Table S3 Editing site sequences for primers

Primers for amplification of short amplicons that include editing sites within a few bases of one of the primers. For some transcripts, editing sites were spaced further than could reliably be captured in a single short read: For those transcripts, multiple primer pairs were used to capture all editing sites. *Italics* indicates sequence/primer closest to editing site(s).

Gene	Forward primer	Reverse primer
Blcap (1)	*CTGCCCGTCCTCCTCATC*	AATCGGAGCAGTGGTACAGG
Blcap (2)	*GCCCGGCAGAGATCATGT*	CAGGAAGACCAGGGCACATA
Cadps	*GGATGTCCTTCGTGATAAGGTC*	CCGATTTCGAATGGTCTCAT
Cyfip2	AGCTGGATGCCAAGAAGAGA	*CTCTTCATAGTGAGCACTGGTCT*
Flna	*CGCCGCCTTACTGTTTCTAGT*	GGATGAAACGCACAGCATAC
Flnb	*GCTGCCTCACTGTTCTGAGC*	CATTCTCATGGGGGATGAAG
Gabra3	GGCTACTTTGTCATCCAGACC	*GAAAATACAAAGGCATAACAGACG*
Gria2 (1)	*TTTCCTTGGGTGCCTTTATG*	CTTTCGATGGGAGACACCAT
Gria2 (2)	*CATCGCCACACCTAAAGGAT*	CAATCAAAGCCACCAGCATT
Gria3	*GCAACCCCTAAAGGCTCAG*	TATAGAACACGCCTGCCACA
Grik1	*TGGAGTTGGAGCTCTCATGC*	ATGGGGGATTCCATTCTTTC
Grik2 (1)	*TCTCCCCTGATATCTGGATGTA*	AGAGCTCCAACTCCAAACCA
Grik2 (2)	ATGGAATGGAATGGTTCGTG	*GCACACAACTGACACCCAAG*
Grik2 (3)	*TGGAGTTGGAGCTCTCATGC*	ATGCGTTCCACAGTCAGAAA
Htr2c (1)	*ATCGCTGGACCGGTATGTAG*	TCACGAACACTTTGCTTTCG
Htr2c (2)	AGATATTTGTGCCCCGTCTG	*GAATTGAACCGGCTATGCTC*
Kcna1	*CATCGCTGGTGTGCTGAC*	CCTGCCTGTAGTGGGCTATG
Kcnma1	CAGCAGTAGCAGCAACATGG	*AAGAAGAGGACGCGTCTAGG*
Neil1	TCTAGAGGCCCTGCAACAGT	*TTCCTCTCCACGCTCTGG*