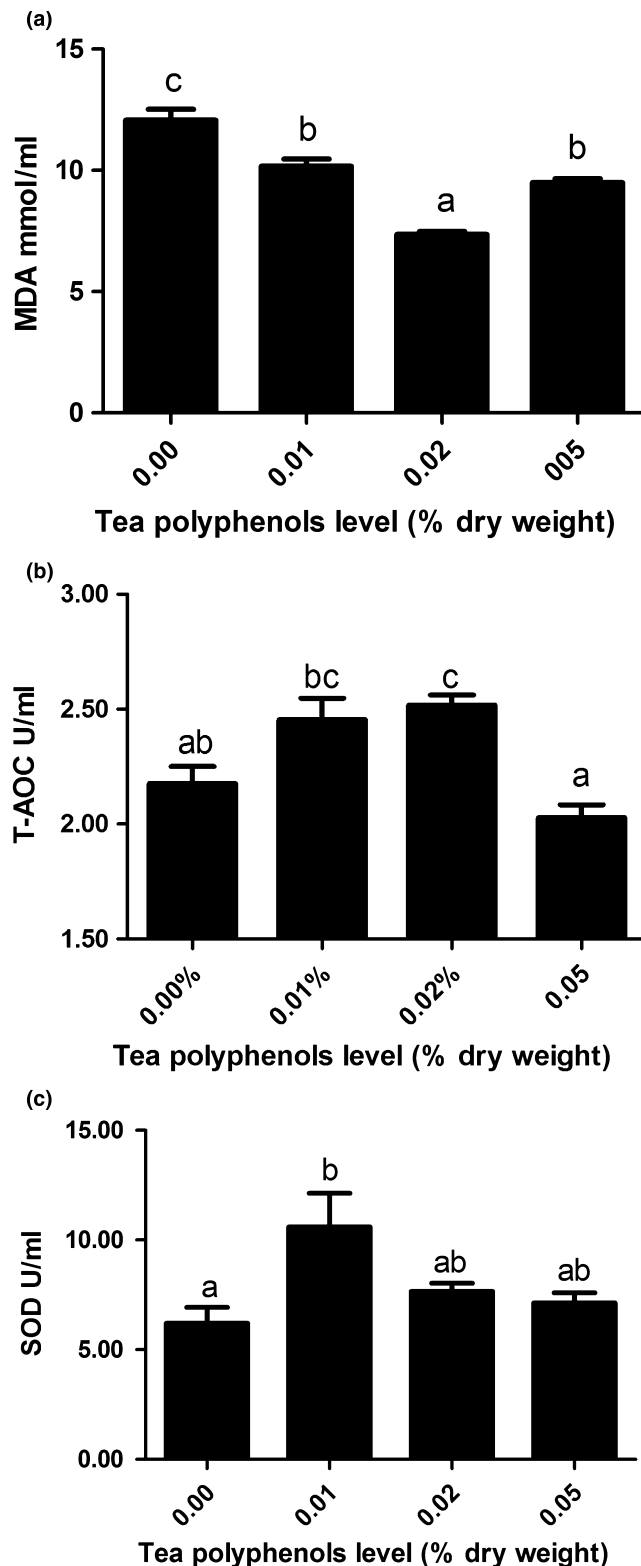


FIGURE 2 Antioxidant indexes (a) MDA. (b) T-AOC. (c) SOD in the serum of large yellow croaker fed the experimental diets. Results are expressed as means + SEM ($n = 3$). Bars bearing with different letters are significantly different by Tukey's test ($p < .05$)

decreased activity of plasma ALT and AST compared with the control group, indicating that TP supplementation could improve liver damage to some extent.

In the liver, with the decrease in hepatic lipid content, SFA significantly decreased, while the n-6 PUFA and n-3 PUFA significantly increased. Studies suggested that SFA and MUFA were more prone to lipolysis in vivo (NRC 2011), and some studies in mammals found that TP can promote the lipid oxidation (Lin & Lin-Shiau, 2006; Murase et al., 2002). Besides, previous studies in mammals have demonstrated that the synthesis of SFA and MUFA were inhibited by TP or tea extract (Yeh, Chen, Chiang, Lin-Shiau & Lin, 2003). The relative content of n-3 PUFA and n-6 PUFA increased with TP increasing from 0.01% to 0.05%, which may be due to the decrease in SFA and MUFA or the absorption of LC-PUFA interfered by TP.

Furthermore, antioxidant capacity and health are related to growth performance. In this study, MDA content was significantly decreased in TP treatment groups compared with the control group. Similar results had been found in humans that TP could decrease MDA content by inhibiting the absorption and accumulation of plasma MDA from the diet (Gorelik, Ligumsky, Kohen & Kanner, 2008). What's more, in this study, T-AOC and the activity of SOD were increased in fish fed diets with 0.01% and 0.02% TP. Therefore, in this study, TP treatment could partly improve hepatocyte injury, the antioxidant activity and health of fish, which may be an explanation for the good growth performance and lower liver lipid content in large yellow croaker.

Previous studies in mammals have shown that tea catechins and EGCG (catechins, epicatechin, epigallocatechin gallate), two of the main components of TP, could restraint the development of fatty liver which induced by HLD and increase the oxidation and lipolysis of lipid (Bose et al., 2008; Murase et al., 2002). However, the lipid metabolism in fish remains to be elucidated, and it is always an interaction of fatty acid transport, synthesis and oxidation which is similar to mammals (Tocher, 2003). LPL involved in hydrolyzing triglyceride and the process of transferring free fatty acids into the cell (Mead, Irvine & Ramji, 2002; Oku, Koizumi, Okumura, Kobayashi & Umino, 2006). In this study, the mRNA expressions of LPL were significantly down-regulated in TP treatment groups compared with the control group, indicating that lipid absorption was reduced. PPAR α has been confirmed to play an important role in increasing oxidation and lipolysis of lipid to reduce lipid accumulation (Goto et al., 2011; Kersten et al., 1999). The activation of PPAR α upregulated gene expressions of CPT1 and ACO, which related to oxidation of fatty acids (Kersten, 2014; Mandard, Müller & Kersten, 2004). Thus, higher CPT1, ACO expression in 0.01% and 0.02% groups may be due to the up-regulation of PPAR α . However, Gene expressions of ACO, CPT1 and PPAR α decreased in fish fed the diet with 0.05% TP treatment group compared with 0.01% and 0.02% TP groups. One possible reason may be that excessive addition of TP may produce a resistance to nutrition, and high doses of caffeine that TP contain inhibited the gene expressions of ACO, CPT1 and PPAR α . Therefore, it could be speculated that dietary TP may improve the growth performance and decreased lipid content in whole body and the liver of large yellow croaker by improving oxidation and lipolysis of lipid.

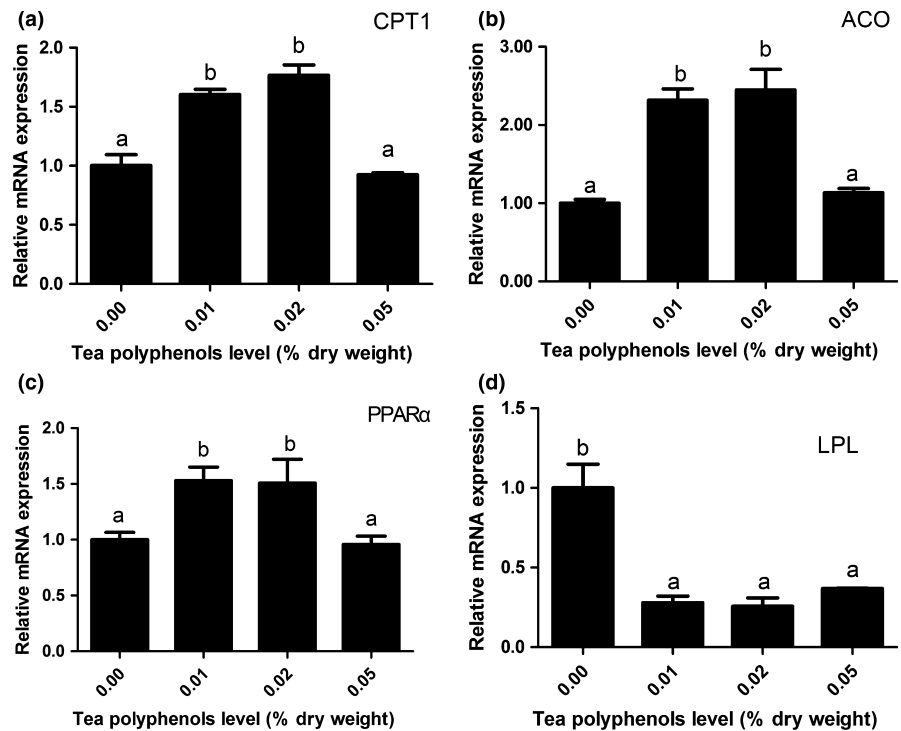


FIGURE 3 Expression of genes related to (a) CPT 1. (b) ACO. (c) PPAR α . (d) LPL in the liver of large yellow croaker fed the experimental diets. Results are expressed as means + SEM ($n = 3$). Bars bearing with different letters are significantly different by Tukey's test ($p < .05$)

In conclusion, results of this study show that 0.01%–0.02% TP supplementation could reduce the deposition of liver lipid of *L. crocea* caused by high-lipid diet, which could be due to the increase in gene expression of lipid oxidation related gene.

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CONFLICT OF INTEREST

There are no conflicts of interest to report.

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