

**Table S7. Genes and primers used for bisulfite pyrosequencing.**

Gene	Primer	Sequence (5' to 3')	Chromosomal localization (bp) <sup>b</sup>	Number and position (relative to TSS) of analyzed CpGs
NANOG	forward	GGAATATGGTTAATAGGAATGGGATAA	chromosome 12	3
	reverse <sup>a</sup>	CCAAACTAATTCAAACCTCCTAACCTTC	(7,941,610 –	(-296, -300, -
	sequencing	TTTTAAAAATTAAGAAAAAGGT	7,941,761)	302)
OCT4	forward	GGGTTTGGAAGTTAGTTAGG	chromosome 6	5
	reverse <sup>a</sup>	CAAACCCTCATTTCACCAA	(31,138,275 –	(+92, +84, +77,
	sequencing	ATTTTATTATTGGAGGGG	31,138,445)	+62, +44)

<sup>a</sup>biotinylated primer

<sup>b</sup>according to Ensembl release 62 - April 2011 (based on: Homo sapiens high coverage assembly GRCh37 from the Genome Reference Consortium)

TSS, transcription start site