

Supplementary Table 4. Primer sequences and PCR conditions for RT-PCR analysis

Gene	Accession Number	RT-PCR					
		Formard primer (5'-3')	Reverse primer (5'-3')	Size (bp)	Annealing (C)	cycles	exons analyzed
ANKRD30A	NM_052997	CAAAGAGGAAGAAGACCATCAACC	TGATGGCATTGTAGAGCCTTCAT	181	55	35	ex1-3
INSL6	NM_007179	AAAAACTCTGCGGCCATGC	CCCAACTGTTTACTGCTTCTTCC	201	55	35	ex1-2
FLJ40201	NM_152607	TGCAGAGCCACAGGTTTGAC	CCTGTGTCCACCTCCTTCATC	181	55	35	ex2-3
SOHLH2	NM_017826.1	TATCTGCCAGGAGCACTGCC	GAATATGCAATCATCCAAAAGCG	187	55	35	ex1-2
DPPA5	NM_001025290	GGAACTCTCCCGGCACGTA	CGTAAACCACGACCTCGGTG	211	55	35	ex1-2
GAPDH	AF261085	TCCCATCACCATCTTCCAG	ATGAGTCCTTCCACGATACC	309	50	25	NA