



Figure S1 (A) Alkaline phosphate assay on biPSC colonies, Cells were weakly positive. (B) Normal bovine karyotype displayed by the biPSC generated.

Table S1 List of primers

Primer	Sequence	Length	Tm	GC%	P.S	t°	Ref
pMxhOCT4	<i>CTAGTTAATTAAGAATCCCAGTG</i>	23	46.6	34.8	300.0	58	Sumer et al. 2011
	<i>CACTAGCCCCACTCCAACCT</i>	20	55.2	60.0			
pMxhSOX2	<i>CTAGTTAATTAAGGATCCCAGG</i>	22	53.1	40.9	300.0	58	Sumer et al. 2011
	<i>TGTTGTGCATCTTGGGGTTCT</i>	21	60.1	47.6			
pMx hKLF4	<i>ACAAAGAGTTCCTCAAGGTG</i>	24	60.8	45.8	250.0	58	Sumer et al. 2011
	<i>TCCAAGCTAGCTTGCCAAACCTACAGG</i>	27	66.8	51.9			
pMxhc-MYC	<i>CTAGTTAATTAAGGATCCCAGTG</i>	23	54.0	39.1	500.0	58	Sumer et al. 2011
	<i>CAGCAGCTCGAATTTCTTCC</i>	20	57.2	50.0			
pMX-hNanog	<i>CTAGTTAATTAAGAATCCCAGTG</i>	23	52.4	34.8	550.0	58	Sumer et al. 2011
	<i>GGGTAGGTAGGTGCTGAGGC</i>	20	61.7	65.0			
SOX2	<i>CATCCACAGCAAATGACAGC</i>	20	57.7	50.0	251.0	58	Sumer et al. 2011
	<i>TTTCTGCAAAGCTCCTACCG</i>	20	58.2	50.0			
POU5F1	<i>GTTCTCTTTGAAAGGTGTTTC</i>	21	49.3	42.9	313.0	58	Sumer et al. 2011
	<i>ACACTCGGACCACGCTTTTC</i>	20	54.2	55.0			
NANOG	<i>GTGTTTGGTGAAGCTCTCTG</i>	20	50.6	50.0	308	58	Sumer et al. 2011
	<i>GGGAATTGAAATACTTGACAG</i>	21	46.8	38.1			
KLF4	<i>GCCCCTAGAGGCCACTT</i>	18	54.7	66.7	455	58	Sumer et al. 2011
	<i>CACAACCATCCCAGTCACAG</i>	20	52.7	55.0			
c-MYC	<i>CGCGGTGCTCCTTCTCGCCAGG</i>	25	68.7	76.0	412	58	Sumer et al. 2011
	<i>GTCCGGGGAAGCGCAGGGC</i>	19	62.6	79.0			
GAPDH	<i>ACCCAGAAGACTGTGGATGG</i>	20	59.0	55.0	247	60	Gad et al, 2012
	<i>ACGCCTGCTTCACCACCTT</i>	19	62.1	57.9			
POU5F1	<i>GTTTTGAGGCTTTCAGCTC</i>	20	58.22	50	182	55	Gad et al, 2012
	<i>CTCCAGGTTGCCTCTCACTC</i>	20	59.75	60			
Oct4 3'UTR	<i>GGGTTTGTACTAGGGCTTTGGG</i>	22	60.88	54.55	124	55	Han et al, 2011
	<i>GCATCATTGAACCTCACCTCCCT</i>	24	61.35	45.83			
Sox2-3'UTR	<i>GCACGGCCATTAACGGCACAC</i>	21	65.25	61.9	126	60	Han et al, 2011
	<i>CTCCATGCTGTTTCTTGCTGCTCCTC</i>	25	64.04	52			
Nanog 3'UTR	<i>TCTGTGTCAGTTTGAGGGACAGG</i>	23	62.07	52.17	234	60	Han et al, 2011
	<i>AACAAGTAAAGCCTCCCTATCCCA</i>	24	61.32	45.83			
Rex1	<i>CACTGCTTCGATTACAACCCAG</i>	25	63.31	52	231	60	Han et al, 2011
	<i>CCACGTACTTACTGCTGGAGATGGG</i>	25	64.62	56			