

S3 Table. Primer pair sequences used for RT-qPCR analysis.

Gene	Genbank access #	Forward primer (5' - 3')	Reverse primer (5' - 3')
RXR α /a	NM_001161551	ATTCAATGGC'ATCTCCTG	GCGGCTTAATATCCTCTG
PPAR γ	NM_131467	GGTTTCATTACGGCGTTCAC	TGGTTCACGTCACTGGAGAA
C/EBP α	NM_131885	AACGGAGCGAGCTTGACTT	AAATCATGCCATTAGCTGC
C/EBP β	NM_131884.2	CTACCTCGCTTCCACAGCA	TCGCTGTAGATTCCCAGTGC
SREBP1	NM_001105129.1	CAGAGGGTGGGCATGCTGGC	ATGTGACGGTGGTGCCGCTG
ChREBP	XM_001338467.2	ATCCCAGGGAGATCGCTCAT	TGCTGCTGGTTAAGGGAGTG
DGAT2	NM_001030196	TGGGGCTTTTTGTAACCTCG	TCTTCCTGGTGCACAGTCC
FASn	XM_682295	ATCTGTTCCTGTTTCGATGGC	AGCATATCTCGGCTGACGTT
ACC α	NM_001271308.1	AGAGAGGGCAGGTTTTACCA	GCCATCATACGAGAGCAACA
ACOX1	BC097101.1	GCACGGATGTGTGTACCGTGC	GCGTCCAGAGCCCCTTGACCT
11 β -HSD2	NM_212720.2	CCTTCAGGTGAGCATCCCTT	CCTCCAGAAGGTTTGGGGAC
11 β -HSD3 α	NM_200323.2	ATGCCATCGAACAGCTTGGA	AATTCACCTCTAGTAGCCATCG
IGF-I	NM_131825.2	GGCAAATCTCCACGATCTCTAC	CGGTTTCTCTTGTCTCTCTCAG
IGF-I α	NM_131433.1	TCGTGTGTGGAGAAGATGGC	ATGGTGTCTTGTGAAGAGCC
β -actin	NM_181601.3	ACTGTATTGTCTGGTGGTAC	TACTCCTGCTTGCTAATCC
rpl8	NM_200713.1	TTGTTGGTGTGTTGCTGGT	GGATGCTCAACAGGGTTCAT