

Table S1. Primer sequences for quantitative PCR.

Primer	Forward sequence (5' to 3')	Reverse sequence (5' to 3')
-2700 Areg TSS	ACATGTAAATGCTCCCTTCCCAG	TCTGTTTGAAGTAAAAGTTGGCTGT
-2100 Areg TSS	AATGAAACCTGTGAGACTCGGA	GGCAATCTGCTTACAGGGCT
-1600 Areg TSS	CAAACATGGTGCTGCTCACT	GTCCTGAACTGTGCCCTCAC
-520 Areg TSS	GAGCACACTTATCCCCAGCA	GCCAGACAGTGGCACTTAGG
-330 Areg TSS	GGAGTCCTTTAGCGCTTCGT	TGGCTTAGTGATGTTAGCCCG
-150 Areg TSS	ACAACGCGAGTGGGAAAGTT	TGGGGATCGAGCCCTGATTA
-50 Areg TSS	AAGTTTCTCCCCGCGTAATCA	GCGGGAGGTTACTACTCCAAA
+40 Areg TSS	TTCCTTCGCTCCCTTCTGAC	TCAGGCTCAGCTGGAGATGT
+240 Areg TSS	GCCGGTGGAACCAATGAGAA	CACCTGAGCCTAAGACCAGC
+370 Areg TSS	CCAAACGTCAGGGAGCAGAT	TCACCCTCCTAGCATGAGCA
Areg	CGAAATGCCTTCTGGCAGTG	GGCTTAATCACCTGTTCAACTCTG
Birc5	TCATCCACTGCCCTACCGAG	CAGGGGAGTGCTTTCTATGCT
Birc7	TCTCCATAGTCTTCGTGCCC	AGGAATGTGCGTACACAGCTA
Btc	GCATCCATGGGAGATGCCGCTT	ACCACTATCAAGCAGACCACCAGG
Ccn1	TTGACCAGACTGGCGCTCT	AGTTTTGCTGCAGTCCTCGT
Ccn2	GGAAAACATTAAGAAGGGCAAAA	CCGCAGAACTTAGCCCTGTA
Ccn3	AGAGTTGTTCTGAGATGAGACCC	CCCTCTGGAACCATGCAAATG
Ccn5	CTCAGCCCAAGGACACCAA	GGATACTCGGGTGGCTATGC
Ccn6	CGTGTGCGCATATCTTGTTG	GTGGGGCTGAAATACTTGGC
Egfr (ErbB1)	TCTGGAAACCGAAATTTGTGCTA	ACGGCCTTGCAGTCTTTCTC
Ereg	CAAGCAGTAGCCGTCCATGT	ACTGTACCGCCTTAGTTCAGA
Lats1	AGCAGCACGTAGAGAACGTC	TCTCATTGATCCTGGGCATCT
Lats2	TGCACTGGATTGAGGTGGACTCA	GAGAATGTGCCAGGCACCTCT
Lhcgr	GCTGGAGTCCATTGAGACGCTCA	AGCATCTGGTTCTGGAGTACATTG
Pgr	TCCAGGTGACCCATGAGGAA	TTGCCTTGATCAATTCGCGG
Ptgs2	CCTGAAGCCGTACACATCATTTGA	AGGCACTTGCATTGATGGTGGCT
Taz	ACTGGCCAGAGATACTTCCTTAATC	AGGCTGATTCATCACCTTCCTG
Tead1	CCTGGCTATCTATCCGCCGT	CCCGTTCTGAGTTTGATGTATCT
Tead2	ACCATCCTCCAGGTTGTGAC	CCTCGTTCACTGGTGGAGAC
Tead3	GCATTAAGGCTATGAACCTGGAC	TTTGGGCAGACGACATGGAT
Tead4	GGTGTATGGAGCCCCGAAAT	CGATCAGCTCATTCCGACCATA
Tnfrsf6	TGAAGGTGGTCTGCTCGCAACC	TCCACAGTTGGGCCAGGTTTCA
Yap1	GACGCTGATGAATTCTGCCTCA	CATGGCAAACGAGGGTCC