

Table S1 The Product Numbers of Commercial Assay kits.

Enzyme	Kit No.	Manufacturer
Amylase (AMS)	C016-1-1, C016-2-1	Nanjing Jiancheng Bioengineering Institute
Glucose-6-phosphate dehydrogenase (G-6-PD)	A027-1-1	Nanjing Jiancheng Bioengineering Institute
Malic enzyme (ME)	BC1125	Solarbio Life Sciences
pyruvate kinase (PK)	A076-1-1	Nanjing Jiancheng Bioengineering Institute
Acyl-CoA oxidase (ACO)	H232	Nanjing Jiancheng Bioengineering Institute
Lipase (LPS)	E1019	Applygen Technologies Inc.
Fatty acid synthase (FAS)	H231	Nanjing Jiancheng Bioengineering Institute
Carnitine-acylcarnitine translocase (CACT)	SEB657Ra	Wuhan Cloud-Clone Corp.
Acetyl-CoA carboxylase (ACC)	H232	Nanjing Jiancheng Bioengineering Institute
Superoxide dismutase (SOD)	A001-3-1	Nanjing Jiancheng Bioengineering Institute
Phospholipid hydroperoxide glutathione peroxidase (GSH-PX)	A005-1-1	Nanjing Jiancheng Bioengineering Institute
Catalase (CAT)	A007-1-1	Nanjing Jiancheng Bioengineering Institute
Lipoprotein lipase (LPL)	A067-1-1	Nanjing Jiancheng Bioengineering Institute
malondialdehyde (MDA)	A003-1-1	Nanjing Jiancheng Bioengineering Institute

Table S2 Primers Used for Detection of DEGs

Gene	Primer name	Primer sequence (5' -3')	Purpose	Amplification efficiency (%)	Correlation coefficient
<i>ghrb</i>	<i>ghrb F</i>	AGCAACCCCATATCCCCTCT	RT-qPCR	95.30	0.998
	<i>ghrb R</i>	AGAGGTAGTGGCAGGACAGG	RT-qPCR		
<i>igfbp4</i>	<i>igfbp4 F</i>	AGCAATATCCGGTGCAGTCC	RT-qPCR	101.35	0.995
	<i>igfbp4 R</i>	AGCAATATCCGGTGCAGTCC	RT-qPCR		
<i>vtgB</i>	<i>vtgB F</i>	CTGGTATGGCAGGTTGCTGA	RT-qPCR	100.08	0.998
	<i>vtgB R</i>	CTCAACCTGGAAATCCAGAAAAC	RT-qPCR		
<i>apoF</i>	<i>apoF F</i>	AGTTCCTGCACTTTCCAGCA	RT-qPCR	96.84	0.998
	<i>apoF R</i>	GTCCTGTGATTTTGTAGTTTCAGCA	RT-qPCR		
<i>apoB-100</i>	<i>apoB-100 F</i>	TCTATTTGGGAAGCAGCTCTGA	RT-qPCR	94.92	0.997
	<i>apoB-100 R</i>	GGAATGTTAGGAAGGAAGGCA	RT-qPCR		
<i>apoE</i>	<i>apoE F</i>	GCAGGGTGAAAGTGCATCGTT	RT-qPCR	103.53	0.996
	<i>apoE R</i>	AGGTTGACTATGCAGAAGGTTGA	RT-qPCR		
<i>apoEb</i>	<i>apoEb F</i>	ATGAGGACACGTTTCGACCC	RT-qPCR	101.77	0.993
	<i>apoEb R</i>	TCAGGGTGGTGATCTTTGCC	RT-qPCR		
<i>ggct</i>	<i>ggct F</i>	AGACTCACCCCTTCCTGGTT	RT-qPCR	99.25	0.999
	<i>ggct R</i>	CAACTGGCAAATCCTTCGGC	RT-qPCR		
<i>gpx1a</i>	<i>gpx1a F</i>	TGCATCAGAAACTCCCCTCAC	RT-qPCR	98.84	0.998
	<i>gpx1a R</i>	ATGGCTGTTGGAGAGATAGCC	RT-qPCR		
<i>gstt3</i>	<i>gstt3 F</i>	GAAGGAACACGGGGGAAGTG	RT-qPCR	94.54	0.992
	<i>gstt3 R</i>	CTCAAAGGGAATCCCGACCG	RT-qPCR		
β -actin	β -actin F	GCAGGAGTACGATGAGTCCG	Reference gene	99.25	0.998
	β -actin R	AGGCATGTATGCAGAGCCAG	Reference gene		

Table S3 Assembly results statistics

Length Range	Transcript	Unigene
200-300	9,934(16.75%)	8,600(23.57%)
300-500	9,177(15.48%)	6,677(18.30%)
500-1000	12,006(20.25%)	6,866(18.82%)
1000-2000	14,881(25.09%)	7,460(20.45%)
2000+	13,303(22.43%)	6,879(18.86%)

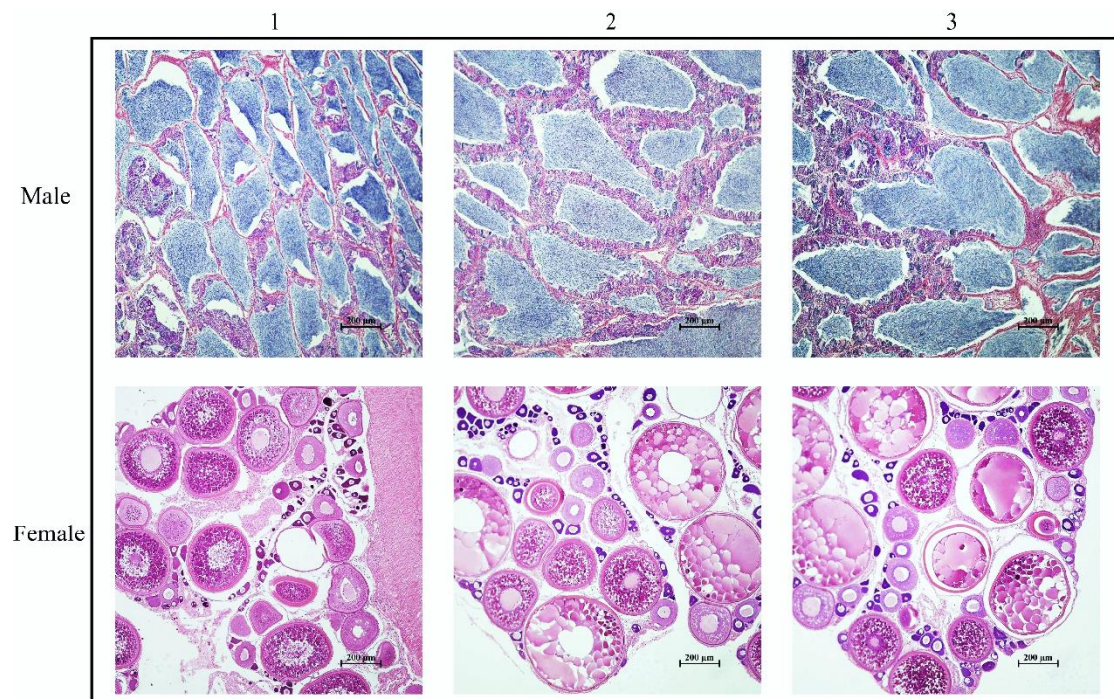


Figure S1: Histological identification of male and female yellowfin tuna gonads. Scale bar: 200 µm.