

S1 Table. Gene accession numbers and primers used in this study

Primer	Vector Base ID	Forward sequence (5' → 3')	Reverse sequence (5' → 3')	Annealing temp. (°C)	Size (bps)
Notch RNAi	AAEL001210	TAATACGACTCACTATAGGGACGTG GTTCCCAGAAGGATT	TAATACGACTCACTATAGGGCCCAG TCTTATCCATGGTTCG	60.5	528
Notch qPCR	AAEL001210	ATCGGGATGCTCAAGATGAC	TGATCATCCGAAATCTGCAC	51	466
Delta RNAi	AAEL011396	TAATACGACTCACTATAGGGGGATA TAGCGGTACTCGATG	TAATACTACTCACTATAGGGGTCAT CTTTCTCCTGCTCAC	60.5	591
Delta qPCR	AAEL011396	ACTACCTCACGATCACCAGA	GGATAGGGTTCACACTCGT	51	366
CSL RNAi	AAEL006419	TAATACGACTCACTATAGGGACGA GAGCGAGTCAGAGGAG	TAATACGACTCACTATAGGGCGTGA AAGTTAGCCCCGTAG	60.5	647
CSL CHK	AAEL006419	CCACCCA ACTATGTGCCTTT	ACATCCGTTCCGGTGTCTTTC	50	606
SPM779 qPCR	AAEL008779	TCCTTGTGGACCCTGTGTAA	AATCTGTGGCTTCAGTTGG	53	499
KLC2.2 qPCR	AAEL021010	CCCGAAGCCATTAAACAC	TGATTCACTGTCACCATTCTGTAC	55	465
S7 qPCR	AAEL009496	GAAAAGATGGCATGGGAAGTG	CTCCTGCTTCCGTTTGATTACC	52	69
Jun qPCR	AAEL003505	CTCCATGCGCGAAAAAGATA	TGTGTAGTTCGTTGTTCGT	54	261
P38 qPCR	AAEL008379	CCCAGTCCTTAAAGGCACG	CGCTTAAGGCCCTTAGAGATT	56	267
EGFR qPCR	AAEL004391	ACCCTTTGGATATAACCCGCG	GCCTATTGGCCCGTGTGGGAAT	56	289