

S3 Table qPCR Primer

(A) RT-qPCR Primer

Gene	Sequence
CCL3	Forward primer: ATGCAGGTCTCCACTGCTG Reverse primer: TTTCTGGACCCACTCCTCAC
DNase1L3	Forward primer: AGGACACCACGGTGAAGAAG Reverse primer: GTGAAGGCCCTTGAAGACTG
ZNF608	Forward primer: TTCAGGGCTCATGGGAAACT Reverse primer: GACACTCTGGCTCTGTCACT
GPR183	Forward primer: ACTACACAGAGACCCGAACG Reverse primer: CCACGAGCCCAATGATGAAG
GPR174	Forward primer: TGTACCCCTTTTCGCTTCCAT Reverse primer: ACAAACCCAATCAACTCGCC
RGS1	Forward primer: TGCAATGGTCTCAATCTCTGG Reverse primer: CAGGGCAAAAGATCAGACTCTG
KLRC3	Forward primer: TCTGGCCAGCATTTTACCTTC Reverse primer: CCACGTACATGTAGCATTGCA
CD69	Forward primer: AACACGTCATGAAGGGTCCT Reverse primer: GGCTGTCTGATGGCATTGAG
LLRN3	Forward primer: GAGGTGAAGCATTCCACAGA Reverse primer: CCCATGCTTCTTCAGTATTTGC
IKZF2	Forward primer: TGAAATGTGACGTCTGTGGC Reverse primer: TTGAACGGCTTCTCTCCAGA
SERPINI1	Forward primer: AGACGAAAGCAGGAACGAGA Reverse primer: GCCCAAGTTCCATCATTCC
PCDH9	Forward primer: TGGGAAAGGGGAAAGGTTGA Reverse primer: TCAGAGCAGCCAACAGGTAA

(B) ChIP-qPCR Primer

	Locus	Sequence
CBF1-independent loci	chr19:42,633,923-42,634,005	Forward primer: TGGCCTCAAGGGAGTGAAC Reverse primer: GGCCACCTCTCTTGTTGTG
	chr16:11,043,370-11,043,449	Forward primer: AGTGCACATTGGCAGAACTC Reverse primer: CTCTGAGCTTGGGTCCTCAC
	chr3:39,189,615-39,189,702	Forward primer: TAGGCAAGCGTGAGAGAAGC Reverse primer: CCCGGAAACATCTGTGAGTC
CBF1-dependent loci	chr4:40,187,701-40,187,810	Forward primer: AGGGAAAGTTAAGACAGGCCTT Reverse primer: TAGAGGATTCCAACCCAATGCC
	chr10:74,057,129-74,057,245	Forward primer: CTTGGCACGAACTCTTCCTTC Reverse primer: AGATGTGAGAATGATGTGGCCG
	chr11:9,623,169-9,623,267	Forward primer: GCTGGGTTTGCGGTTTAATG Reverse primer: TTGCTGCTCCCTAAGGTTTG
Control loci	(+) chr8:24224967-24225059	Forward primer: CTTCATGGCTACAGACTCTTGG Reverse primer: CCTATGTCTCGCTTCCTGCT
	(-) chr8:99055763-99055882	Forward primer: CTGGTCTGACGCTCCTGACT Reverse primer: CAGTGCCCAGAATTCCAGAT