

Supplementary Data

SUPPLEMENTARY TABLE S1. SEQUENCE OF THE PRIMERS USED IN THIS STUDY

<i>Sl no.</i>	<i>Gene name</i>	<i>Accession no.</i>	<i>Size (bp)</i>	<i>AT (°C)</i>	<i>Primer sequences</i>
1	<i>OCT-4</i>	EU926737	232	58	F, GATATACCCAGGCCGATGTG R, TCGATACTCGTCCGTTTCT
2	<i>CDX-2</i>	NM_001206299.1	121	58	F, GTCTGGAGCTGGAGAAGGAG R, CTGCGGTTCTGAAACCAAAT
3	<i>DNMT3a</i>	NM_001206502.1	188	58	F, GTGCTGTCTCTATTCGATGG R, CCATTCTGGATATGCTTCTG
4	<i>DNMT1</i>	NM_182651.2	138	58	F, CTCAGAAGGGAGATGTGGAG R, TAGTAGTCACAGTAGCTGAGGA
5	<i>NANOG</i>	DQ487022	211	57	F, GGGGAAGGGTAATGAGTCCAA R, AGCCTCCCTATCCCAGAAAA
6	<i>SOX-2</i>	DQ487021	162	60	F, TCCACATTGAGATCAGCAA R, CATGAGCGTCTTGGTTTTCC
7	<i>CASPASE3</i>	NM_001077840.1	158	58	F, TGGTATTGAGACAGACAGTGG R, AGCATCTCACAAGAAGCCTG
8	<i>BCL-XL</i>	AF245488.1	187	58	F, CTTGGATGGCCACTTACCTG R, AAGAGTGAGCCCAGCAGAAC
9	<i>GATA2</i>	XM_583307.3	140	58	F, GAGGACTGTAAGCGTAAAGG R, AAGAACCAAGTCTCCCAT
10	<i>GAPDH</i>	GU32429.1	235	58	F, ATCACCTCGCTTCTACTGG R, AGGAGACAACCTGGTCCTCA

AT, Annealing temperature; F, forward; R, reverse.