

Table S1. Oligonucleotide sequences used in RT-PCR analysis.

Target gene	Direction	Sequence (5'-3')	Annealing temperatures	GenBank accession
<i>Pparγ</i>	forward	GC GG GCTGAGAAGTCACGTT	58 °C	NM_001127330.3
	reverse	CCATCACGGAGAGGTCCACA		
<i>C/ebpα</i>	forward	TACCGAGTAGGGGGAGCA	55 °C	NM_001287514.1
	reverse	TCATTTTCTCACGGGGCC		
<i>Fasn</i>	forward	AGAAGCCATGTGGGAAGATT	56 °C	NM_007988.3
	reverse	AGCAGGGACAGGACAAGACAA		
<i>Acly</i>	forward	CTTGGGCCGGAACAAAAA	55 °C	NM_001199296.1
	reverse	GCCGAGGTGGTGCAGAT		
<i>Acca</i>	forward	AGAAACCCGAACAGTGGAACT	55 °C	NM_133360.3
	reverse	AGGTAGCCCTTCACGGTTAAA		
<i>Fabp4</i>	forward	GGGAACCTGGAAGCTTGTCT	55 °C	NM_001409514.1
	reverse	ACTCTCTGACCGGATGGTGA		
<i>Scd1</i>	forward	TTCTTGCATACACTCTGGTGC	55 °C	NM_009127.4
	reverse	CGGGATTGAATGTTCTGTCGT		
<i>Dgat1</i>	forward	GGCCC AAGGTAGAAGAGGAC	55 °C	NM_010046.4
	reverse	GATCAGCATCACCAACACACC		
<i>Pgc1α</i>	forward	AAGTGGTAGCGACCAATCG	55 °C	NM_008904.3
	reverse	AATGAGGGCAATCCGTCTCA		
<i>Cpt1a</i>	forward	CTCCGCCTGAGCCATGAAG	55 °C	NM_013495.2
	reverse	CACCAGTGATGCCATTCT		
<i>Rplp0</i>	forward	AGGTCCCTCCTGGTGAAC	52 °C	NM_007475.5
	reverse	GTGCTGATGGCAAGAAC		
<i>Gapdh</i>	forward	AGTATGACTCCACTCACGGCAAAT	56 °C	NM_001411843.1
	reverse	GTCTCGCTCCTGGAAGATGGT		
<i>nDNA H19</i>	forward	GAACAGAACGATTCTAGGCTGG	55 °C	NR_130974.1
	reverse	TTCTAAGTGAATTACGGTGGGTG		
<i>MT-Cyb</i>	forward	ACGCAAACGGAGCCTCAATA	55 °C	NC_005089.13
	reverse	TGTGGCTATGACTGCGAACAA		
<i>MT-Atp6</i>	forward	CCAATGGCATTAGCAGTCGG	55 °C	NC_005089.1
	reverse	AATGGTAGCTGTTGGTGGC		

Note: peroxisome proliferator-activated receptor- γ , Ppar- γ ; CCAAT/enhancer-binding protein α , C/ebp α ; fatty acid synthase, Fasn; ATP-citrate lyase, Acly; acetyl-coA Carboxylase α , Acca; fatty acid-binding protein 4, Fabp4; stearoyl-Coenzyme A desaturase 1, Scd1; diacylglycerol O-acyltransferase 1, Dgat1; Peroxisome proliferator-activated receptor γ coactivator 1 α , Pgc1 α ; carnitine palmitoyltransferase 1A, Cpt1a; Ribosomal protein lateral stalk subunit P0, Rplp0; Glyceraldehyde-3-phosphate dehydrogenase, Gapdh. Nuclear DNA H19, nDNA H19; Mitochondrially Encoded Cytochrome B, MT-Cyb; Mitochondrially Encoded ATP Synthase Membrane Subunit 6, MT-Atp6.