



Effects of dietary terrestrial oils supplemented with L-carnitine on growth, antioxidant capacity, lipid metabolism and inflammation in large yellow croaker (*Larimichthys crocea*)

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Abstract

The present study was conducted to determine the effects of dietary terrestrial oils (TO) supplemented with L-carnitine on growth performance, biochemical and antioxidant response, lipid metabolism and inflammation in large yellow croaker (*Larimichthys crocea*). Three iso-nitrogenous and iso-lipidic experimental diets were formulated with FO (fish oil, the control group), 75 % TO (75 % FO was substituted by the oil mixture with equal amounts of soyabean oil, linseed oil and pork lard) and 75 % TOC (75 % TO supplemented with 800 mg/kg L-carnitine). Compared with the control group, feed efficiency ratio and specific growth rate were significantly increased in fish fed diets with 75 % TO and 75 % TOC. Hepatic lipid content, serum TAG level, LDL-cholesterol level and the mRNA expression of pro-inflammatory genes (*tnfa* and *ifnγ*) were significantly increased in fish fed the diet with 75 % TO compared with the control group. However, the supplementation of 800 mg/kg L-carnitine in the 75 % TO diet repressed hepatic lipid content, serum LDL-cholesterol level and the mRNA expression of *tnfa* and *ifnγ* in fish compared with fish fed the diet with 75 % TO. Total antioxidant capacity, the activity of superoxide dismutase, the mRNA expression of *cpt-1* and the activity of CPT-I were significantly increased in fish fed the diet with 75 % TOC compared with 75 % TO. In conclusion, these results suggested that the supplementation of 800 mg/kg L-carnitine in the diet with TO mixture could increase growth, antioxidant capacity and fatty acid oxidation and decrease the expression of inflammatory genes in large yellow croaker.

Key words: *Larimichthys crocea*: Terrestrial oils: L-Carnitine: Lipid metabolism: Inflammation

The rapidly developing aquaculture industry is accompanied by a growing demand for fish oil (FO)⁽¹⁾. Due to the high price and limited supply of FO, terrestrial oils (TO), which mainly consist of vegetable oil and animal fat, are commonly used to replace FO in aquafeed industry⁽²⁾. Many studies have demonstrated that TO can partially replace FO without significantly decreasing growth performance or feed utilisation of European sea bass (*Dicentrarchus labrax* L.)⁽³⁾, gilt-head sea bream (*Sparus aurata*)⁽⁴⁾, Atlantic salmon (*Salmo salar*)⁽⁵⁾, turbot (*Scophthalmus maximus* L.)⁽⁶⁾ and silvery-black porgy (*Sparidentex basta*)⁽⁷⁾. However, the decrease in growth and non-specific immunity has also been repeatedly reported in fish when dietary FO is excessively replaced by TO, probably due to the unbalanced dietary fatty acid profile^(8,9).

All fish species have an essential demand of both *n*-3 and *n*-6 PUFA⁽¹⁰⁾. Generally, freshwater and diadromous fish species require linoleic acid (LNA) and α -linolenic acid (ALA), whereas marine fish species strictly require long-chain PUFA (LC-PUFA), such as EPA, DHA and arachidonic acid^(11,12). Furthermore, an appropriate ratio of *n*-3:*n*-6 PUFA is important for the growth and health performance of Malabar grouper (*Epinephelus malabaricus*)⁽¹³⁾, salmon^(14,15) and yellow catfish (*Pelteobagrus fulvidraco*)⁽¹⁶⁾. In addition to the need for PUFA, fish also need an appropriate amount of SFA and MUFA, which are good dietary energy sources for fish^(2,17). Animal fat contains a high percentage of SFA and MUFA, and it is a cheap alternative lipid resource for aquafeed industry⁽¹⁸⁾. However, few studies have investigated the effects of a relatively appropriate dietary fatty acid

Abbreviations: *aco*, acyl-CoA oxidase; ALA, α -linolenic acid; *arg-1*, arginase-1; *cd36*, cluster of differentiation 36; *cox-2*, cyclo-oxygenase-2; *cpt-1*, carnitine palmitoyltransferase 1; *fas*, fatty acid synthase; *fatp1*, fatty acid transport protein 1; FO, fish oil; *gapdh*, glyceraldehyde-3-phosphate dehydrogenase; *hl*, hepatic lipase; *ifnγ*, interferon γ ; LNA, linoleic acid; *mtp*, microsomal TAG transfer protein; *sreb1*, sterol-regulatory element binding protein 1; T-AOC, total antioxidant capacity; TO, terrestrial oils.

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profile on growth performance, metabolism and health of fish when dietary FO is replaced by a high level of TO.

Large yellow croaker (*Larimichthys crocea*) is an economically important fish in China. When dietary FO is excessively replaced by TO, inflammatory response and abnormal lipid deposition in large yellow croaker often cause huge economic losses^(19,20). Progress is urgently needed to find effective additives to reduce inflammation and abnormal lipid deposition in fish fed the diet with a high level of TO. L-Carnitine plays a crucial role in shuttling long-chain fatty acids into mitochondria⁽²¹⁾ and reduces abnormal lipid deposition^(22,23). L-Carnitine can also increase the immunocompetence of several fish species to better cope with biotic and abiotic stresses^(24–26). However, few studies have elucidated the effects of a high level of dietary TO supplemented with L-carnitine on fish. Therefore, the purpose of this study was to determine the effects of a high level of dietary TO supplemented with L-carnitine on growth, lipid deposition and inflammation of large yellow croaker for the wide usage of TO in aquafeed industry.

Methods

Animal ethics

The protocols for animal care and handling were conducted in strict accordance with the Management Rule of Laboratory Animals (Chinese order no. 676 of the State Council, revised 1 March 2017). The present study was approved by the Institutional Animal Care and Use Committee of the Ocean University of China.

Diets formulation

Three iso-nitrogenous (42% crude protein) and iso-lipidic (12% crude lipid) experimental diets were formulated, which included FO (the control group), 75% TO (75% FO was substituted by TO, dietary lipid source consisted of equal amounts of FO, soyabean oil, linseed oil and pork lard) and 75% TOC (75% TO supplemented with 800 mg/kg L-carnitine). 75% FO was substituted by TO to preserve an appropriate fatty acid profile in the 75% TO diet. The addition of L-carnitine was based on Sang's⁽²⁷⁾ study. Ingredients were purchased from Qingdao Great Seven Biotechnology Co. Ltd. Each diet was supplemented with vitamin and mineral premixes, an attractant mixture and a mold inhibitor. Procedures for making diets and storage of experimental diets were in accordance with the protocol described by Ai *et al.*⁽²⁸⁾. Ingredients of the three experimental diets (Table 1) and fatty acid profile of oil sources and diets (Table 2) were given in detail.

Fish and experimental procedures

Large yellow croaker juveniles were obtained from Ningde Fufa Fishery Co. Ltd. Before the formal experiment, experimental fish were hand-fed twice daily (05.30 and 17.30 hours) to acclimatise the fish to the diet and environment for 2 weeks. Fish of disease-free and homogenous sizes (initial body weight 15.84 (SEM 0.12) g) were fasted for 24 h. Then, 720 fish were randomly allocated to twelve floating net cages (1.0 m × 1.0 m × 2.0 m).

Table 1. Formulation and chemical proximate analysis of the experimental diets (% dry weight)

Ingredients	Experimental diets		
	FO	75% TO	75% TOC
White fishmeal	35	35	35
Soyabean meal	28	28	28
Wheat meal	23.8	23.8	23.72
Soyabean lecithin	1.5	1.5	1.5
Vitamin premix	2	2	2
Mineral premix*	2	2	2
Attractant mixture†	0.1	0.1	0.1
Mold inhibitor‡	0.1	0.1	0.1
Fish oil	7.5	1.875	1.875
Soyabean oil	0	1.875	1.875
Linseed oil	0	1.875	1.875
Pork lard	0	1.875	1.875
L-Carnitine	0	0	0.08
Total	100	100	100
Proximate analysis (DM %, mean values)			
Crude protein	42.29	42.49	42.07
Crude lipid	12.23	12.34	11.98

FO, fish oil; 75% TO, 75% fish oil substituted by terrestrial oils; 75% TOC, 75% TO supplemented with 800 mg/kg L-carnitine.

* The mixture of mineral mixture and vitamin mixture was purchased from Qingdao Master Biotechnology Co. Ltd.

† Attractant: mixture of 50% glycine acid and 50% betaine by weight.

‡ Mold inhibitor: mixture of 50% calcium propionic acid and 50% fumaric acid by weight.

Fish were hand-fed to apparent satiation, twice daily for 70 d. Each diet was randomly allocated to four replicate cages (sixty fish per cage). During the rearing period, the dissolved oxygen content was approximately 7.0 mg/l. Salinity ranged from 25.0 to 32.0 g/l and water temperature from 21.5 to 31.0°C.

Sample collection and analysis of growth data

At the termination of the experiment, fish were starved for 24 h and anaesthetised with MS222 (1:10 000; Sigma). Then, the number and weight of fish were determined for analyses of survival rate and final body weight. The wet weight of the liver, visceral and body and the length of six fish per cage were determined for analyses of morphometric parameters. The muscle and liver samples of the above six fish per cage were collected into 5 ml centrifuge tubes and stored at –80°C for compositional and fatty acid profile analyses. Blood samples of six fish per cage were collected from the caudal vein with a 1 ml syringe and clotted at room temperature to obtain serum samples for biochemical analyses. Liver samples were collected from twelve fish per cage, which contained six fish after serum collection. Samples were immediately frozen in liquid N₂ and stored at –80°C for analyses of antioxidant capacity, enzyme activity and gene expression.

Proximate composition and fatty acid profile analyses

DM was analysed by drying whole fish samples to a constant weight at 105°C. Crude protein and crude lipid contents of ingredients, diets and fish bodies were measured in accordance with the standard methods of the Association of Official Analytical Chemists (AOAC, 1995)⁽³⁰⁾. Fish tissue and diet samples were freeze-dried in a lyophilised chamber (Alpha 1-4 LDplus;

Table 2. Fatty acid profile of oil sources and the experimental diets (% total fatty acids)* (Mean values)

Fatty acid (% total fatty acids)	Fish oil†	Soyabean oil†	Linseed oil†	Pork lard	Experimental diets		
					FO	75% TO	75% TOC
14:0	8.77	0.08	0.07	1.23	6.84	2.38	2.31
16:0	19.15	10.32	5.39	22.41	22.92	14.63	14.41
18:0	4.26	3.95	3.76	15.53	5.61	5.39	5.44
20:0	1.18	0.27	0.17	0.83	1.45	0.98	1.28
∑SFA	33.36	14.62	9.39	40.01	36.82	23.38	23.44
16:1n-7	11.85	0.09	0.10	1.33	8.92	3.54	3.59
18:1n-9	9.75	26.10	20.45	34.40	13.80	19.17	19.40
∑MUFA	21.60	26.19	20.55	35.73	22.72	22.71	22.99
18:2n-6	1.54	49.65	15.52	18.37	11.62	23.23	23.65
20:4n-6	1.30	0.00	0.00	0.24	1.25	0.53	0.59
∑n-6 PUFA	2.84	49.65	15.52	18.61	12.87	23.76	24.24
18:3n-3	0.76	4.92	53.02	0.84	1.50	10.88	10.97
20:5n-3	12.34	0.00	0.00	0.00	7.22	3.55	3.23
22:6n-3	7.30	0.00	0.00	0.00	5.36	3.23	3.07
∑n-3 PUFA	20.41	4.92	53.02	0.84	14.08	17.66	17.26
n-3:n-6 PUFA	7.18	0.10	3.42	0.05	1.09	0.74	0.71
∑n-3 LC-PUFA	19.64	0.00	0.00	0.00	12.58	6.77	6.30

FO, fish oil; 75% TO, 75% fish oil substituted by terrestrial oils; 75% TOC, 75% TO supplemented with 800 mg/kg L-carnitine; LC-PUFA, long-chain PUFA.

* The contents of some fatty acids are minor and not detected, such as C20:0, C22:0, C24:0, C14:1, C20:1n-9, C22:1n-11, C20:2n-6, C18:3n-6, C20:3n-6, C18:4n-3 and C22:5n-3.

† Fatty acid profile of fish oil, soyabean oil and linseed oil is according to Li *et al.*⁽²⁹⁾ supplemental data.

Table 3. Sequences of the PCR primers used in this study

Target gene	Forward primers (5'-3')	Reverse primers (5'-3')	Accession number
<i>acc1</i>	GACTTGGCGGAATACCTACTGG	GCTTGCTGGATGATCTTTGCTT	XM027273319
<i>aco</i>	AGTGCCAGATGATCTTGAAGC	CTGCCAGAGGTAACCATTTCT	JX456348
<i>apoB100</i>	AGAGTGTGTCCAGGATAAAGATGC	CAGGGCTCAGGGTCTCAGTC	KM593126
<i>arg1</i>	AACCACCCGCAGGATTACG	AACTCACTGGCATCACCTCA	XM019269015
<i>β-Actin</i>	GACCTGACAGACTACCTCATG	AGTTGAAGGTGGTCTCGTGGA	GU584189
<i>cd36</i>	GAGCATGATGGAAAATGGTTCAAAG	CTCCAGAACTCCCTTTACCTTAG	KM593122
<i>cox-2</i>	CTGGAAGGCAACACAAGC	CGGTGAGAGTCAGGGACAT	KP259877
<i>cpt-1</i>	GCTGAGCCTGGTGAAGATGTTT	TCCATTTGGTTGAATTGTTACTGTCC	JX434612
<i>dgat2</i>	TTCGGTGCTTTCTGCAACTTCG	AAGGATGGGGAAGCGGAAGT	MG013506
<i>elovl4</i>	GGGCTCTTATTGGCTATGCT	TGCTTCTCCCGTTATCCTC	XM010739301
<i>elovl5</i>	ATCACCTTCTTTCATCTATCACC	CGGTGAGAGTCAGGGACAT	JQ320377
<i>fad6</i>	TTCGCTTCTCTGCTGCTATG	CCAGTCACGGTCTTCTCG	JX434611
<i>fas</i>	CAGCCACAGTGAGGTCATCC	TGAGGACATTGAGCCAGACAC	JX456351
<i>fatp1</i>	CAACCAGCAGGACCCATTACG	CATCCATCACCAGCACATCACC	KM593124
<i>gapdh</i>	GACAACGAGTTCGGATACAGC	CAGTTGATTGGCTTGTGGG	XM010743420
<i>hl</i>	TCCGTCCATCTATTCATTGACTCTC	GCCACTGTGAACCTTCTTGATATTG	JX456350
<i>ifnγ</i>	TCAGACCTCCGCACCATCA	GCAACCATTGTAACGCCACTTA	KM501500
<i>il-1β</i>	CATAGGGATGGGACAACGA	AGGGGACGGACACAAGGGTA	KJ459927
<i>il-10</i>	AGTCGGTTACTTTCTGTGGTG	TGATGACGCAATATGGTCTG	XM010738826
<i>il-6</i>	CGACACACCCACTATTTACAAC	TCCCATTTTCTGAACTGCCTCT	KU140675
<i>mtp</i>	ATGTCCAAAATGTTCTCCATGTCTG	ATGTCAATAGCCAACCTCCTTG	KP027412
<i>scd1</i>	AAAGGACGCAAGCTGGAAC	CTGGGACGAAGTACGACACC	KP202156
<i>srebp1</i>	TCTCCTTGACAGTCTGAGCCAAC	TCAGCCCTTGGATATGAGCCT	KP342262
<i>tnfr</i>	ACACCTCTCAGCCACAGGAT	CCGTGTCCTACTCCATAGTT	EF165623

acc1, acetyl-CoA carboxylase 1; *aco*, acyl-CoA oxidase; *arg-1*, arginase-1; *cd36*, cluster of differentiation 36; *cox-2*, cyclo-oxygenase-2; *cpt-1*, carnitine palmitoyltransferase 1; *dgat2*, diacylglycerol acyltransferase 2; *elovl4*, long-chain fatty acid protein 4; *elovl5*, elongation of very long-chain fatty acid protein 5; *fad6*, fatty acyl desaturase 6; *fas*, fatty acid synthase; *fatp1*, fatty acid transport protein 1; *gapdh*, glyceraldehyde-3-phosphate dehydrogenase; *hl*, hepatic lipase; *ifnγ*, interferon γ; *mtp*, microsomal TAG transfer protein; *scd1*, stearoyl-CoA desaturase 1; *srebp1*, sterol-regulatory element binding protein 1.

Christ) to analyse the moisture content. Lipid contents of freeze-dried liver and muscle samples were determined by chloroform/methanol (v/v, 2:1) following the procedures described by Folch *et al.*⁽³¹⁾. Fatty acid profile of oils and freeze-dried samples was measured according to procedures described by Metcalfe *et al.*⁽³²⁾ with minor modifications⁽³³⁾. Fatty acid profile was separated and determined by a HP6890 gas chromatograph (Agilent

Technologies Inc.). Results were shown as the percentage of each fatty acid to total fatty acid content.

Serum parameters and antioxidant capacity analysis

Contents of serum total cholesterol, total TAG, LDL-cholesterol and HDL-cholesterol and activities of serum alanine

Table 4. Growth and somatic parameters of large yellow croakers (Mean values with their standard errors; *n* 4)

	FO		75 % TO		75 % TOC	
	Mean	SEM	Mean	SEM	Mean	SEM
Initial body weight (g)	15.82	0.25	15.84	0.24	15.85	0.16
Final body weight (g)	35.60 ^b	0.32	38.22 ^a	0.46	37.93 ^a	0.35
Specific growth rate (%/d)*	1.16 ^b	0.01	1.26 ^a	0.03	1.25 ^a	0.00
Survival rate (%)†	87.92	1.97	90.83	0.83	89.17	1.60
Feed intake (%/d)‡	2.13	0.02	2.12	0.02	2.10	0.02
Feed efficiency ratio§	0.52 ^b	0.00	0.56 ^a	0.01	0.56 ^a	0.01
Viscerosomatic index (%)	6.92	0.23	7.18	0.21	7.05	0.30
Hepatosomatic index (%)¶	1.98 ^b	0.04	2.14 ^a	0.03	2.07 ^{a,b}	0.05
Condition factor (%)**	1.02	0.03	1.05	0.02	1.00	0.04

FO, fish oil; 75 % TO, 75 % fish oil substituted by terrestrial oils; 75 % TOC, 75 % TO supplemented with 800 mg/kg L-carnitine.

^{a,b} Mean values in a row sharing the same superscript letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

* Specific growth rate (%/d) = $(\ln(\text{final body weight}) - \ln(\text{initial body weight})) \times 100 / \text{duration of experimental days}$.

† Survival rate (%) = $100 \times \text{final fish number} / \text{initial fish number}$.

‡ Feed intake (%/d) = $100 \times \text{dry feed fed in g} / ((\text{final body weight} + \text{initial body weight}) / 2) / \text{duration of experimental days}$.

§ Feed efficiency ratio = $\text{wet weight gain in g} / \text{dry feed fed in g}$.

|| Viscerosomatic index (%) = $100 \times \text{visceral wet weight} / \text{final body weight}$.

¶ Hepatosomatic index (%) = $100 \times \text{liver wet weight} / \text{final body weight}$.

** Condition factor (%) = $100 \times \text{final body weight} / (\text{final body length}^3)$.

transaminase and aspartate aminotransferase were analysed with matching commercial reagent kits (Mindray Bio Medical Co. Ltd) by an automatic biochemical analyzer (BS180; Mindray). Activities of catalase, superoxide dismutase, total antioxidant capacity (T-AOC) and the content of malondialdehyde in the liver of fish were determined by commercial kits (Nanjing Jiancheng Bioengineering Institute).

Activities of carnitine palmitoyltransferase-I and acyl-CoA oxidase

Mitochondria was extracted, and its integrity was assessed according to procedures described by Suarez & Hochachka⁽³⁴⁾ and Aprille & Asimakis⁽³⁵⁾, respectively. CPT-I activity was determined according to the method of Bieber & Fiol⁽³⁶⁾ with some modifications^(37,38). Briefly, CPT-I activity was determined by the CoA-SH formation in the 5',5'-dithio-bis-(2-nitrobenzoic acid) (Sigma) reaction from palmitoyl-CoA (Sigma) at 412 nm. Results were expressed as 1 μmol of product formed per min per mg of mitochondrial protein at 25°C. Protein concentration of extracted samples was determined according to procedures of Bradford Protein Assay Kit (Beyotime Biotechnology).

Peroxisomal ACO activity was determined by the H_2O_2 -dependent oxidation of 2',7'-dichlorofluorescein (Sigma) according to the method of Small *et al.*⁽³⁹⁾. The reaction medium consisted of 11 mM potassium phosphate buffer (pH 7.4) (Sangon biotech), 50 μM horseradish peroxidase type II (Sigma), 0.05 mM 2',7'-dichlorofluorescein (Sigma), 40 mM 3-Amino-1,2,4-triazole (Sigma) and 0.02 % Triton-X 100 (Sangon biotech), and the reaction was initiated by the addition of 30 μM palmitoyl-CoA (Sigma). The assay medium contained 10–40 mg of protein in a total volume of 1 ml.

Total RNA extraction, cDNA synthesis and real-time quantitative PCR

Total RNA was extracted from the liver of experimental fish using TRIzol reagent (Takara), and the quality and quantity of RNA

were detected by a 1.2 % denaturing agarose gel and a NanoDrop spectrophotometer (Thermo Scientific), respectively. The extracted RNA was reversely transcribed to cDNA by PrimeScript™ RT reagent Kit (Takara). β -Actin and glyceraldehyde-3-phosphate dehydrogenase (*gapdh*) were used as reference genes to normalise the expression levels of genes⁽⁴⁰⁾. The real-time quantitative PCR primers (Table 3) of candidate genes were designed based on nucleotide sequences of large yellow croaker. The real-time quantitative PCR was performed in a quantitative thermal cycler (Mastercycler ep realplex; Eppendorf). The volume and procedure of real-time quantitative PCR were carried out as described previously⁽⁴¹⁾. Only one PCR product in these reactions was confirmed by melting curve analysis at the end of each reaction. Standard curves were generated with 4-fold serial dilutions (in triplicate) of cDNA sample, and the amplification efficiency was analysed by the equation: $E = 10^{(-1/\text{slope})} - 1$ ⁽⁴²⁾. The amplification efficiencies of all detected genes ranged between 0.95 and 1.04. The expression levels of all genes were calculated using the comparative Ct method ($2^{-\Delta\Delta\text{Ct}}$ method) as described by Livak & Schmittgen⁽⁴³⁾.

Statistical analyses

All data analyses were processed by SPSS 19.0 (IBM). Results were shown as mean values with their standard errors. Statistics were performed to a one-way ANOVA and followed by Tukey's test. Statistics with $P < 0.05$ were considered to be significant.

Results

Survival and growth performance

The survival rate of fish ranged from 87.92 to 90.83 %, but no significant differences were found among dietary treatments ($P > 0.05$). The dietary oil source markedly affected specific growth rate and feed efficiency ratio of fish in the present study, with specific growth rate and feed efficiency ratio significantly

Table 5. Body composition analysis of large yellow croakers (Mean values with their standard errors; n 4)

	FO		75 % TO		75 % TOC	
	Mean	SEM	Mean	SEM	Mean	SEM
Whole body (% wet weight)						
Moisture	74.99	0.40	75.06	0.73	75.96	0.80
Lipid	7.16	0.19	6.91	0.37	6.54	0.17
Protein	14.74	0.15	14.86	0.28	14.32	0.17
Liver (% wet weight)						
Moisture	59.64 ^a	1.08	53.10 ^b	1.11	56.80 ^{a,b}	1.53
Lipid	22.10 ^b	0.66	27.13 ^a	0.63	23.32 ^b	0.94
Muscle (% wet weight)						
Moisture	73.31 ^b	0.61	72.53 ^b	0.48	74.48 ^a	0.64
Lipid	7.95	0.37	8.53	0.34	8.60	0.38

FO, fish oil; 75 % TO, 75 % fish oil substituted by terrestrial oils; 75 % TOC, 75 % TO supplemented with 800 mg/kg L-carnitine.

^{a,b} Mean values in a row sharing the same superscript letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

increased in fish fed diets with 75 % TO and 75 % TOC compared with the control group ($P < 0.05$). Hepatosomatic index was dramatically increased in fish fed the diet with 75 % TO compared with the control group ($P < 0.05$). When the 75 % TO diet was supplemented with 800 mg/kg L-carnitine, a decreasing trend (from 2.14 to 2.07) was observed in hepatosomatic index of fish compared with fish fed the diet with 75 % TO. There was no significant difference in feed intake, viscerosomatic index or condition factor of fish among dietary treatments ($P > 0.05$) (Table 4).

Body composition analysis

There was no significant difference in the moisture, lipid or protein contents of whole fish body among dietary treatments ($P > 0.05$). Fish fed the diet with 75 % TO had markedly higher lipid content of the liver than the control group ($P < 0.05$). However, the lipid content of the liver was decreased in fish fed the diet with 75 % TOC compared with 75 % TO ($P < 0.05$). No significant difference was found in lipid content of the liver in fish fed the diet with 75 % TOC and the control group. The moisture was markedly decreased in the liver of fish fed the diet with 75 % TO compared with the control group ($P < 0.05$). However, the moisture was dramatically increased in the muscle of fish fed the diet with 75 % TOC compared with 75 % TO and the control group ($P < 0.05$) (Table 5).

Fatty acid profile in the liver

The contents of C14:0, C16:0, C16:1 n -7, C20:4 n -6 (arachidonic acid), C20:5 n -3 (EPA), C22:6 n -3 (DHA), SFA, n -3 LC-PUFA and the ratio of n -3: n -6 PUFA were dramatically decreased in the liver of fish fed diets with 75 % TO and 75 % TOC compared with the control group ($P < 0.05$). Furthermore, the contents of C18:1 n -9, C18:2 n -6, C18:3 n -3 and n -6 PUFA were significantly increased in the liver of fish fed diets with 75 % TO and 75 % TOC compared with the control group ($P < 0.05$). The supplementation of 800 mg/kg L-carnitine in the 75 % TO diet significantly decreased the content of C18:3 n -3, DHA and n -3 LC-PUFA in the liver of fish compared with fish fed the diet with 75 % TO (Table 6).

Serum biochemical indexes

The content of TAG was dramatically increased in fish fed the diet with 75 % TO compared with the control group ($P < 0.05$). The content of HDL-cholesterol was significantly decreased in fish fed the diet with 75 % TO compared with the control group ($P < 0.05$). The content of LDL-cholesterol was significantly decreased, but the content of HDL-cholesterol was significantly increased in fish fed the diet with 75 % TOC compared with 75 % TO ($P < 0.05$). No significant difference was found in activities of serum aspartate aminotransferase or alanine transaminase in fish among dietary treatments ($P > 0.05$) (Table 7).

Antioxidant capacity

A decreasing trend (from 1.16 to 1.03) was found in the activity of T-AOC (Fig. 1(C)) in fish fed the diet with 75 % TO compared with the control group, but no significant difference was found between the two treatments ($P > 0.05$). The supplementation of 800 mg/kg L-carnitine in the 75 % TO diet significantly increased the activity of superoxide dismutase (Fig. 1(B)) and T-AOC (Fig. 1(C)) in fish compared with fish fed the diet with 75 % TO ($P < 0.05$). No significant differences were found in the content of malondialdehyde (Fig. 1(A)) or the activity of catalase (Fig. 1(D)) in fish among dietary treatments ($P > 0.05$).

Enzyme activities of CPT-I and ACO

The activity of CPT-I was dramatically increased in fish fed the diet with 75 % TOC compared with fish fed the diet with 75 % TO and the control group ($P < 0.05$). However, no significant difference was found in the activity of ACO among dietary treatments ($P > 0.05$) (Fig. 2).

Expression of genes related to lipid metabolism

The mRNA expression of *diacylglycerol acyltransferase 2* (*dgat2*) and *acetyl-CoA carboxylase 1* (*acc1*) was dramatically increased in fish fed the diet with 75 % TO compared with fish fed the diet with 75 % TOC and the control group ($P < 0.05$) (Fig. 3(C)). Expression of genes related to fatty acid synthesis (*elongation of very long-chain fatty acid protein 4* (*elovl4*), *elongation of very long-chain fatty acid protein 5* (*elovl5*) and *fatty*

Table 6. Fatty acid profile (% total fatty acids) in the liver of large yellow croakers (Mean values with their standard errors; n 4)*

	FO		75 % TO		75 % TOC	
	Mean	SEM	Mean	SEM	Mean	SEM
14:0	4.31 ^a	0.26	1.77 ^b	0.04	1.85 ^b	0.02
16:0	22.88 ^a	0.57	17.69 ^b	0.44	18.13 ^b	0.14
18:0	6.41	0.49	6.50	0.48	6.64	0.51
20:0	1.30	0.11	1.44	0.05	1.50	0.13
∑SFA	34.90 ^a	0.66	27.40 ^b	0.67	28.12 ^b	0.56
16:1 n -7	12.93 ^a	0.36	6.84 ^b	0.16	7.12 ^b	0.09
18:1 n -9	22.16 ^b	1.16	27.85 ^a	0.64	29.46 ^a	0.54
∑MUFA	35.09	0.94	34.69	0.57	36.58	0.47
18:2 n -6	7.90 ^b	0.54	18.21 ^a	0.67	15.46 ^a	1.03
20:4 n -6	0.50 ^a	0.03	0.25 ^b	0.00	0.22 ^b	0.01
∑ n -6 PUFA	8.41 ^b	0.53	18.46 ^a	0.67	15.69 ^a	1.03
18:3 n -3	1.48 ^c	0.23	7.46 ^a	0.35	6.09 ^b	0.08
20:5 n -3 (EPA)	3.34 ^a	0.15	0.78 ^b	0.02	0.48 ^b	0.04
22:6 n -3 (DHA)	2.08 ^a	0.10	0.83 ^b	0.01	0.54 ^c	0.04
∑ n -3 PUFA	6.90 ^b	0.26	9.07 ^a	0.32	7.11 ^a	0.07
n -3: n -6 PUFA	0.83 ^a	0.07	0.49 ^b	0.02	0.46 ^b	0.03
∑ n -3 LC-PUFA	5.42 ^a	0.23	1.61 ^b	0.04	1.02 ^c	0.08

FO, fish oil; 75 % TO, 75 % fish oil substituted by terrestrial oils; 75 % TOC, 75 % TO supplemented with 800 mg/kg L-carnitine; LC-PUFA, long-chain PUFA.

^{a,b,c} Mean values in a row sharing the same superscript letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

* Some fatty acids, of which the contents are minor, trace amount or not detected, such as C20:0, C22:0, C24:0, C14:1, C20:1 n -9, C22:1 n -11, C20:2 n -6, C18:3 n -6, C18:4 n -3, C22:5 n -3, are not listed in the table.

Table 7. Serum biochemical indexes and enzyme activities of large yellow croakers (Mean values with their standard errors; n 4)

	FO		75 % TO		75 % TOC	
	Mean	SEM	Mean	SEM	Mean	SEM
TC (mmol/l)	2.79	0.29	3.45	0.29	2.89	0.30
TAG (mmol/l)	3.71 ^b	0.44	5.09 ^a	0.40	3.80 ^{a,b}	0.32
HDL-cholesterol (mmol/l)	0.79 ^a	0.03	0.56 ^b	0.01	0.77 ^a	0.06
LDL-cholesterol (mmol/l)	0.75 ^{a,b}	0.07	0.93 ^a	0.11	0.63 ^b	0.03
ALT (U/l)	55.76	2.21	60.40	2.63	57.86	3.09
AST (U/l)	5.33	0.47	6.12	0.60	5.65	0.50

FO, fish oil; 75 % TO, 75 % fish oil substituted by terrestrial oils; 75 % TOC, 75 % TO supplemented with 800 mg/kg L-carnitine; TC, total cholesterol; ALT, alanine transaminase; AST, aspartate aminotransferase.

^{a,b} Mean values in a row sharing the same superscript letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

acyl desaturase 6 (fad6) was dramatically increased in fish fed diets with 75 % TO and 75 % TOC compared with the control group ($P < 0.05$) (Fig. 3(D)). The mRNA expression of *cpt-1* and *cluster of differentiation 36 (cd36)* was significantly increased in fish fed the diet with 75 % TOC compared with fish fed the diet with 75 % TO and the control group ($P < 0.05$) (Fig. 3(A) and (B)). The mRNA expression of *stearoyl-CoA desaturase 1 (scd1)* and *hepatic lipase (hl)* was significantly increased in fish fed the diet with 75 % TOC compared with the control group ($P < 0.05$) (Fig. 3(A) and (D)). There was no significant difference in the mRNA expression of *sterol-regulatory element binding protein 1 (srebp1)*, *fatty acid synthase (fas)*, *aco*, *fatty acid transport protein 1 (fatp1)*, *microsomal TAG transfer protein (mtp)* or *apob100* among dietary treatments ($P > 0.05$) (Fig. 3(A)–(C)).

Expression of genes related to inflammation

The mRNA expression of pro-inflammatory genes (*tnfa* and *interferon γ (ifn γ)*) was significantly increased in fish fed the diet with 75 % TO compared with the control group ($P < 0.05$), while

they both were repressed in fish fed the diet with 75 % TOC compared with 75 % TO ($P < 0.05$) (Fig. 4(B)). The mRNA expression of anti-inflammatory genes (*il-10* and *arginase-1 (arg1)*) was significantly increased in fish fed the diet with 75 % TOC compared with fish fed the diet with 75 % TO and the control group ($P < 0.05$) (Fig. 4(A) and (B)). There was no significant difference in the mRNA expression of *il-1 β* , *il-6* or *cyclo-oxygenase-2 (cox-2)* of fish among dietary treatments ($P > 0.05$) (Fig. 4(A) and (B)).

Discussion

In the present study, specific growth rate significantly increased in fish fed diets with 75 % TO and 75 % TOC compared with the control group. The growth-promoting effect was possibly due to the balance of fatty acids with the supplementation of various TO. In the present study, the content of n -3 LC-PUFA in 75 % TO and 75 % TOC diets was approximately 0.83 %, which was supplied in an appropriate level for large yellow croaker. A previous study has reported that large yellow croaker fed

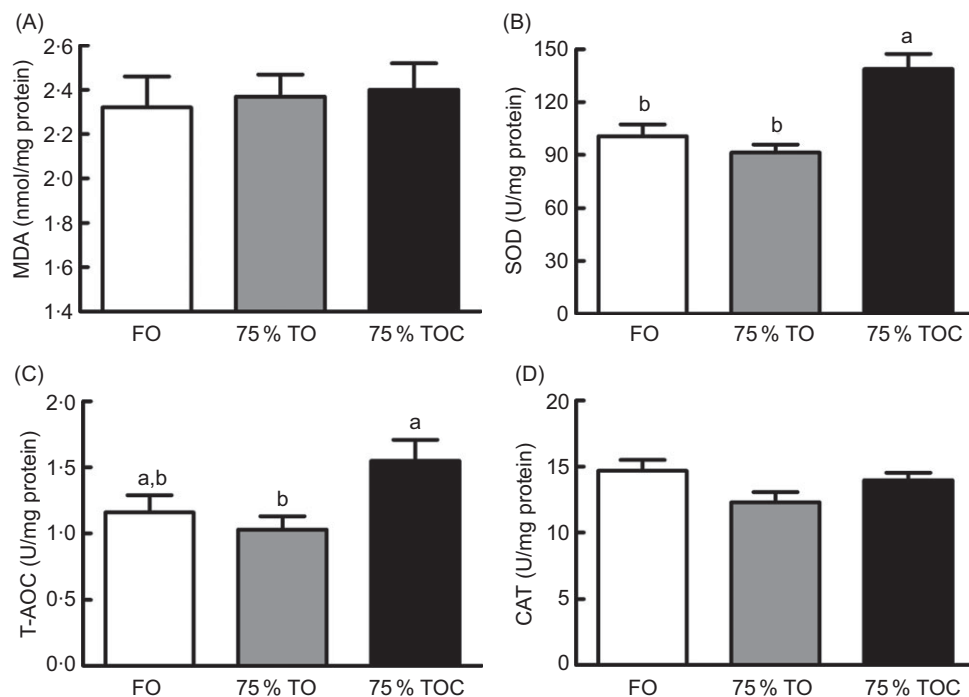


Fig. 1. Concentration of (A) malondialdehyde (MDA) and activities of (B) superoxide dismutase (SOD), (C) total antioxidant capacity (T-AOC) and (D) catalase (CAT) in the liver of large yellow croakers. FO, fish oil; 75 % TO, 75 % fish oil substituted by terrestrial oils; 75 % TOC, 75 % TO supplemented with 800 mg/kg L-carnitine. Values are means (n 4), with their standard errors represented by vertical bars. ^{a,b} Columns sharing the same letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

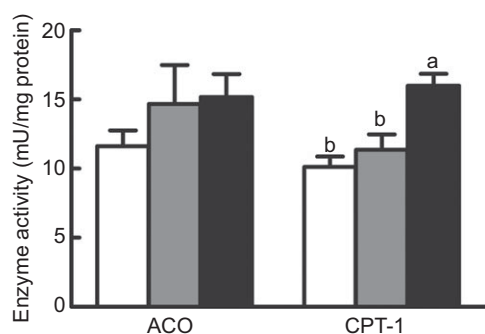


Fig. 2. Enzyme activities of carnitine palmitoyltransferase 1 (CPT-I) and acyl-CoA oxidase (ACO) in the liver of large yellow croakers. (□), Fish oil (FO); (▒), 75 % FO substituted by terrestrial oils (75 % TO); (■), 75 % TO supplemented with 800 mg/kg L-carnitine. Values are means (n 4), with their standard errors represented by vertical bars. ^{a,b} Columns sharing the same letter or absence of letters are not significantly different determined by Tukey's test ($P > 0.05$).

0.60–0.98 % dietary n -3 LC-PUFA have better growth performance, nonspecific immune responses and disease resistance than the control group (0.15 % n -3 LC-PUFA)⁽³³⁾. Also, the ratio of ALA:LNA in 75 % TO and 75 % TOC diets was kept in an appropriate value (0.47) for large yellow croaker, while the corresponding ratio in the FO diet (0.13) was lower than the aforementioned ratio in TO diet. A previous study has shown that the ratio of dietary ALA:LNA in 0.5 is beneficial for the growth of large yellow croaker⁽⁴⁴⁾.

Previously, Turchini *et al.*⁽²⁾ found that 60–75 % of dietary FO could be replaced by alternative TO in almost all finfish species

when the requirement of essential fatty acid was met. However, a high level of TO might increase abnormal lipid deposition of fish⁽⁴⁵⁾. In this experiment, significantly higher contents of LDL-cholesterol and TAG as well as hepatic lipid content were found in fish fed the diet with 75 % TO than the control group. Similar results were also observed in Malaysian mahseer (*Tor tambroides*)⁽⁴⁶⁾, rainbow trout (*Oncorhynchus mykiss*)⁽⁴⁷⁾ and turbot⁽⁶⁾. Previous studies have demonstrated that oleic acid is more likely to increase steatosis and plasma lipid concentrations than palmitic acid^(48,49) and PUFA^(50,51). Accordingly, the increased lipid deposition of experimental fish could be due to the increase of dietary oleic acid in the 75 % TO diet compared with the FO diet. However, hepatic lipid deposition and the content of LDL-cholesterol were decreased in fish fed the diet with 75 % TOC compared with 75 % TO. Sang *et al.*⁽²⁷⁾ reported that the supplementation of 800 mg/kg L-carnitine significantly improved growth performance and reduced serum TAG and total cholesterol levels in large yellow croaker compared with the control group ($P < 0.05$). The hypolipidemic effect was probably due to the role of L-carnitine in facilitating the transport of long-chain fatty acids into mitochondria for β -oxidation^(52,53), as contents of C18:3 n -3, DHA and total n -3 LC-PUFA were dramatically decreased, and the content of n -6 PUFA was also numerically lower in the liver of fish fed the diet with 75 % TOC compared with 75 % TO.

The decreased content of PUFA could be attributed to changes of fatty acid oxidation and synthesis in fish fed the diet with 75 % TOC compared with 75 % TO. As for fatty acid utilisation and oxidation, CPT-I is a rate-limiting enzyme that controls the mitochondrial uptake of long-chain acyl-CoA and facilitates

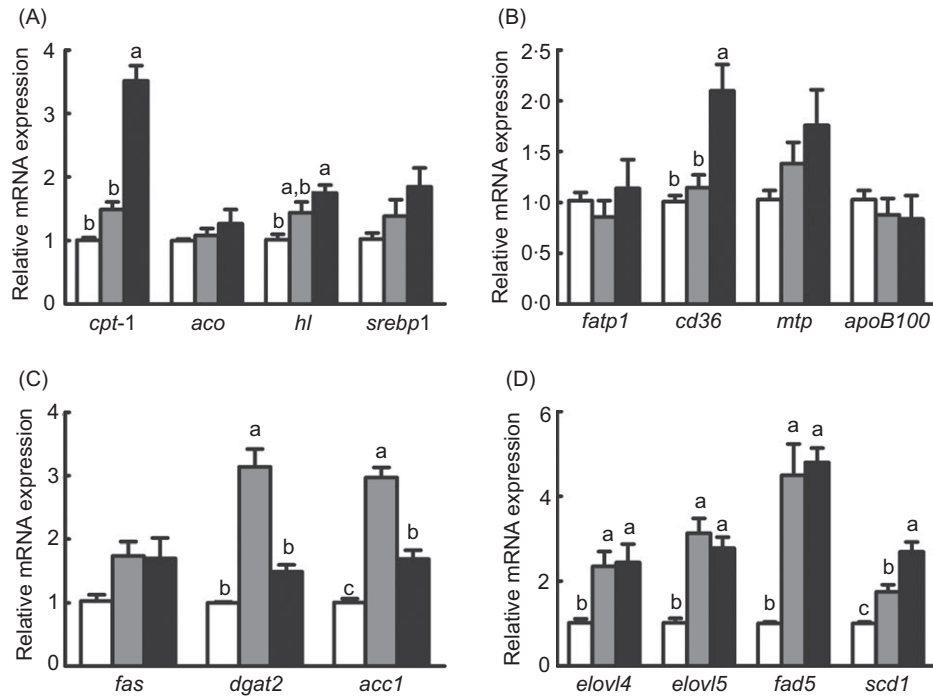


Fig. 3. Expression of genes related to lipid metabolism in the liver of large yellow croakers. *cpt-1*, Carnitine palmitoyltransferase I; *aco*, acyl-CoA oxidase; *hl*, hepatic lipase; *srebp1*, sterol-regulatory element binding protein 1; *fas*, fatty acid synthase; *fatp1*, fatty acid transport protein 1; *cd36*, cluster of differentiation 36; *mtp*, microsomal TAG transfer protein; *dgat2*, diacylglycerol acyltransferase 2; *acc1*, acetyl-CoA carboxylase 1; *elovl4*, long-chain fatty acid protein 4; *elovl5*, elongation of very long-chain fatty acid protein 5; *fad6*, fatty acyl desaturase 6; *scd1*, stearoyl-CoA desaturase 1. (◻), Fish oil (FO); (◼), 75 % FO substituted by terrestrial oils (75 % TO); (◼), 75 % TO supplemented with 800 mg/kg L-carnitine. Values are means (*n*4), with their standard errors represented by vertical bars. ^{a,b,c} Columns sharing the same letter or absence of letters are not significantly different determined by Tukey's test (*P* > 0.05).

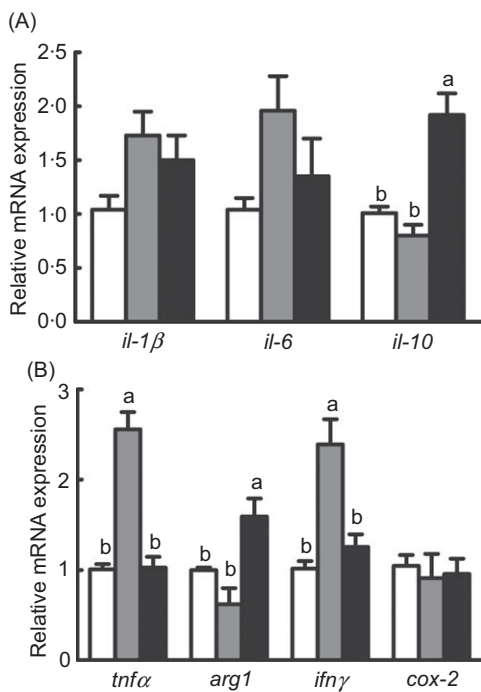


Fig. 4. Expression of genes related to inflammation in the liver of large yellow croakers. *arg-1*, Arginase-1; *ifnγ*, interferon γ ; *cox-2*, cyclo-oxygenase-2. (◻), Fish oil (FO); (◼), 75 % FO substituted by terrestrial oils (75 % TO); (◼), 75 % TO supplemented with 800 mg/kg L-carnitine. Values are means (*n*4), with their standard errors represented by vertical bars. ^{a,b} Columns sharing the same letter or absence of letters are not significantly different determined by Tukey's test (*P* > 0.05).

long-chain fatty acids for β -oxidation⁽⁵⁴⁾. CD36 is a key NEFA uptake transporter that involves transferring fatty acids into cells⁽⁵⁵⁾ or mitochondria^(56,57) for utilisation. HL is a pivotal TAG lipase that contributes to vascular lipoprotein degradation, TAG hydrolysis and the uptake of lipoprotein into the liver⁽⁵⁸⁾. In the present study, the mRNA expression of *cpt-1*, *cd36* and *hl* as well as the activity of CPT-I was significantly increased in fish fed the diet with 75 % TOC compared with 75 % TO. These results showed that L-carnitine facilitated the utilisation of lipid in the diet and promoted fatty acid oxidation. As for fatty acid synthesis, *elovl4*, *elovl5* and *fad6* were key genes for the biosynthesis of arachidonic acid, EPA and DHA from LNA and ALA in fish⁽¹²⁾. No significant difference was observed in the mRNA expression of *elovl4*, *elovl5* or *fad6* in fish fed diets with 75 % TOC and 75 % TO, which meant that the supplementation of L-carnitine had no significant effect on the capacity of LC-PUFA biosynthesis. Further, vertebrates cannot synthesise C18: 2*n*-6 and C18: 3*n*-3 from C16: 1*n*-7 and C18: 1*n*-9⁽¹²⁾, and the capacity of biosynthesis of C20 and C22 LC-PUFA from C18 PUFA is very limited in most marine fish species⁽⁵⁹⁾. Above all, the unchanged PUFA synthesis and increased PUFA oxidation could be attributed to the decreased content of PUFA in fish fed the diet with 75 % TOC compared with 75 % TO. Similarly, dietary L-carnitine promoted the utilisation of fatty acid and decreased the content of PUFA in red porgy (*Pagrus pagrus*, L.)⁽⁶⁰⁾, cobia (*Rachycentron canadum*)⁽⁶¹⁾ and beluga sturgeon (*Huso buso*)⁽⁶²⁾.

Antioxidant capacity is one of the most commonly used indexes to evaluate the physical condition of fish⁽⁶³⁾. In this study, the activity of T-AOC showed a decreasing trend (from

1.16 to 1.03) in fish fed the diet with 75 % TO compared with the control group. When the 75 % TO diet was supplemented with 800 mg/kg L-carnitine, the activity of superoxide dismutase and T-AOC was dramatically increased in fish compared with fish fed the diet with 75 % TO. A previous study has reported that L-carnitine induces Nrf2/Keap1 pathway activation *in vivo* and *in vitro* and increases the antioxidant enzyme activities (catalase, T-superoxide dismutase, GSH-PX) in *Rhynchocypris lagowskii*⁽⁶⁴⁾. Further, L-carnitine played some crucial physiological roles in the inhibition of superoxide radical formation⁽⁶⁵⁾. The increased antioxidant capacity could be attributed to the modulatory effects of L-carnitine on critical antioxidant enzymes in fish fed the diet with 75 % TOC compared with 75 % TO.

In addition to antioxidant capacity, the inflammation in fish was also an important indicator to evaluate the health status. In the present study, the mRNA expression of pro-inflammatory genes (*tnfa* and *ifn γ*) was significantly higher in fish fed the diet with 75 % TO than the control group. However, the mRNA expression of the pro-inflammatory genes was decreased in fish fed the diet with the 75 % TOC compared with 75 % TO. The mRNA expression of anti-inflammatory genes (*il-10* and *arg-1*) was increased in fish fed the diet with the 75 % TOC compared with 75 % TO. Previous studies have demonstrated that dietary L-carnitine can improve the immunity and disease resistance of juvenile narrow clawed crayfish (*Astacus leptodactylus leptodactylus*)⁽⁶⁶⁾, black carp (*Mylopharyngodon piceus*)⁽⁶⁷⁾ and common carp (*Cyprinus carpio* L.)⁽⁶⁸⁾. L-Carnitine exerted protective effects by inhibiting the generation of ammonia and xenobiotics in rat hepatocytes⁽⁶⁹⁾. Future research is required to explore the specific mechanism how L-carnitine decreased the inflammation of fish fed the diet with 75 % TOC compared with 75 % TO.

In conclusion, dietary TO mixture (75 % TO and 75 % TOC diets) increased the growth of large yellow croaker compared with the control group, which was probably due to the appropriate level of *n*-3 LC-PUFA and the appropriate ratio of ALA:LNA. Although the growth was not significantly different, the supplementation of 800 mg/kg L-carnitine in the 75 % TO diet (75 % TOC) could increase antioxidant capacity, fatty acid oxidation and decrease lipid abnormal deposition and the expression of inflammatory genes compared with 75 % TO.

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Q. A. and K. M. designed the research; X. L., Q. C. and Q. C. conducted the research; X. L. analysed the data and wrote the paper. We appreciate Yunqiang Zhang, Jiamin Li, Weiwei Dai, Xiufei Cao, Ye Gong, Tao Ding and Md Golam Sajed Riar for their help in revising the article.

The authors declare that there are no conflicts of interests associated with the manuscript.

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