



Synergistic effects of dietary cholesterol and taurine on growth performance and cholesterol metabolism in juvenile turbot (*Scophthalmus maximus* L.) fed high plant protein diets

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饲料胆固醇和牛磺酸对大菱鲆幼鱼生长表现和胆固醇代谢的协同作用（高植物蛋白饲料中）

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ABSTRACT

本研究是调查胆固醇，牛磺酸和二者在饲料中组合对于大菱鲆生长和胆固醇代谢的影响，在饲喂高植物蛋白饲料基础之上。基础饲料（对照组）是14.5%鱼粉和50%粗蛋白。其他三组等氮和等脂的实验饲料，由基础饲料中添加1%胆固醇，1%牛磺酸和1%胆固醇和1%牛磺酸的组合。结果表明WGR在这三组中显著高于对照组。TC组在所有处理中有最高的生长表现。血浆总胆固醇，游离胆固醇和低密度脂蛋白胆固醇水平在TC组明显比C组低。C组的CYP7A1活性显著比对照组高，并且CYP7A1的活性在T组中是最高的。兼并引物扩增的HMG-CoAr的PCR产物是331bp。HMG-CoAr的mRNA水平在TC组显著比C组低。这些结果表明1%胆固醇和1%牛磺酸的组合有助于大菱鲆幼鱼在高植物蛋白的饲料中获得更好的生长而没有负面效果。

of cholesterol, taurine and combination of dietary cholesterol and cholesterol metabolism in juvenile turbot (*Scophthalmus maximus* L.) with 14.5% fish meal and 50% crude protein was formulated. Experimental diets were prepared with the supplementation of 1.0% cholesterol and 1.0% taurine to the basal diet, which The results showed that the weight gain rate in fish fed C-1.0% than that in fish fed the control diet. Especially, fish fed TC diet data among dietary treatments. The plasma total cholesterol, cholesterol levels were significantly lower in fish fed TC diet compared with fish fed C-1.0% diet showed significantly higher activity of CYP7A1 in fish fed T-1.0% diet ($P < 0.05$), and activity of CYP7A1 in fish fed T-1.0% diet. The PCR product of 3-hydroxy-3-methylglutaryl-Coenzyme A synthase primers was 331 bp. The HMG-CoAr mRNA levels were that in fish fed C-1.0% diet ($P < 0.05$). These results suggested that taurine is helpful for juvenile turbot fed high plant protein diets to improve growth performance.

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兼并引物

1. Introduction

Taurine is an essential amino acid derivative and was first identified in bile (Park and Lee, 1998). It is an important antioxidant (Thurston et al., 1996). HMG-CoAr synthesis is rate-limiting and activity sizes (Yokoyama et al., 1997)证明虹鲟能由半胱氨酸合成牛磺酸。L-半胱氨酸脱羧酶是肝牛磺酸合成的限速酶 (Kim et al., 2005b), 而且这种酶的活性在不同种类和不同大小的鱼中也是不同的 (Yokoyama et al., 2001)。当移走鱼粉时, 常常补充EAA, 脂肪和矿物质。但是很多必需营养经常被忽视。例如, 多数植物原料缺乏牛磺酸 (Yamamoto et al., 1998)。植物饲料中牛磺酸补充提高了牙鲆的WGR和FER (Kim et al., 2005a; Park et al., 2002) 虹鲟 (Gaylord et al., 2006), 军曹鱼 (Lunger et al., 2007) 和黄尾鱼 (Takagi et al., 2008)。因此, 在饲喂植物饲料时可能需要牛磺酸的补充。

牛磺酸是氨基酸衍生物, 首次从牛的胆汁中分离出来 (Park and Lee, 1998)。在哺乳动物和鱼中, 已知牛磺酸在膜稳定、抗氧化、解毒 (Wright et al., 1986)、渗透调节 (Thurston et al., 1996)、与胆汁酸结合 (Goto et al., 1996) 起到重要的生理作用。但是, 虽然牛磺酸能由鱼部分合成, 合成的牛磺酸不能满足大部分鱼类的需求 (Gaylord et al., 2006; Kim et al., 2005b; Yokoyama et al., 1997, 2001)。Yokoyama et al. (1997) 证明虹鲟能由半胱氨酸合成牛磺酸。L-半胱氨酸脱羧酶是肝牛磺酸合成的限速酶 (Kim et al., 2005b), 而且这种酶的活性在不同种类和不同大小的鱼中也是不同的 (Yokoyama et al., 2001)。当移走鱼粉时, 常常补充EAA, 脂肪和矿物质。但是很多必需营养经常被忽视。例如, 多数植物原料缺乏牛磺酸 (Yamamoto et al., 1998)。植物饲料中牛磺酸补充提高了牙鲆的WGR和FER (Kim et al., 2005a; Park et al., 2002) 虹鲟 (Gaylord et al., 2006), 军曹鱼 (Lunger et al., 2007) 和黄尾鱼 (Takagi et al., 2008)。因此, 在饲喂植物饲料时可能需要牛磺酸的补充。

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often supplemented when fish meal (FM) is removed. However numerous essential nutrients are often overlooked. For example, most plant-based ingredients are deficient in taurine (Yamamoto et al., 1998). Dietary taurine supplementation improved weight gain rate (WGR) and feed efficiency rate in Japanese flounder (Kim et al., 2005a; Park et al., 2002), rainbow trout (Gaylord et al., 2006), cobia (Lunger et al., 2007) and yellowtail (Takagi et al., 2008) fed plant-based diets. Therefore, taurine supplementation may be required when fish are fed plant-based diets.

Cholesterol is an essential dietary nutrient for a variety of marine crustaceans. Cholesterol is many marine crustaceans' essential nutrient (Hernandez et al., 2004; Holme et al., 2006; Teshima, 1997). However, due to the limited investigation about the potential demand for cholesterol in crustaceans (Deng et al., 2010; National Research Council, 1993; Sealey et al., 2001). Fish meal is rich in cholesterol, but in most plant-based diets, the level is low (Cheng and Hardy, 2004; Deng et al., 2010). In addition, dietary cholesterol supplementation significantly improved the FI and WGR of Twibell and Wilson (2004) in SBM diet, and in yellowtail (Chen, 2006) fed 41% SBM diet.

(Twibell and Wilson, 2004) fed soybean meal (SBM)-based diet and Japanese flounder fed diet including 41% SBM (Chen, 2006).

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centrifugation (4000 g for 10 min) at 4 °C and immediately stored at -80 °C until analysis. Liver samples were frozen in liquid nitrogen and stored at -80 °C for subsequent determination of lipid content, activity of CYP7A1, and RNA isolation. The feces collection method was the same as described in Deng et al. (2010).

During the feeding trial, water temperature was controlled at 19–22 °C, pH 7.5–8.0, salinity 30–33‰. Ammonia nitrogen was lower than 0.4 mg/L, nitrite nitrogen was lower than 0.1 mg/L, and dissolved oxygen was higher than 6.0 mg/L. Each tank was aerated 24 h daily.

2.3. Chemical analyses

2.3.1. Body composition and energy assays

Dry matter was analyzed for ingredients, crude protein, crude fat, ash, and energy. Crude protein was determined using the Kjeldahl method (Association of Official Analytical Chemists, AOAC, 1995) and multiplied by 6.25. Crude fat was determined using the Soxhlet method (Parr 1281 automatic bomb calorimeter, Parr, Moline, IL, USA). Duplicate analyses were conducted for each sample.

2.3.2. Cholesterol, taurine and total bile acid assays

The concentration of total cholesterol (TC), high-density lipoprotein cholesterol (HDL-C) and low-density lipoprotein cholesterol (LDL-C) in plasma were determined according to the method described by Deng et al. (2010). The concentration of free cholesterol (FC) in plasma was measured by peroxidase-anti-peroxidase (PAP) complex method using commercial kits (Shanghai Mingdian Bioengineering Co., Ltd., Shanghai, China). The concentration of total cholesterol, high-density lipoprotein, and low-density lipoprotein in plasma was measured by peroxidase-anti-peroxidase (PAP) complex method using commercial kits (Shanghai Mingdian Bioengineering Co., Ltd., Shanghai, China). The concentration of total cholesterol, high-density lipoprotein, and low-density lipoprotein in plasma was measured by peroxidase-anti-peroxidase (PAP) complex method using commercial kits (Shanghai Mingdian Bioengineering Co., Ltd., Shanghai, China). The concentration of total cholesterol, high-density lipoprotein, and low-density lipoprotein in plasma was measured by peroxidase-anti-peroxidase (PAP) complex method using commercial kits (Shanghai Mingdian Bioengineering Co., Ltd., Shanghai, China).

2.3.3. Microsome preparation, total protein content, and CYP7A1 activity

Approximately 500 mg liver tissue were minced in Tris-HCl buffer (0.1 M Tris-HCl, 0.1 M NaCl, 0.1 M NaHCO3, 0.1 M NaH2PO4, 0.1 M NaOH, pH 7.4) and homogenized in 4 volumes of the same buffer containing 100 mg/ml of protease inhibitors (Roche Diagnostics, Mannheim, Germany). The homogenate was centrifuged at 1000g for 5 min, and the supernatant was centrifuged at 105000g for 1 h. The supernatant was dialyzed into Tris-HCl buffer (0.1 M Tris-HCl, 0.1 M NaCl, 0.1 M NaHCO3, 0.1 M NaH2PO4, 0.1 M NaOH, pH 7.4) and stored at -80 °C. The protein concentration was determined using Bradford's method (Bio-Rad, Richmond, CA, USA). CYP7A1 activity was determined using a CYP7A1 ELISA kit (Sigma-Aldrich, St. Louis, MO, USA).

activity of CYP7A1 was measured using a commercial fish CYP7A1 ELISA kit (purchased from R&D, Minneapolis, USA). Duplicate analyses were conducted for each sample.

2.3.4. RNA extraction, cDNA synthesis and partial sequence cloning of HMG-CoAr gene

Total RNA was extracted from turbot liver using Trizol Reagent (Invitrogen, USA). The quantity and quality of isolated RNA was determined by spectrophotometry with a Nano Drop® ND-1000 spectrophotometer (Nano-Drop Technologies, Wilmington, DE, USA), and electrophoresed on a 1.2% denaturing agarose gel to test the integrity. Three microgram (3 µg) of total RNA was subjected to reverse transcription by PrimerScript® RT Enzyme using Oligo-dT primer (Takara, Japan) in 10 µl of reaction mixture. The cDNA was amplified using HMG-CoAr specific primers (5'-GCACATCTACTTCCARTTYCA RA A-3' and 5'-CGGCATGCAGCCGAAR CARCACA-3') and Taq polymerase (Takara, Japan). The PCR products were purified using a PCR purification kit (Qiagen, Crawley, UK) and sequenced using the same primers. The sequence was compared with the GenBank database (http://www.ncbi.nlm.nih.gov/blast) using the BLAST program. The HMG-CoAr gene was cloned into the pMD-18T vector (Takara, Japan) and transformed into the competent cells of Escherichia coli DH5α. The recombinants were identified through blue-white color selection in ampicillin-containing LB plates and confirmed by PCR. Three positive clones in each PCR fragment were sequenced in both directions and these resulting sequences were verified and subjected to cluster analysis in NCBI.

2.3.5. Real-time PCR analysis of HMG-CoAr mRNA expression

Total RNA from individual liver samples, using 3 individuals per tank, was extracted from turbot liver using Trizol Reagent (Invitrogen, USA). The quantity and quality of isolated RNA was determined by spectrophotometry with a Nano Drop® ND-1000 spectrophotometer (Nano-Drop Technologies, Wilmington, DE, USA), and electrophoresed on a 1.2% denaturing agarose gel to test the integrity. Three microgram (3 µg) of total RNA was subjected to reverse transcription by PrimerScript® RT Enzyme using Oligo-dT primer (Takara, Japan) in 10 µl of reaction mixture. The cDNA was amplified using HMG-CoAr specific primers (5'-GCACATCTACTTCCARTTYCA RA A-3' and 5'-CGGCATGCAGCCGAAR CARCACA-3') and Taq polymerase (Takara, Japan). The PCR products were purified using a PCR purification kit (Qiagen, Crawley, UK) and sequenced using the same primers. The sequence was compared with the GenBank database (http://www.ncbi.nlm.nih.gov/blast) using the BLAST program. The HMG-CoAr gene was cloned into the pMD-18T vector (Takara, Japan) and transformed into the competent cells of Escherichia coli DH5α. The recombinants were identified through blue-white color selection in ampicillin-containing LB plates and confirmed by PCR. Three positive clones in each PCR fragment were sequenced in both directions and these resulting sequences were verified and subjected to cluster analysis in NCBI.

for each PCR assay. A four-fold serial dilution was used to assess PCR efficiencies for each assay, quantifying six concentrations. The primer amplification efficiency was optimized for each pair of primers, resulting in: 0.9847 for HMG-CoAr and 0.9772 for β -actin. The absolute ΔC_T value (HMG-CoAr C_T – β -actin C_T) of the slope is 0.0187, which indicated the $\Delta\Delta C_T$ calculation for the relative quantification of target genes might be used. The expression level of HMG-CoAr was calculated by $2^{-\Delta\Delta C_T}$ method, and the value stood for n-fold difference relative to the calibrator (Livak and Schmittgen, 2001).

2.4. Calculations and statistical methods

Growth parameters were calculated as follows:

Weight gain rate, WGR (%) = $100 \times [(\text{final body weight} - \text{initial body weight}) / \text{initial body weight}]$.

Feed intake, FI (%/d) = $100 \times \text{total amount of the feed consumed (g)} / [(\text{initial body weight} + \text{final body weight}) / 2] / \text{days}$.

WGR
FI
FCR
PPV
EPV
SR
CF
HSI
VSI
SPSS11.5软件用来数据评估。所有数据用单因素方差分析(ANOVA)然后T检验。当P<0.05时为差异显著。

Viserosomatic index, VSI (%) = $100 \times (\text{visceral weight} / \text{body weight})$.

Software SPSS, 11.5 was used for all statistical evaluations. All data were subjected to one-way analysis of variance (ANOVA) followed by Tukey's test. Differences were regarded as significant when $P < 0.05$.

3. Results

3.1. Survival rate, growth performance and somatic indexes

Survival rate was higher than 96% in all treatments (Table 2), and no significant difference was found among dietary treatments ($P > 0.05$). FI of fish fed T-1.0% diet was significantly lower than that of

存活率高于96%，处理间无显著差异。
T组FI显著低于其他组，C组FI显著高于其他组。
FBW和WGR在补充了胆固醇，牛磺酸或二者后显著增加，TC组的生长最好。
FCR在T组合TC组显著低于对照组。PPV,EPV,CF,VSI在各处理中无显著差异。

fish fed the control diet ($P < 0.05$). No significant differences were detected among dietary treatments with respect to PPV, EPV, CF, or VSI ($P > 0.05$).

3.2. Body composition

The whole-body lipid content in fish fed C-1.0% diet was significant. C组的总体脂肪含量显著高于对照组，而T组显著低于对照组。这几组的鱼体灰分含量显著低于对照组。在几个饲料处理之间的总体蛋白含量没有显著差异。

3.3. Plasma, liver and feces cholesterol

Fish fed C-1.0% diet showed significantly higher plasma TC, FC, cholesterol esters, HDL-C and LDL-C levels compared to the control diet, ($P < 0.05$, Table 4). The plasma TC, FC and LDL-C levels were

Table 2
Growth performance and survival of turbot fed the experimental diets (n = 3).

	Control	C-1.0%	T-1.0%	TC	Pooled S.E	F value	P value
IBW ¹	5.83	5.84	5.84	5.84	0.0048	0.083	0.967
FBW ²	24.3 ^a	26.7 ^{bc}	26.1 ^b	27.7 ^c	0.4096	12.853	0.002
WGR ³	316.9 ^a	357.1 ^{bc}	346.4 ^b	374.2 ^c	7.0227	10.763	0.004
FI ⁴	1.89 ^b	1.96 ^c	1.83 ^a	1.91 ^b	0.0151	17.948	0.001
FCR ⁵	0.97 ^b	0.97 ^b	0.93 ^a	0.93 ^a	0.0064	8.133	0.008
PPV ⁶	31.4	31.2	31.0	32.7	0.3126	1.824	0.221
EPV ⁷	26.1	26.0	25.6	26.9	0.3019	1.629	0.247
SR ⁸	99.2	99.2	96.7	96.7	0.8613	0.632	0.615
CF ⁹	3.21	3.29	3.28	3.32	0.0381	0.285	0.835
HSI ¹⁰	1.39 ^{ab}	1.76 ^b	1.09 ^a	1.50 ^{ab}	0.1072	2.207	0.165
VSI ¹¹	5.62	5.67	5.51	5.50	0.1035	0.124	0.943

Values in the same row with no common superscripts are significantly different ($P < 0.05$).

¹ IBW, initial body weight.

² FBW, final body weight.

³ Weight gain rate, WGR (%) = $100 \times [(\text{FBW} - \text{IBW}) / \text{IBW}]$.

⁴ Feed intake, FI (%/d) = $100 \times \text{total amount of the feed consumed (g)} / [(\text{IBW} + \text{FBW}) / 2] / \text{day}$.

⁵ Feed conversion rate, FCR = total amount of the feed consumed (g) / weight gained (g).

⁶ Protein productive value, PPV (%) = $100 \times (\text{whole-body protein gain} / \text{protein consumption})$.

⁷ Energy productive value, EPV (%) = $100 \times (\text{whole-body energy gain} / \text{energy consumption})$.

⁸ Survival rate (SR) (%) = $100 \times (\text{final fish number} / \text{initial fish number})$.

⁹ Condition factor, CF = $100 \times \text{fish weight} / (\text{body length})^3$.

¹⁰ Hepatosomatic index, HSI (%) = $100 \times (\text{liver weight} / \text{body weight})$.

¹¹ Viserosomatic index, VSI (%) = $100 \times (\text{visceral weight} / \text{body weight})$.

C组比对照组表现出更高的血浆TC,FC,胆固醇酯, HDL-C,LDL-C水平。血浆TC,FC,LDL-C水平在TC组中显著比C组低。肝TC和胆固醇酯水平在C组和TC组中显著比对照组高。肝FC水平在各组中没有显著差异。C组和TC组粪便TC显著高于对照组，在各饲料处理中粪便中TBA没有显著差异。

3.4. Activity of CYP7A1

The fish fed T-1.0% diet had the highest activity of CYP7A1 ($P < 0.05$). T组的CYP7A1活性最高，C组CYP7A1活性显著高于对照组。在大菱鲆肝脏中的CYP7A1活性在对照组和TC组中没有显著差异。

3.5. Cloning of partial cDNA sequence of the HMG-CoAr and expression of HMG-CoAr in liver

The PCR product amplified by the degenerate primers was 331 bp, and its nucleotide sequences was significantly homologous to yellowtail *Seriola quinqueradiata* (GenBank accession number: AB218826)

Table 3

Proximate composition in whole body of turbot fed the experimental diets (% diet on wet basis; n = 3).

	Control	C-1.0%	T-1.0%	TC	Pooled S.E	F value	P value
Moisture	76.5	76.2	76.9	76.4	0.1209	2.548	0.129
Crude protein	15.1	15.1	15.0	15.2	0.0756	0.374	0.774
Crude lipid	4.9 ^b	5.4 ^c	4.6 ^a	4.9 ^b	0.1146	3.163	0.086
Ash	3.5 ^b	3.2 ^a	3.3 ^a	3.2 ^a	0.0391	6.435	0.016

Values in the same row with no common superscripts are significantly different ($P < 0.05$).

Table 4
Lipid profiles in plasma, liver and feces of turbot fed the experimental diet (n = 3).

	Control	C-1.0%	T-1.0%	TC	Pooled S.E	F value	P value
Plasma (mmol/L)							
Total cholesterol	3.27 ^a	7.13 ^c	2.66 ^a	5.63 ^b	0.5594	67.025	0.000
Free cholesterol	2.03 ^a	3.59 ^c	1.51 ^a	2.89 ^b	0.2566	18.958	0.001
Cholesterol esters	1.23 ^a	3.54 ^b	1.15 ^a	2.74 ^b	0.3287	18.506	0.001
HDL-C ¹	2.09 ^{ab}	4.46 ^c	1.85 ^a	3.42 ^{bc}	0.3699	7.664	0.010
LDL-C ²	1.43 ^a	2.88 ^b	1.42 ^a	2.10 ^a	0.2080	8.484	0.007
HDL-C/LDL-C	1.46	1.52	1.31	1.66	0.1847	0.376	0.798
Liver (g/kg wet liver)							
Total cholesterol	2.97 ^a	4.03 ^b	3.44 ^{ab}	4.07 ^b	0.1799	3.720	0.061
Free cholesterol	1.22	1.38	1.22	1.41	0.0388	2.340	0.150
Cholesterol esters	1.75 ^a	2.65 ^b	2.22 ^{ab}	2.66 ^b	0.1501	3.510	0.069
Faeces (g/kg dry matter)							
Total cholesterol	5.0 ^a	19.8 ^b	5.1 ^a	19.9 ^b	2.2487	134.14	0.000
Bile acid	0.72	0.82	0.73	0.85	0.0299	1.240	0.357

Values in the same row with no common superscripts are significantly different (P<0.05).

¹ HDL-C, high-density lipoprotein cholesterol.

² LDL-C, low-density lipoprotein cholesterol.

(identities number: (GenBank HMG-CoA compared

用兼并引物扩增的PCR产物是331bp, 并且其核苷酸序列与黄尾鱼, 欧洲鲈和大西洋鲑的同源性很高, HMG-CoAr mRNA水平在TC组显著低于C组。

expression S. salar 80%). TC diet

4. Discussion

FM contains approximately 500–700 mg taurine per 100 g dry matter, while plant proteins, such as SBM, contain only trace amounts of taurine (Yamamoto et al., 1998). Therefore, replacement of FM with plant protein source results in lower dietary levels of taurine. It has been reported that growth and feed efficiency of red sea bream (Takagi et al., 2006), cobia (Lunger et al., 2007), rainbow

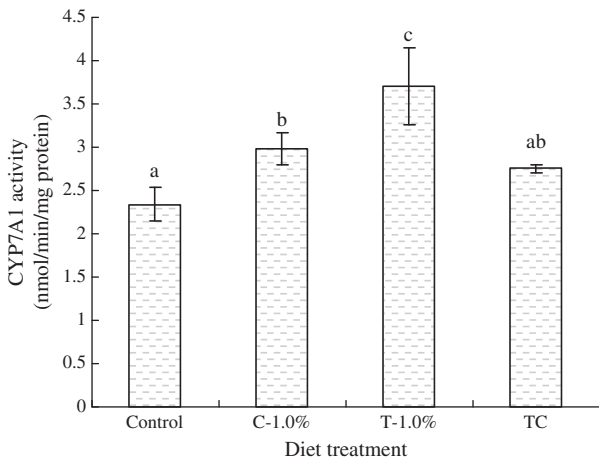


Fig. 1. Activity of cholesterol 7 α hydroxylase (CYP7A1) in liver of turbot *Scophthalmus maximus* L. fed the experimental diets. Data of the four experimental diets are presented as mean \pm S.E. (n = 3). Different letters above the bars denote significant differences between diet groups at the P<0.05 level.

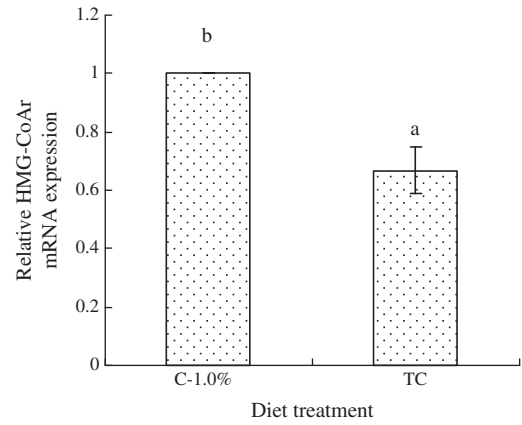


Fig. 2. Relative 3-hydroxy-3-methylglutaryl-Coenzyme A reductase (HMG-CoAr) mRNA levels in liver of turbot *Scophthalmus maximus* L. after being fed the experimental diets including C-1.0%, and TC diets. HMG-CoAr mRNA levels were evaluated by real-time quantitative PCR and expressed relative to β -actin levels. Data of the two experimental diets are presented as mean \pm S.E. (n = 3). Different letters above the bars denote significant differences between diet groups at the P<0.05 level.

trout (Gaylord et al., 2006, 2007) and yellowtail (Takagi et al., 2008) fed low levels of FM in diets based on high plant protein sources are improved by dietary taurine supplementation. In the present study, fish fed high plant protein diets with 1.14% taurine (T-1.0%) had significantly higher WGR compared to fish fed diet with 0.11% taurine (control). Thus, juvenile turbot appears to have a requirement for taurine when fish was fed high plant protein including SBM and WGM. This result was supported by the findings of Conceição et al. (1997) who reported that growth rates were positively correlated with taurine levels in fish. 鱼粉每100g干物质中包含大约500-700mg牛磺酸, 但是植物蛋白, 如SBM, 只含有很少的牛磺酸(Yamamoto et al., 1998)。因此, 用植物蛋白源替代鱼粉会导致饲料牛磺酸水平下降。已有报道, 真鲷 (Takagi et al., 2006), 军曹鱼(Lunger et al., 2007), 虹鳟 (Gaylord et al., 2006, 2007) 和 黄尾鱼(Takagi et al., 2008) 饲喂低鱼粉高植物蛋白饲料的生长和饲料效率由于饲料牛磺酸的补充而提高。在本研究中, 高植物蛋白饲料含1.14%牛磺酸WGR显著高于饲料含0.11%牛磺酸(对照组)。由此, 大菱鲆幼鱼似乎在当饲喂高植物蛋白包括SBM和WGM时对牛磺酸有需求。这个结果由Conceição et al.(1997)的发现所支持, 他报道在大菱鲆幼鱼中GR与牛磺酸水平很相关。牛磺酸是氨基酸衍生物并且拥有刺激鱼类摄取的特点, 例如低分子量, 氮含量, 水溶性和酸性(Carr, 1982)。牛磺酸在欧洲鲈(Martinez et al., 2004), 虹鳟(Gaylord et al., 2006), 黄尾鱼(Takagi et al., 2008), 牙鲆(Kim et al., 2005b)中用作主要的诱食剂。但是, 在本实验中, T组的FI显著低于对照组。这就可能是牛磺酸的促进生长作用不是由于饲料摄入的增多。生长的促进可以归因于一些与辅助作用相关的生物现象, 如膜建立, 抗氧化, 解毒作用(Wright et al., 1986), 渗透调节(Thurston et al., 1980), 与胆汁酸结合(Goto et al., 1996)。但是, 当与饲喂鱼粉饲料时, 如之前报道的在虹鳟(Gaylord et al., 2006)和牙鲆(Chatzifotis et al., 2008)中(一样, 牛磺酸补充并没增加生长。似乎表明饲料牛磺酸的水平, 与一些内源性产物结合, 对于维持生长是充足的。 is sufficient to maintain growth.

Similarly, cholesterol is rich in FM, but only low levels are present in most plant sources (Cheng and Hardy, 2004; Deng et al., 2010). 同样的, 胆固醇在鱼粉中也很丰富, 在植物源中只存在少量 (Cheng and Hardy, 2004; Deng et al., 2010)。也有报道表明, 在基于植物蛋白源的饲料中叉尾鲷(Twibell and Wilson, 2004), 牙鲆(Chen, 2006; Deng et al., 2010)的生长, 能由饲料胆固醇补充提高。但是, 大西洋鲑(Bjerkeng et al., 1999), 杂交条纹鲈(Sealey et al., 2001)和牙鲆(Deng et al., 2010)在鱼粉饲料上补充胆固醇抑制了生长。因此, 胆固醇补充可能是在植物蛋白基础上有需求。另外, 在本实验中, 胆固醇好牛磺酸的组合比T组和C组或对照组提高了大菱鲆幼鱼的生长。这样, 观察到的胆固醇和牛磺酸系统促生长效果可以被解释为使含SBM和WGM的饲料(缺乏牛磺酸和胆固醇)又达到营养平衡。 bot compared to fish fed other diets containing either 1-1.0% (P<0.05) or

C-1.0% ($P>0.05$) or the control ($P<0.05$) (Table 2). Thus, the observed synergistic growth-promotion effect of combined cholesterol and taurine can be explained by a re-equilibration of the diet containing SBM and WGM, which was short of taurine and cholesterol.

In the present study, the whole-body lipid content in fish fed C-1.0% diet was significantly higher than the control diet while that in fish fed T-1.0% diet was significantly lower than the control diet. Dietary addition of 1.0% cholesterol significantly resulted in higher overall lipid content, but T group significantly lower than control. In SBM diet, addition of 1% cholesterol significantly led to higher apparent digestibility and liver lipid content (Chen, 2006). In this study, high plant protein diet containing 1.25% cholesterol can improve the utilization of total lipid, and compared with control, Gaylord et al. (2006) reported that addition of 0.5–1.5% cholesterol to plant-based diet had no effect on total lipid content. In plant-based diet, addition of cholesterol to total lipid storage and theoretical need for further research.

Taurine may interfere with lipid metabolism of fish, which had been shown in other animals where taurine exhibits hypolipidemic effects (Militante and Lombardini, 2004). Furthermore, the plasma cholesterol-lowering effects of taurine supplementation have been well demonstrated in rats and mice (Militante and Lombardini, 1999), mice (Murakami et al., 1996; Yokogoshi et al., 1999), and mice with increased cholesterol (Murakami et al., 2002), and humans (Mizushima et al., 1996) in which taurine supplementation significantly increased CYP7A1 mRNA expression, which is the rate-limiting step in bile acid synthesis (Murakami et al., 1999, 2005; Yokogoshi et al., 1999). In this study, taurine supplementation significantly increased CYP7A1 mRNA expression, which is the rate-limiting step in bile acid synthesis (Murakami et al., 1999, 2005; Yokogoshi et al., 1999). This suggests that taurine supplementation may increase bile acid synthesis by improved activity of CYP7A1 because of unchanged bile acid concentration and CYP7A1 activity (Table 4, Fig. 1).

The liver cholesterol-lowering effects of taurine supplementation have been well established in rats or mice fed high cholesterol diet (Chen et al., 2005; Gandhi et al., 1992; Park and Lee, 1998; Yan et al., 1993). The cholesterol metabolism is affected by *de novo* synthesis of cholesterol in liver of fish based on fish fed high cholesterol diet. A possible mechanism is that taurine supplementation led to a prior storage of cholesterol in liver of fish fed high plant protein based on high cholesterol diet, and then caused plasma cholesterol-lowering effects.

In conclusion, the results of the present study showed that supplementation of either cholesterol or taurine and their combination in diet significantly improved growth performance of fish fed high cholesterol diet. In high cholesterol diet, addition of 1% cholesterol significantly improved growth performance of fish fed high cholesterol diet. In high cholesterol diet, addition of 1% cholesterol significantly improved growth performance of fish fed high cholesterol diet. In high cholesterol diet, addition of 1% cholesterol significantly improved growth performance of fish fed high cholesterol diet.

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