

Supplemental Table S1 (Boije et al, Sonic hedgehog signalling patterns the developing chicken comb as revealed by exploration of the Pea-comb mutant)

Supplemental Table S1: List of qRT-PCR Primers

Genes	Primer sequence (5'-3') F- forward, R-reverse	Ensemble ID
<i>ARHGEF12</i>	F-CGACTTCGATCCGACAGATAGC R-CCGTTGGATAACAACACACCG	ENSGALG00000006663
<i>BMP4</i>	F-TTTACGAAGTGATGAAGCCGCT R-CTCACATCAAAGGTCTCCCAGC	ENSGALG00000012429
<i>cKIT</i>	F-TCATCGGTGCCATCCTTTGA R-CTTGTCACCATTTGCTGGATGC	ENSGALG00000013925
<i>COL1A2</i>	F-GCTTTGTGGATACGCGGATTT R-TGTGGCCCTTTGTCTCCTCTAG	ENSGALG00000009641
<i>COL2A1</i>	F-TTCACCTACAGCGTCTTGGAGG R-GATATCTACAATGGACAGGCGCG	ENSGALG00000013587
<i>DCT</i>	F-TGAGGAAGCCACCAGTTGTCA R-GGTGGATGGTAGTCTTTGCAC	ENSGALG00000016899
<i>EFEMP1</i>	F-TGCCCCGTATCAAACACCAT R-TGGTTGCCTGAATCTGGAAGAT	ENSGALG00000007908
<i>ETS1</i>	F-GCAGAAAGAAGAGGCAAAACCA R-GCGTGCTCGATGCCATAACTA	ENSGALG00000001143
<i>FEZF2</i>	F-GTGCACCATTTGCAACAAAGC R-TCCTTTGCCGCAAGTGACA	ENSGALG00000007184
<i>FOXD3</i>	F-ACAACCTCTCGCTCAACGACTG R-CGTTGTCTGAACATGTCTCAGA	ENSGALG00000012644
<i>FZD1</i>	F-CCATGAGCCCTGACTTCACTGT R-AGTTGAGCGTTTTCCCCGA	ENSGALG00000009064
<i>GLI1</i>	F-CCTCTGACAGCCAAATACCCAG R-TTTCCCATCCTCCTTCTCCAG	ENSGALG00000014639
<i>GLI2</i>	F-CTGCAGACGATGATCAGGACCT R-GGCAGATAAATGGCCGTAGGA	ENSGALG00000011630
<i>HAS2</i>	F-TGCGTACAGTGCATCAGTGGA R-AGCCCAGACTTAGCACTCGGTT	ENSGALG00000016394
<i>HAS3</i>	F-CAGTTCATCCACACGGAGAAGC R-GGTGCTCCAGGAAAGCAAAGA	ENSGALG00000000630
<i>IHH</i>	F-GCAACAAGTACGGCATGCTTG R-TACCTGACTTGACGGAGCAGTG	ENSGALG00000011347
<i>ITGB3</i>	F-CAACCTTCGCATCGGCTTT R-GGCAAGCACTTTTCCCCAAT	ENSGALG00000000379
<i>MEOX2</i>	F-GAGGAAAAGCGACAGCTCAGAT R-TCTCTGATTTGCTCCTTGGTGA	ENSGALG00000010794
<i>MITF</i>	F-ACATGGACTGTCCCTTGTTCCA R-GGTCTTGGTTGCAGTTGTCCAG	ENSGALG00000007679

<i>MMP1</i>	F-AGTTCCAGTTTGGATCCTCGTGC R-TGCAGTGTGCATGTGGAAAAG	ENSGALG00000019061
<i>MMP13</i>	F-TGCTGGTCTTCAAGGACAGGTT R-TTTGTTTGGAAGCTCCGGC	ENSGALG00000017183
<i>MMP2</i>	F-TAGATGATGCCTTTGCCCGAG R-TGTTCCCATCGGCCAAAATT	ENSGALG00000003580
<i>PAX3</i>	F-CCCAGTCTGATGAAGGCTCTGA R-TTTCCAGCTCTTCCAGTTGCTC	ENSGALG00000005246
<i>PAX7</i>	F-CACATCCGCCATAAGATCGTG R-GGTACCTGCAGAGGATTTTGGA	ENSGALG00000003782
<i>PLAU</i>	F-AGGAATGGCAGAAGATCCAGG R-CCAGTCCAAGTTGCAGAGCAT	ENSGALG00000005086
<i>PTCH1</i>	F-TGCAGAGCCACGATCAGAAAT R-AGCAAATCCCTTGTTGAGCTTG	ENSGALG00000012620
<i>RHOA</i>	F-AGCGTGGCTTCAGTGCTATCA R-GCTGAATGGAACGTGGTCATTT	ENSGALG00000003806
<i>RUNX2</i>	F-GCTAAACCCAAACTTGCCCAA R-GGACTCATCCATCCTGCCACTA	ENSGALG00000016706
<i>S100B</i>	F-TGAAGAAATCGGAGCTGAAGGA R-TCCAGTGCCTCCATGACTTTG	ENSGALG00000006217
<i>SHH</i>	F-CCCCAAATTACAACCCTGACAT R-TTCATCACCGAGATCGCCA	ENSGALG00000006379
<i>SMO</i>	F-CATCGCTACGCAGACCAATAAA R-ATGCCAGTGCCAAACATGG	ENSGALG00000007910
<i>SNAIL</i>	F-CTGCACCCATCTGGACATCTTT R-GCGAATGCCAACACTGTGATC	ENSGALG00000008018
<i>SNAIL2</i>	F-TCTCCAGACCCTGGCTACTTCA R-CCCTCAGATTGGATCTGTCTGC	ENSGALG00000015241
<i>SOX5</i>	F-AGGAACAGATTGCAAGACAACA R-CTGCTGGATCTGTTGCTGAAG	ENSGALG00000013204
<i>SOX6</i>	F-ACTGTGGCTGAAGCTCGAGTCT R-TCCTTAGCCCAAACCATGAAAG	ENSGALG00000006074
<i>SOX9</i>	F-TCCCCACATCGATTTCGA R-GCAGGTATTGGTTCGAACTCGTT	ENSGALG00000004386
<i>SOX10</i>	F-CCCCCTGTCCCATCTGTTTAA R-AAGGCTAAGGCTGACAGTGCAG	ENSGALG00000012290
<i>SPAG6</i>	F-AGATGATCCTGAACCCCGATG R-GCTTCAACCACCAACTCTGCA	ENSGALG00000007892
<i>TWIST</i>	F-TGAGCCACTGAGGAGAGGAAGT R-GCCAGTTATCCCCCCTAAAAAA	ENSGALG00000010862
<i>VCAN</i>	F-CAATATGAGAACTGGCGACCAA R-TCTTGCAGGTGTAGGTCAGGTG	ENSGALG00000015624