Supplemental Table 1. Primers used in the present study.

	Forward	Reverse
Sequence analysis		
exon 3a	TGTGTGTGTGTGTGTCACC	GGCTTCGTCCCCAGTATCTT
exon 3b	AGAGGCCTTGCCTTGAAGAT	CTCTTGGACTGTGCCATTGA
exon 3c	TTCCAGCAAGGAGTTTGCTT	GCAAGAATGGAGCTCTGAGG
exon 3d	CCTTGTCGGGTAGCACTCTC	CAGGTAGTGGAGCAGGGAAG
exon 3e	AGCCTCACTCCCACCAGTAA	TAGCCTGGATTTACCCTCCA
exon 4	CGGTGTGGAGATAGGCAAAG	ATGATTTGCATGACCCCAAT
exon 5	AGAACCATTTTGTGGCCAAG	CTGTGTTCCTCCCAACAACC
exon 6	GTGGAATTTTGCTCGAGAGG	TGAATTCTCACCCCACTTCC
RT-PCR		
exons 3 and 6	TCAGTGGCCTCAGATTCCAT	TCAGGGGTCATTGCCATAAG
△exon 4	CTCTGCCCAGACAGACTCAT	TCAGGGGTCATTGCCATAAG
△exon 5	TCAGTGGCCTCAGATTCCAT	TCTGTTCCTCTTGAATGTTAGACG
GAPDH	GCACCGTCAAGGCTGAGAAC	TGGTGAAGACGCCAGTGGA
cDNA cloning		
	GTCGACGGCTGATGGGGGCTACCCTAAT	CGGTACCGGGCATCCTGCTCAGATTTA
Mutagenesis		
g.IVS4−2A>G	CTAACATTACAAAGCCTGCAAGCTTG	GGCTTTGTAATGTTAGACGGTGACAC
△exon 5	TAACATTCAAGAGGAACAGAGATCA	TCCTCTTGAATGTTAGACGGTGACAC