

On the utilization of dietary glycerol in carnivorous fish - Part II: insights into lipid metabolism of rainbow trout (*Oncorhynchus mykiss*) and European seabass (*Dicentrarchus labrax*)

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Supplementary Material

Supplementary Table S1 Primer sequences and accession numbers for qPCR analysis on liver of rainbow trout (*O. mykiss*).

Trout	Forward primer (5'-3')	Reverse primer (5'-3')	Accession number (Genoscope, GenBank, Sigenae)	Previously reported
Reference gene				
<i>ef1a</i>	TCCTCTTGGTCGTTTCGCTG	ACCCGAGGGACATCCTGTG	AF498320	Song et al. 2018
Lipogenesis				
<i>fas</i>	GTGATGTCGAGCTTCGTGCT	CTCCAGTGCTGACGCACCT	tcaa0001c.m.06_5.1.om.4	Song et al. 2018
<i>dgat</i>	CCGAAGTTACAGAAGGCTCC	CCGAAGTTACAGAAGGCTCC	LOC110537459	No
Lipolysis				
<i>hoad</i>	GGACAAAGTGGCACCAGCAC	GGGACGGGGTTGAAGAAGTG	tcad0001a.i.15_3.1.om	Song et al. 2018
<i>cpt1a</i>	GATGGCTTGACGTTCTCCT	AGATCCAGGAGAGTCCTCC	GSONMG00076843001	Song et al. 2018
<i>cpt1b</i>	AGGGACAGTTCGAAAATGGAG	CCAGAGAGGGAAGATGAAGA	GSONMG00076841001/GSONMG00066036001	No
Long chain PUFA biosynthesis				
$\Delta 6$ - <i>fad</i>	AGGGTGCCTCTGCTAACTGG	TGGTGTGGTGATGGTAGGG	AF301910	Song et al. 2018
<i>elovl2</i>	TGTGGTTTCCCCGTTGGATGCC	ACAGAGTGGCCATGTCCTTGT	FYV3OTN01A4WMI.s.om.10	Song et al. 2018
<i>elovl5</i>	GAACAGCTTCATCCATGTCC	TGACTGCACATATCGTCTGG	AY605100	Song et al. 2018
<i>ef1a</i> (elongation factor 1 α); <i>fas</i> (fatty acid synthase); <i>dgat</i> (diglyceride acyltransferase); <i>hoad</i> (3-hydroxyacyl-CoA dehydrogenase); <i>cpt1a</i> and <i>cpt1b</i> (carnitine palmitoyltransferase I a and b, respectively); $\Delta 6$ - <i>fad</i> (acyl-CoA 6-desaturase); <i>elovl2</i> and <i>elovl5</i> (elongation of very long-chain fatty acid protein 2 and 5, respectively).				

Song, X., Marandel, L., Skiba-Cassy, S., Corraze, G., Dupont-Nivet, M., Quillet, E., Geurden, I., and Panserat, S. (2018). Regulation by Dietary Carbohydrates of Intermediary Metabolism in Liver and Muscle of Two Isogenic Lines of Rainbow Trout. *Frontiers in Physiology* 9. doi: <https://doi.org/10.3389/fphys.2018.01579>

Supplementary Table S2 Primer sequences and accession numbers for qPCR analysis on liver of European seabass (*D. labrax*).

Seabass	Forward primer (5'-3')	Reverse primer (5'-3')	Accession number (Genoscope, GenBank, Sigenae)	Previously reported
Reference gene				
<i>ef1a</i>	GCTTCGAGGAAATCACCAAG	CAACCTTCCATCCCCTGAAC	AJ866727	Castro et al. 2015
Lipogenesis				
<i>fas</i>	ATCCTGAAGCGTAGCCTGAA	CCAAGTGGCACAGGGTACTT	MF566098.1	No
<i>dgat</i>	CAAGCCAACATCCGAGTTTT	GCTGATGACCCACTGTAGCA	FL486573	Castro et al. 2015
Lipolysis				
<i>hoad</i>	ACACTTCTGCACTGGACGTG	CTTGCTCGAGGTATGGCTTC	JQ938805.1	No
<i>cpt1a</i>	GTCTGCTCGTCTGTGGATGA	TTCTTTAGCCAAGCCCTTCA	KF857302.1	No
Long chain PUFA biosynthesis				
$\Delta 6$ - <i>fad</i>	CCTTCACTGCTTTTCATCCCAA	CCCAGGTGGAGGCAGAAGAA		Geay et al. 2010
<i>elovl5</i>	GCACGGTGGCTACAATTCT	TGTCCATGAACTCGATGAGC		Castro et al. 2015

ef1a (elongation factor 1 α); *fas* (fatty acid synthase); *hoad* (3-hydroxyacyl-CoA dehydrogenase) and C) *cpt1* (carnitine palmitoyltransferase I); $\Delta 6$ -*fad* (acyl-CoA 6-desaturase); *elovl5* (elongation of very long-chain fatty acid protein 5).

Castro, C., Corraze, G., Panserat, S., and Oliva-Teles, A. (2015). Effects of fish oil replacement by a vegetable oil blend on digestibility, postprandial serum metabolite profile, lipid and glucose metabolism of European sea bass (*Dicentrarchus labrax*) juveniles. *Aquaculture Nutrition* 21, 592-603. doi: <https://doi.org/10.1111/anu.12184>

Geay, F., Santigosa I Culi, E., Corporeau, C., Boudry, P., Dreano, Y., Corcos, L., Bodin, N., Vandeputte, M., Zambonino-Infante, J.L., Mazurais, D., and Cahu, C.L. (2010). Regulation of FADS2 expression and activity in European sea bass (*Dicentrarchus labrax*, L.) fed a vegetable diet. *Comparative Biochemistry and Physiology Part B: Biochemistry and Molecular Biology* 156, 237-243. doi: <https://doi.org/10.1016/j.cbpb.2010.03.008>