Shanghai 2016/7/27--2016/7/28

The 2nd workshop on the study of lipid nutrition and metabolism in aquatic animals



21 participating units, nearly 70 participants;

17 invited lectures, 22 abstracts, 29 publications



2. Invited lectures

1) Sustainable feeds and n-3: impacts and sloutions——Douglas Tocher

- 2) Recent advances in the study of regulatory mechanisms of LC-PUFA biosynthesis in rabbtifish——Li Yuanyou
- 3) Functional characterization and nutritional regulation of putative ElovI5 and ElovI4 elongases in large yellow croaker——Ai Qinghui
- 4) Is ARA an effective nutrient in depressing lipid accumulation and improving health status in grass carp—Ji Hong
- 5) Mechanisms and metabolic regulation of PPARa activation in Nile tilapia——Du Zhenyu
- 6) The role of fatty acid desaturases on highly unsaturases fatty biosynthesis in loach——Gao Jian

- 7) n-3 fatty acids impact on cardiometabolic diseases and obesity ——Michel Nance
- 8) n-3 essential fatty acids in Nile tilapia: Bioconversion of dietary linolenic aicd—Pan Qing
- 9) From marine environment to freshwater: in vivo synthesis of highly unsaturated fatty acids in *Litopenaeus vannamei*——Li Erchao
- 10) Effects of partial replacement of fish oil with repeseed oil on growth, fatty acid profile and expression of immune related gene for two sizes of yellow croaker—Huang Xuxiong
- 11) The research of changed energy metabolism pattern of IGF-I overexpression Crucian Carp——Li Dongliang
- **12)** The application of Maxi-mil in aquatic feed——Han Zejian

- 13) Effects and mechanism of endoplasmic reticulum stress and the related signaling pathway on minerla-induced alteration in lipid metabolism in fish——Luo Zhi
- 14) n-3 rich diet as a nutritional strategy to counteract overactivation

Focusing on the fatty acids metabolism (13); Focusing on the n-3 PUFA (9); Focusing on the n-3 PUFA biosynthesis (6)

16) Cloning, tissue expression of the FABP gene, and effects of dietary phospholipid levels on FABP and vitellogenin mRNA expression in the female swimming crab—Zhou Qicun

17) The relationship among the host intestinal microbiota and environment——Zhang Meiling





Tired

Gratified

The 2nd workshop on the study of lipid nutrition and metabolism in aquatic animals

Thanks for your attention!

Reading report





Biochimica et Biophysica Acta (BBA) -Molecular and Cell Biology of Lipids

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Mechanisms and metabolic regulation of PPARα activation in Nile tilapia (Oreochromis niloticus)

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1. Abstract



Objective: to illuminate the molecular mechanisms and metabolic regulation of PPARα activation in fish

- **Contents:** a. Gene cloning and tissue expression of mRNA of Nile tilapia (Nt) PPARα.
 - b. Influence of the agonist (fenofibrate) and fasting on NtPPARα expression in vitro and in vivo.
 - c. Investigating the metabolic regulatory effects of NtPPAR α (feed with fenofibrate or fasted).
 - d. Investigating the global regulatory effects of fenofibrate and fasting by hepatic transcriptomic study



To spare the protein consumption, high-lipid diets are currently being used in aquaculture.





In mammals, PPAR α has been recognized as a master regulator of lipid metabolism, and a transcriptional factor activated by binding of ligands.

PPARa is a key regulator in lipid catabolism



(Costet, P., 1998)





Fibrate-induced PPARa activation in fish

PPARa ligands, such as fibrates, have moderate lipid-lowering effects and induce PPARa mRNA expression in torafugu, yellow catfish, rainbow trout gill-W1 cells, and zebrafish primary cells. However, bezafibrate and clofibric acid have no effect on PPARa expression in fathead minnow and grass carp.



(Du et al., 2004; Liu et al., 2005; Ibabe et al., 2005; Kondo et al., 2007; Du et al., 2008; Weston et al., 2009; Guo et al., 2015; Zheng et al., 2015;)



PPARα molecules have been cloned in many teleosts, but its activation mechanisms have not been well demonstrated.

PPARa has been cloned in some fish species

Species	Highly expression	Literatures
牙鮃 Bastard halibut	Stomach, Liver, Intestine	(Cho et al. 2012)
章鱼Grass carp	Liver	(He et al. 2012)
虹崎 Rainbow trout	Adipose tissue, Muscle, Ovary	(東京取得, 2012)
军曹鱼 Cobia	Muscle, liver, Heart	(Taal et al. 2008)
解節 Mullet	Liver	(Raineeand et al. 2006
团头粉Black bream	Muscle, Muscle, Liver, Heart, Brain, Intestine	(2hao et al. 2011)
黄颜值 Yellow catfish	Liver, Heart, Muscle	
明形系統的第 Pompano	Brain, Kidney, Intestine, Pleen	(方科科, 2015)
尼罗罗非鱼 Nile Tilapia	Liver	(Ning at al., 2016)

The mechanisms of the PPARα activation and the regulatory effects on lipid metabolism has not been thoroughly investigated in fish.





Why choose the Nile tilapia as fish model for lipid metabolism?



Nile tilapia (Oreochromis niloticus) is an important aquaculture species worldwide and a good fish model for metabolic studies because of its rapid growth, high disease and stress resistance, and because its entire genome is available.







2.2. Gene cloning, identification and tissue expression of mRNA of NtPPARα

2.3. Catabolism rate of intraperitoneally injected [1-14C] palmitate

At the end of 4 week, four fishes from each gro intraperitoneal injection Of [1-14C] palmitate (DMS)



2.4. Mitochondrial and peroxisomal palmitate oxidation in liver and muscle homogenates

At the end of the feeding trial, pieces of liver and muscle were collected from each group, diluted, homogenized, measured.





2.5. Culture of tilapia primary hepatocytes

2.6. Quantitative real time PCR and western blot analyses

2.7. Transcriptomic analysis

2.8. Histological study





3.1. NtPPAR α structure and features

NtPPAR α has 475 deduced amino acid (AA), and share high identity to known PPAR α molecules in other teleosts

The highest expression level was detected in the brain, followed by the liver and heart



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3.2. Activation of NtPPARα by fenofibrate and fasting in primary hepatocytes of Nile tilapia



(A) The mRNA expression of NtPPAR α at 12 h and 24 h. (B) The protein express and phosphorylation of NtPPAR α using western blotting. (C–D) Statistical analysis of the western blotting results

3.3. Effects of fenofibrate and fasting on body and serum composition



(A) Weight gain rate; (B) hepatic-somatic index (HSI); (C) mesenteric fat index; (D) hepatic triglyceride (TG) content; (E) hepatic glycogen content; (F) absolute amount of fatty acids in liver; (G) histological characteristics of liver.



3.3. Effects of fenofibrate and fasting on body and serum composition



These results indicate that fenofibrate has hypolipidemic effects in tilapia and mainly targets the liver.



A) Triglyceride; B) free acids (FFAs); C) β-hydroxybutyrate; D) glucose; E) insulin; F) lactate.



3.4. The hypolipidemic effects caused by fenofibrate are due to increased fatty acid β -oxidation in tilapia



(A-C) The mitochondrial and peroxisomal β -oxidation of [1-14C] palmitate in the homogenates of liver and muscle; (D) The conversion of intraperitoneally injected [1-14C] palmitate into carbon dioxide; (E) The relative copy number of mtDNA cytochrome b

The results suggest that increased lipid breakdown and decreased adipogenesis are the main metabolic consequences after NtPPARα is activated, the magnitude of NtPPARα activation differed between the fenofibrate and fasting treatments, but dephosphorylation of NtPPARα was likely the common activation * mechanism in both treatments.



A–C) Lipid catabolism-related genes: PPARα, CPT1a and ACO;D–E) Lipid anabolism related genes: PPARγ and FAS; F) western-blot of PPARα, pPPARα (S12) and GAPDH; G) relative WB quantification of PPARα; H) relative WB quantification of pPPARα (S12).

3.6. Global regulatory effects of fenofibrate and fasting on tilapia in a hepatic transcriptomic study



(A) Venn diagram representing mRNA transcripts differentially expressed (DEGs) in the liver from fasting and fenofibrate treatment compared to that of contorl. (B–C) Distribution by categories of differentially expressed genes and in the liver when compared fenofibrate and fasting treatment to that of control.





Specifically, significant changes were detected in genes of key enzymes for lipid hydrolysis (ATGL), adipogenesis (DGAT), and FA β -oxidation (ACS and CPT-I) in the transcriptomic measurements for the fasting group, whereas none of these genes changed significantly in the fenofibrate group.



4. Discussion



4.1. PPAR α activation and its mechanisms in different species

In mammals, two potential mechanisms for ligand-induced PPARα activation have been proposed, including increased receptor expression at the transcriptional or protein level and modifications of phosphorylation status (phosphorylation or dephosphorylation).

In the present study, in vitro and in vivo evidence for the first time, illustrate that NtPPARα was activated by fenofibrate or endogenous ligands, such as fasting-induced FFAs, through increased mRNA and protein expression and decreased phosphorylation.



4. Discussion



4.2. Hypolipidemic effects and fatty acid β -oxidation through PPAR α activation in different species

The hypolipidemic effects of PPAR α activation have been widely reported and one of the main mechanisms involves increasing mitochondrial and peroxisomal FA β -oxidation. However, these induction effects vary greatly among species.

Species	effects of fenofibrate	
species	mitochondrial activity	peroxisomal activity
Rodents	increased	increased
Human	mildly induce	mildly induce
Monkey	mildly induce	mildly induce
Rainbow trout	increased	increased
Grass carp	increased	increased
Yellow catfish	increased	increased
Tilapia	increased	no

4. Discussion



4.3. NtPPAR α activation affects carbohydrate metabolism

Until now, there has been limited direct evidence linking pharmacological PPARa activation and glucose homeostasis. Our results show higher serum levels of glucose, insulin, and lactate in the fenofibrate group than those in the fasting group, indicating that glucose utilization decreased in response to fenofibrate. Those studies suggest that utilization of lipid and carbohydrate is balanced in fish.

5. Conclusions



Conclusions

- PPARα was activated in response to fenofibrate and fasting in Nile tilapia;
- NtPPARα activation mainly targets the liver and is relatively moderate;
- Dephosphorylation is the basal NtPPARα activation mechanism;
- NtPPARα activation increased activity and the number of hepatic mitochondria.





Thanks for your attention!