



RT followed by 35 cycles PCR suggests that both Promoters are used and all 2/3 A-G exons are skipped

Apbb1							
P1:1/2B		P2:1/2B		2/3 AF			
				A	59		
				В	171		
				C	80		
				D	29		
				E	84		
				F	40		
W/ 1/2 B (115 nt)	269	W/ 1/2 B	146	G	13		
Skip	154	Skip	131	Skip	184		

D. cDNAs sequenced

Alt P1	Exon 2 801 nt	Exon 3
Alt P2	Exon 2 801 nt	Exon 3