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TISSUE DISTRIBUTION OF LIPID AND FATTY ACID METABOLISM AND TRANSCRIPTION FACTORS GENES IN ADULT ATLANTIC BLUEFIN TUNA (Thunnus thynnus L.)

Betancor M.B.¹, A. Ortega², F. de la Gándara², D.R. Tocher¹ & G. Mourente³

- 1 Institute of Aquaculture, University of Stirling, FK9 4LA, Stirling, Scotland, UK
- 2 Planta Experimental de Cultivos Marinos, Instituto Español de Oceanografía (IEO), 30860, Puerto de Mazarrón (Murcia), Spain.
- 3 Departamento de Biología, Facultad de Ciencias del Mar y Ambientales, Universidad de Cádiz, 11510 Puerto Real (Cádiz), Spain (gabriel.mourente@uca.es).

To determine expression of the major lipid pathways in tissues of adult bluefin tuna 8 individuals were used for collecting samples for tissue expression of key lipid metabolism genes. Triplicate sets of samples of brain, gills, heart, kidney, spleen, liver, intestine, white muscle, red muscle, adipose tissue ovary and testis were collected. Expression of genes was determined by qPCR.

Tissue expression profiles showed that PUFA biosynthetic pathway genes were expressed in all tissues examined, highest expression in brain, liver and testis. Elongase *elovl5* showed higher expression than desaturase *fads2d6* in all other tissues, with low expression of in red muscle and ovaries.

Transcription factors, ppara and ppary showed parallel expression, with adipose tissue with the highest relative copy number, followed by intestine>testis>liver. The expression of lxr was low in liver, with highest expression in testis, brain and kidney. Similarly, rxr was poorly expressed in liver with higher expression in muscle, spleen and brain. The rank order of expression of srebp1 was brain, testis, ovary, intestines, kidney, gill, liver, white muscle, spleen, heart and red muscle. For srebp2 the highest expression was shown in brain, testis and adipose with lowest expression in heart and white muscle. Expression of fabp2 was highest in intestine, brain and heart with lower levels in liver, red muscle, adipose and kidney. fabp4 showed highest expression in ovaries with liver showing the lowest. fabp7 showed highest expression levels in brain and testis with lowest values in liver. cptl expression was highest in brain and lowest in liver. Similarly, the relative copy number for fas was highest in brain followed by gonads, gill and liver, with white muscle showing lowest expression. The expression of aco was highest in adipose and intestine, followed by liver, kidney and brain. The expression of hmgcl was highest in ovary followed by adipose, brain and testis, with lowest expression in liver. Expression of IpI was highest in testis and lowest in ovary, with liver and white muscle, adipose, heart, gills and red muscle, kidney, intestine showing intermediate levels of expression.

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