

<i>Pparg1a</i>	19017	peroxisome proliferative activated receptor, gamma, coactivator 1 alpha (Pparg1a)	GGGACATGTGCAGCAAGA	AAGAGGCTGGCTCACAA	74	93
<i>Pparg1b</i>	170826	peroxisome proliferative activated receptor, gamma, coactivator 1 beta (Pparg1b)	GGGACATGTGCAGCAAGA	AAGAGGCTGGCTCACAA	74	94
<i>Prdm16</i>	70673	PR domain containing 16 (Prdm16)	CAGCAGGGTGAAGCATTCT	GGGTGATCCGCTTTGG	87	97
<i>Ptgs2</i>	19225	prostaglandin-endoperoxide synthase 2 (Ptgs2)	CAGCCAGGCAGCAATCCTT	AGTCCGGGTACAGTCACT	100	94
<i>Plk2b_gDNA</i>	19229	Protein tyrosine kinase 2 beta (Plk2b)	TGGTAATCAAGTCTGAGATGGTGT	CCITTCGCAATGAAGTGTGT	100	100
<i>Plk2b_RNA</i>	19229	Protein tyrosine kinase 2 beta (Plk2b)	AAGACCTGAAGAAGACTGACCC	CAAGATTCATCATATCATGCCTC	76	100
<i>Rb1</i>	19645	retinoblastoma 1 (Rb1)	CCCTTGATGGCTTCAGATT	ATGGTTACCTGGAGAGCA	119	93
<i>Rplp0</i>	11837	ribosomal protein, large, P0 (Rplp0)	CCTATAAAGGCACACGCGG	ACGATGCTCACTCAACGAGG	67	102
<i>Runx111</i>	12395	runt-related transcription factor 1; translocated to, 1 (cyclin D-related) (Runx111)	GGACTCGCTTTTGTGTCT	CCACAGATCAACTTTTCCGC	116	96
<i>Rxra</i>	20181	retinoid X receptor alpha (Rxra)	CAACCTGACTTAAGGGCT	ACTTGAGCCGATGGAACG	169	101
<i>Sirt1</i>	93759	sirtuin 1 (Sirt1)	GATGACAGAAGTCAACAGC	AACAAAAGTATGGACTATCCGC	72	95
<i>Sirt2</i>	64383	sirtuin 2 (Sirt2)	CGGACTCTGACCTCTG	CGAGCTGAATCTGAGCCTCC	65	94
<i>Sirt3</i>	64384	sirtuin 3 (Sirt3)	GCTACATGCACGGTGTCTGAA	CAATGTCGGGTTCAACGCC	115	90
<i>Slc25a4</i>	11739	solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4 (Slc25a4)	CCGTTTGACACTGTTCTGCG	ATAATATCAGCCCTTCCGGC	62	101
<i>Slc27a1</i>	26457	solute carrier family 27 (fatty acid transporter), member 1 (Slc27a1)	GGGATCTCTCTCCACCCC	GAGCTTGCTGATCAGCTT	106	99
<i>Slc29a1</i>	63959	solute carrier family 29 (nucleoside transporters), member 1 (Slc29a1)	CACCAGCCTCAGGACAGTAT	GTCAGCGGTTTGTGAA	120	97
<i>Slc2a4</i>	20528	solute carrier family 2 (facilitated glucose transporter), member 4 (Slc2a4)	CTTATTGACGGCCTGAGTC	GGGGTCCCATCTGTCAGA	86	95
<i>Slit2</i>	20563	slit homolog 2 (Drosophila) (Slit2)	AGTTTGTCTGCAGTGATGAGG	CTACAGGTACAAGCAGCGGG	91	96
<i>Sp100</i>	20684	nuclear antigen Sp100 (Sp100)	TGATGGAGGGAACCCAACTC	CTTCTTGAAGATAGTGGCAC	69	96
<i>Src</i>	20779	Rous sarcoma oncogene (Src)	GAAGTGGATGTGAGAGAGGG	GGATGAACCGTCTGCC	71	97
<i>Srebf1</i>	20787	sterol regulatory element binding transcription factor 1 (Srebf1)	CACAGCGGTTTTGAACGACA	CATAGGGGGCGTCAACAGG	89	97
<i>Taz</i>	66826	tafazzin (Taz)	AGCCCTTCAGTACCTCCT	CGCATCTCACAGTCACTT	67	99
<i>Tbp</i>	21374	TATA box binding protein (Tbp)	TTTGGCTAGTTTCTCGGT	TGAAATAGTATGCTGGGCACT	59	95
<i>Tbp_gDNA_Control</i>	21374	TATA box binding protein (Tbp)	TTCCAAAATCCGGGATGGC	AACGATTCCGCACAGTCTT	81	96
<i>Tbx1</i>	21380	T-box 1 (Tbx1)	GGCAGGCAGCAAGTTC	TTGTCATCTACGGGCACAAG	102	96
<i>Tcf7l2</i>	21416	transcription factor 7 like 2, T cell specific, HMG box (Tcf7l2)	TTTGGAAAGACGGCCAGA	CTTGGACATCAAGCAGTGGC	173	101
<i>Tfam</i>	21780	transcription factor A, mitochondrial (Tfam)	TCCTCTGCTATCAGCTTG	GGATAGCTACCATGCTGAA	61	100
<i>Tmem26</i>	327766	transmembrane protein 26 (Tmem26)	GCTCACCTCAAGTCAA	GGGTGCTGCAACTGTT	131	91
<i>Tnfrsf9</i>	21942	tumor necrosis factor receptor superfamily, member 9 (Tnfrsf9)	CGTGCAAGACTCTGTGATAAC	GTCCACTATGCTGGAGAGG	104	97
<i>Tsc22d3</i>	14605	TSC22 domain family, member 3 (Tsc22d3)	TTACTGAAATGGCAGCTTGT	CTCGGAGCTAGCTTTTGG	62	92
<i>Twist1</i>	22160	twist basic helix-loop-helix transcription factor 1 (Twist1)	CCATGTCGCGTCCCACTA	ACAATGACATCTAGTCTCCGC	70	98
<i>Ucp1</i>	22227	uncoupling protein 1 (mitochondrial, proton carrier) (Ucp1)	CACCTCCCGTGGACACT	CCTAGACACCTTTATACCTAATGG	91	96
<i>Vdr</i>	22337	vitamin D receptor (Vdr)	ACTTTGACCGGAATGTGCTT	CATGCTCCGCTGAAGAAC	113	100
<i>Wnt1</i>	22408	wingless-type MMTV integration site family, member 1 (Wnt1)	GATGGTGGGCATCGTGAAC	GGGTTCTGTGGATCAGTCG	130	97
<i>Wnt10b</i>	22410	wingless-type MMTV integration site family, member 10B (Wnt10b)	AGCCGCATCTCAAGCG	AGAAGGAGAAAGCACTCTCAG	46	101
<i>Wnt3a</i>	22416	wingless-type MMTV integration site family, member 3A (Wnt3a)	TGGCTCTCTCGGATACCTC	GCCAAGGACCACAGATCG	79	97
<i>Wnt5a</i>	22418	wingless-type MMTV integration site family, member 5A (Wnt5a)	TGCTTGGATTGCCCCCAA	AATGGGCTTCTCATGGCGAG	119	97
<i>Wnt5b</i>	22419	wingless-type MMTV integration site family, member 5B (Wnt5b)	GCTACCGCTTGGCAAGGATT	CATTTCAGGCGACATCGCA	154	100