

Supplementary Table S1. Primer sequences used for qPCR analysis of specific genes

Gene name	Primer sequence (5' to 3')	Annealing temperature (°C)	Amplicon size (bp)
<i>SOX2</i>	F: TGGTCTGTCTTCATGGTGTCT R: GTGGGAGGTTAACCATGGGA	59	126
<i>OCT4</i>	F: CGGAAGAGAAAGCGGACAAG R: CGATGTGGCTGATCTGCTG	58–59	103
<i>c-Kit</i>	F: CTTGGACGGCTTGCTAACG R: AAATCCACCGTGGGTATGGG	59	145
<i>PPAR<math>\gamma</math></i>	F: TACCAAAGTGCCATCAAA R: TGGAGTGGAAATGCTGGA	56	111
<i>ACTB</i>	F: GTCATGGTGGGTATGGGTCA R: CGTTGTAGAAGGTGTGGTGC	59	151

Supplementary Table S2. Average diameter and number of spermatogonial stem cell (SSC) colonies on different days of culture

Sl. No	Days of culture	Average colony diameter ( $\mu\text{m}$ )	SE	Mean <i>n</i> of SSC colonies per 1 161 px surface area	SE
1	day 9	68.92 <sup>a</sup>	1.20	99.17 <sup>a</sup>	4.10
2	day 30	139.71 <sup>b</sup>	1.67	69.67 <sup>b</sup>	2.38
3	day 65	213.53 <sup>c</sup>	12.52	30.50 <sup>c</sup>	1.13

SE = standard error

<sup>a–c</sup>means in a column with different superscripts differ significantly ( $P < 0.01$ )