

Table S1. Primers used for DNA methylation analysis and expression analysis.

Gene	Type	Forward primer sequence	Reverse primer sequence	Annealing temperature (°C)	Reference
HE6	M	CGGTGGGACGTCGGGGTCTGC	CTAAAATACGCACCGATAACCGCG	72	1)
SB5	M	GGGATCGATTAACTCGC	CAAACCTAACCGCCACGTCCG	72	1)
SA9	M	GATTCGAAGAAATACGAGCGGATC	AACGTTATCGATAAAACTCGCCTCG	72	1)
SD2	M	GCGGGGGTTTCGAGGTT	GCCATCACCTCCCACATCGTAAACG	72	1)
SD2	U	TGTGAGTGTGGGGTTTGAGGTT	ACACCATCACCTCCCACATCAAACA	62	1)
B2 repeat	N	AGATGGTGTGAGTTATTATGTG	CTATTCTCCAAAAATCTAAAT	51	1)
HE6	BS	GTATTTAGGGTTGAGGATAATAGA	AAACACTCAAACATCCCTTC	60	This study
SA9	BS	AGGATTGGTGTAGGATT	CTAACCCAACAAAAACCCCTTC	54	This study
Cd14	RT	TGTCCAACAAAGTTCCGGCACT	TCCAGITCCCTGCGGCTGTACCTT	69	1)
Cd3g	RT	CGAGCGTGTTCAGTGTCAAGG	GCGAAGAAGAAGCCGAATATGGT	54	1)
Ela2	RT	GCTACCATCACGCCAACGTG	GCTGGGTAGGGTCGGTTGTGC	69	1)
Il1b	RT	AACAGTGGCAGGATATAGTTGA	CCCATCTGGCAGAGGACTAAGG	66	1)
Nos2	RT	CGAGCCAAGGTCTACGTCAA	CGCACGCTCCGCAGACATAGAG	60	1)
Tnf	RT	GCATGGAGCTCAGGGACAACCGAG	GGCTGACGGTGTGCGTGAGGA	69	1)
Npx2	RT	CCGCATCACCTCCCACATCGT	AAAGCGGCCACAGAAAGCGTTG	62	1)

M, primers specific to methylated DNA; U, primers specific to unmethylated DNA; N, primers for normalization of qMSP; BS, primers for deep bisulfite sequencing; RT, primers for quantitative RT-PCR; and 1) Niwa et al., Cancer Res, 70:1430-1440, 2010.