

**Supplementary Table S6:** Primer sequences used in real-time qRT-PCR analyses

Probeset ID	Gene symbol	Forward (5'-3')	Reverse (3'-5')
	18s rRNA_v1	ccgtcggcatgtattagctc	accgtctcctacctggttga
Ssc.30843.1.A1_at	ADAMDEC1	ttctgcttgccctgactcgta	aggtgagccatgcagacttc
Ssc.18933.1.A1_at	ADH4	tggatgcactggtgtctcat	tcatggctcattgttcatca
Ssc.4891.1.A1_at	AGPAT9	cgactcaagcgggagtaaga	gtcgttacgctacggagtc
Ssc.6943.1.A1_at	ANGPT1	tgatattttggatgcagttttca	catgtgtattgttttaagggcttg
Ssc.8980.1.A1_at	ANGPTL4	actcttcacctgcctgagga	gcacccacatacaacagcac
Ssc.1039.1.S1_at	APOC3	gacacctcccttctggacaa	ggactccttcacgtggtta
Ssc.833.1.S1_at	C1QA	cttccaggtggtgtccaagt	tggatccagacctgtctcc
Ssc.17026.1.Sa_at	C4A	gcctctttgagacgagcatc	ctaagtgaaggcggcaagag
Ssc.10460.1.S1_at	CFB	aaggaaagctgcctgtgaga	agtgttggggcagcatagg
Ssc.864.1.S1_at	CFH	cagtctatcctccgggaaca	gcattttggtggttctgacc
Ssc.16157.1.S1_at	CRP	taggaacctgaggggtgagg	ggggcctcaatcggaactaca
Ssc.2033.1.S1_at	CRY1	ttaggacaaaattttctgaaatc	aaatttgggtcaaaacaagca
Ssc.19692.1.S1_at	CXCL2	caggaggaggagttctctcaa	atgtccaagggaacgtcaa
Ssc.208.1.S1_at	CYP1A1	ctccatctacatgccact	agagcctgttccaaagaatg
Ssc.203.1.S1_s_at	CYP3A22 / CYP3A39	cgccctgaaagggtcagtaa	ggggcgctctgtgtttcttta
Ssc.203.1.S1_at	CYP3A22 / CYP3A39	cctgaaactaagctctcaagga	caaggtctctgatgctctgg
SscAffx.21.1.S1_at	CYP7A1	caattggcacaacgttttctt	gcgtgatgtgtccttcaaaa
Ssc.21145.1.S1_at	DEFB1	gctgactgtctgcctcctct	acttggccttgcactgttaa
Ssc.16236.1.S1_at	DPP4	acatgggtcttccaactcca	aatccactccagcatccac
Ssc.30467.1.A1_at	ELOVL2	tgatgagcagaccagcactt	ccgccccaatcatgatacta
Ssc.13778.1.S1_at	FCGR1A (IgG)	caagtgcagggtcaacaacg	cgttgctcttccactcaaca
Ssc.16377.1.A1_s_at	GSTA1	tgttgaaagagccatggacaa	ttcagcagagggaagtgg
Ssc.390.1.A1_at	HIF1A	tggggcatggtaaaagaaag	gtaatgctccccatccaa
Ssc.47.1.S1_at	IGFBP2	catggcctgtacaacctcaa	tggctgtggttactgcatc
Ssc.15588.1.S2_at	IGFBP3	agcccctgtaccgtaatcct	ggcagaacgtctgaagatcc
Ssc.286.1.S1_s_at	IRG6 (RSAD2)	caccatgcgggaattctact	tggaaaaatgcctgacaatga
Ssc.14340.1.S1_at	LITAF	ccggttggtatgtttggaaag	aaaccaactctggcttgtgg
Ssc.670.1.S1_at	LYZ	aaggagcagaatgccagatg	cttcatggtcagcaaaatgc
Ssc.548.1.S1_a_at	MMP7	tggccattcttaggtctgg	tgaatgtgtgttctccctgaa
Ssc.600.1.S1_s_at	MT1A	actgctctgccccacag	acagcagctgcactgtcc
Ssc.9372.1.A1_at	NFKBIZ	ggaaacatgctttgctggtt	ccaaatacatcttgggaca
Ssc.13870.1.A1_at	NR1D2	tcaagtatccttttctttcca	tgttcggatcaaatcaaga
Ssc.15928.1.A1_at	NR3C1	ccaggctcactcccatcact	gtctcagccagactccaagc
Ssc.6323.1.S1_at	PLIN2	tctgatggcctcctcagttc	tctcaggttctgtgggtctct
Ssc.8177.1.A2_at	PLP1	agaagcttccaggtgcata	taggaggtctccagggaagg
Ssc.14490.1.S1_at	PTGR1	catgaacacatcacggaagg	tggactctcgtgtcgtcttg
Ssc.2381.1.A1_at	S100A9	caaatgtcgcagatggaatg	caggctctccaggatgtggt
Ssc.7493.1.A1_at	SIRT1	cgcagagaagcatttattttcc	ctcttctccggcagtaatgg