

**Supplementary Table 1: RT-PCR primers for Wnt signaling components in the mouse**

Gene	Primer pair (5'-3')	Product (bp)	Span intron or not	Annealing temperature (°C)
<i>Wnt1</i>	F: CTTGGCAAGATCGTCAAC R: CTGCCTGTTGTGAAG	281	√	55
<i>Wnt2</i>	F: TCCGAAGTAGTCGGGAATC R: CCTGAAGTCAGCCATGGC	380	√	53
<i>Wnt2b</i>	F: GAGAAGAGGCTTAAGGATGC R: ACCTGCAGCCTTGCCAAG	330	√	55
<i>Wnt3</i>	F: AGCACACAATGAAGCAGGC R: TGAAGAGCGCTACTTAGCC	237	√	58
<i>Wnt3a</i>	F: TGTTCGGACAAAGCCACC R: TTCATGGCAGAGCGGGCA	252	√	58
<i>Wnt8a</i>	F: CAAGGCATCTGACCTAC R: GTCATACTGGCCTTTAGG	541	√	53
<i>Wnt8b</i>	F: CACCCTGACTAGAACTGC R: CAGCAGGTCCACCTTGAG	370	√	55
<i>Wnt9a</i>	F: CTGCTCAAGCGAGGCTTC R: ATCACCTTACACCCACG	281	√	55
<i>Wnt9b</i>	F: GCATCAAGGCTGTGAAGAG R: TGGCACTCCACGTAGCAG	439	√	56
<i>Wnt10a</i>	F: GCTCCTGTTCTTCTACTG R: TGGATGGCGATCTGGATG	210	√	52
<i>Wnt10b</i>	F: AAGCAATGAGATTCTGGGCC R: ATCCCGAGAGAACTTCTCTC	556	√	55
<i>Fzd1</i>	F: GCTGCTGCTGCTGCTTTG R: GCAGTTGGGCATGATGG	251	–*	55
<i>Fzd2</i>	F: GTAGTGTGCTTGACATCCC R: ACAAGAGCCTGCCAGAG	621	–*	56
<i>Fzd3</i>	F: GGAATATGGACGTGTACAC R: ACCATCATGTAGCAGACTGC	474	√	52
<i>Fzd4</i>	F: GCTACAACGTGACCAAGATG R: GAATTGCTTCCCACGGAGT	394	√	52
<i>Fzd5</i>	F: GAAGGAAGAGAAGGCAGTG R: CACTCAGTCCACACCAGATAG	272	√	58
<i>Fzd6</i>	F: CATGATGGCCGGAACCAAG R: TTGCTTCCAACCCAGAAGAC	287	√	53
<i>Fzd7</i>	F: CCCACCTTCACTGCGATG R: TGCCACCATGAAGTAGCAAC	406	–*	60
<i>Fzd8</i>	F: GCAGGACATGAGAAAGTGG R: ACAAGGAGAAGACCACCG	316	–*	56

**Supplementary Table 1: Contd...**

Gene	Primer pair (5'-3')	Product (bp)	Span intron or not	Annealing temperature (°C)
<i>Fzd9</i>	F: GAACGCCTCAACATGGACT R: TTAGTCATGTGCAAGACCAC	335	–*	52
<i>Fzd10</i>	F: AATGCTCTGACCGGCTTCG R: GCTGCATACGTGTTGCCAG	494	–*	56
<i>Gsk3α</i>	F: GGACAAAGGTGTTCAAATC R: TCAGTTAAAGCTTGTGAGG	298	√	53
<i>Gsk3β</i>	F: CTTTGGAAAGTGCAAAGCAG R: CCAACTGATCCACACCAC	200	√	53
<i>Dkk1</i>	F: ACACCAAAGGACAAGAAGG R: GGCTTGATGGTGATTTTTC	224	√	53
<i>Dkk2</i>	F: TGAACCAAGGACTGGCTTTCG R: GCAGCGGGTACCAGGGCAAC	565	√	56
<i>Dkk3</i>	F: GTACACCTGCCAGCCATG R: CCTCTGGTTGTCACAGATG	142	√	56
<i>Dkk4</i>	F: AAAGCAGTAAGGGACAGGAG R: CCTCTGGAGCAGACTTGT	134	√	56
<i>Dkk11</i>	F: GAGGAGAACCAGGAGCAC R: ATGATCCAGAAGGCCACC	254	√	53
<i>Axin1</i>	F: CATGCAGTGGATCATTGAGG R: AGCAGCTCCTTGAAGTGGC	546	√	56
<i>Axin2</i>	F: CAGCTGAAAACGGATTCAGG R: CAGTTTCAGTTTCTCCAGCC	353	√	56
<i>Dvl1</i>	F: GAGAACATGAGCAATGACG R: ACTGGAGCCACTGTTGAG	615	√	55
<i>Apc</i>	F: CTTGCTGATCTTGACAAAG R: TCAAAACACTGGCTGTTTC	445	√	56
<i>Kremen1</i>	F: CTGATCGCATCAATCAGGC R: TGAAGTGTGAGGATGAGGAG	249	√	53
<i>Kremen2</i>	F: GGCTTCGCGCTCACCTAC R: TACCACACAGCCAGCTTC	308	√	53
<i>Ck1α1</i>	F: TAGCTGACCAGATGATCAG R: GTATCGGGCAGTGCCAGTG	299	√	53
<i>Ctnnb1</i>	F: ATCCAAAGAGTAGCTGCAGG R: TCATCCTGGCGATATCCAAG	311	√	53
<i>Lef1</i>	F: GCATCCCTCATCCAGCTATT R: GCTTCTCTCTTCTCTTCTTC	378	√	56
<i>Tcf4</i>	F: CACTTTCCTAGCTCCTTCTTC R: GTTCGTGTGGTCAGGAGAATAG	388	√	58
<i>Gapdh</i>	F: GAACGGATTTGGCCGTATTG R: TTTGGCTCCACCCTTCAAG	328	√	56

Contd....

\*Represents gene with only one exon. RT-PCR: reverse transcription polymerase chain reaction