

Table S2. RT PCR primers for Notch signaling targets and modifiers.

Gene	Exons Detected	Forward	Reverse
<i>Hey1</i>	All	CCCAGACTACAGCTCCTCAGA	ATGCTCAGATAACGGGCAAC
<i>Hey2</i>	All	ATGAAGCGCCCTTGTGAG	AGCCACTTCTGTCAAGCACTC
<i>Hes5</i>	All	ATGCTCAGTCCCAAGGAGAA	AGCAGCTTCATCTGCGTGT
<i>Hes1</i>	All	AAACGAAAATGCCAGCTGAT	GTCACCTCGTTCATGCACTC
<i>HeyL</i>	All	TGTGGGTCAAGAGAACGATCTTA	CCCCAGGTATCTTATGACCTCAG
<i>Lfng</i>	1-6	CTGCTGTTTCGAGACCTGGAT	CAATGTAGCCAATGGTGCAG
	4-8	AGCTATCCCCACACCCAAG	TACAGGTGGCAATGGACAGA
<i>Mfng</i>	All	GTCTCCAGGATCAGGCAACA	CTGGGTAGAGGAGACAATGGA
	processed transcript	GCTTTCCTGTCGTCTTGCTC	GCTCTGCAGAACAGTTGGTG
	4-7 (retained intron)	TTCCTGACAGCCCAGATGA	GCACCATCTTCAAAGCCAGT
	5-7 (retained intron)	GAACGCCTCCAGGAGAGACT	GGTGGCTAGGAGTTTGGTCA
<i>Neur1a</i>	1-3 (201)	CAGATCACAAAGGAACACTATCCA	CACAGGCTCACCGACAGG
	2-5	GACGTCTACGGCCTCACG	AGGTGAGTTGGGGGCTGT
	1-3 (202)	ACTTCTCCAGTGTCTCCTCTCTG	GCACAGGCTCACCGACAG
	1-2 (201)	GAAGGAGGAGGGGAAGTGG	TTCAGCCTGACTTGCTCGTA
	2-3	GCTCTTCCATCCCCACACTA	GACCCCACTGAAGAAAAGCA
	3-4	AGCATTGCCTGAGGAGTTTG	GGCTGTGCTGAGAATTGAGC
	4-5	GAGCTGCACCTGAGTCACAA	GACTGTGTCCACTGCGTGTT
	1-2 (202)	GCACACACTTGACACCTTT	GAGGATCTGGGAGCCCTTAG
<i>MamlD1</i>	3-6	TGATTCTATGCCTGCTCTGC	ATCTTTTGCTGGTTGGATGG
<i>Maml2</i>	All	TTATCAAGCCAGCCTTTGCT	TGAGGCTGGAGTTTGGAGTT
<i>Maml3</i>	2-5	AACGCATGCTTCTCATTAAAGC	AGCCACCTGGTTTCTCACAG
<i>Maml1</i>	2-5	TCAGAACCCCGTGAATAACC	GAGGACAGCTGGAGTTGGAC
<i>Numb</i>	2-8	GGCGGAGGTAGAGTCTGAGG	GGCCTTCACTGCTTTCTTTC
	7-11	CAGTTGCAAGATGCCAAGAA	GAGAGGCAGCACCCAGAAGAC
<i>Numbl</i>	1-6	CACAGCAGCAGCAGCACT	CAGGAGCACAGAAGGAGACC
	Retained intron	CACAGCAGCAGCAGCACT	TGGAATCGGCTCAGTTAAGG
<i>Rbpj</i>	1 (202)-4	ACTCGAGGCGTCTGCTTAAC	CCAACCACTGCCCATAAGAT
	1 (201)-4	GGACTACTCGGAGGGCTTGT	CCAACCACTGCCCATAAGAT
	1 (203)-4	TGGAGTTTTGGCGAGAGTTT	CCAACCACTGCCCATAAGAT
	3-7	AACGAGGGGATCAAACAGTG	CTGCCCGTAATGGATGTAGC
	6-11	GAATTTCCACGCCAGTTCAC	GCAGAAATGTCTGGGACCAC