

Table S3

TableS3. Aberrantly spliced cellular transcripts as detected by 5'RACE.

target genes:	no. of reads	putative target genes:	no. of reads
EEF1G	7	SIRPB1	3
STAT5B	5	ZNF251	1
FAM126b	3		
COBRA1	2		
CHKA	2		
POLA2	1		
PRPL22	1		
ZNF426	1		
CLCN7	1		
FBX038	1		
MECP2	1		
COX6B1	1		
GATAD2B	1		
MAP4K2	1		
SAPS2	1		
TRABD	1		
ZBTB43	1		
HMHA1	1		
PRMT1	1		
CPSF1	1		