

Branch seq Splice acceptor seq.	gene exon seq. exons translation	CDS location exon # CDS location	Gene <intron length> Splice acceptor seq. Intron phase
FLJ32785 -471 .GCACCGTCCTTCCCG	fusion intron 1 -470 -455 GTTAATAAAATGCCCT	fusion intron 2 -454	fusion intron 2 -454 gtacgcttcacg.
ADAM15	1 exon 1 79 ATGCGGCTGGCGCT.. GCTCCCAAATATAG M R L A L L P N I G	80 <1205> gtgagtcctccgcct. phase 1	
.ccctcag... 1284 .tcctgggacctgcag	80 exon 2 186 GTGGCACTGAGGAG.. AAAAGGTGCTTCAG G T E E K V L Q	1392 <637> gtgagctctcactcc. phase 0	
.aggtgac... 2028 .ctctgggtgccacag	187 exon 3 263 ACCAGTCTGCCTGA.. CTGCTACAGAATAG T S L P E L L Q N R	2106 <416> gtaatagtgatggtg. phase 2	
.ttagcac... 2521 .ttcttctactccacag	264 exon 4 342 GGAGTTGGTCCCAG.. AGGGACACACTTTG E L V P G G H T L	2601 <161> gtgagtcaggccctc. phase 0	
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.ctctgac... 2925 .tttcttttttcttag	420 exon 6 612 AGGCTTGGTGGTCC.. GCCACATTCGCCGG G L V V L H I R R	3119 <1283> gtgagatgaatggc. phase 0	
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.ccccag... 5799 .tgccaccctccccag	1149 exon 12 1323 CTTCTACCAGGCC.. GTGGCTTCTGGAT F L P G L G F L D	5975 <274> gtgagccccctttccc. phase 0	
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	21b 2566 TTAATGGTGACAG -	10730 <127> gtttgtttgcagaca. phase -	
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.ctctgac... 11071 .cttcccgaatccag	2550 exon 23 2865 ACCAGCGCCACCGC.. TTGCAATAAACGTGACATCTTGGGAGCGTT P A P P P -		