

A microarray configuration to quantify expression levels and relative abundance of splice variants

Pascale Fehlbaum, Caroline Guihal, Laurent Bracco, Olivier Cochet.

Supplementary material

Table S1: List of genes and identified splice events for which probes were designed. The LF/SF column indicates the long and short form for each transcript.

Gene	ACC#	Isoform	LF/SF
HSPC111	NM_016391	Wild type	LF
		Alternative acceptor site exon 4	SF
APEX	NM_001641	Wild type	LF
		Alternative donor site exon 1	SF
BIN1	NM_004305	Wild type	LF
		Skipping exon 12	SF
		Skipping exon 12 + 13	SF
		Skipping exon 12 + 13 + 14	SF
		Skipping exon 13	SF
MYBL2	NM_002466	Wild type	LF
		Skipping exon 8	SF
FKBP1A	NM_00801	Wild type	SF
		intron 4 retention	LF
DUSP3	NM_004090	Wild type	SF
		intron retention 3' UTR	LF
PIIB	NM_00942	Wild type	SF
		intron 4 retention	LF
NUDC	NM_006600	Wild type	FL
		Skipping exon 3 +4	SF
		Skipping exon 5 +6	SF
		Skipping exon 3+4+5+6	SF
APP	NM_000484	Wild type	LF
		Skipping of exon 7 & 8	SF
GSTZ1	NM_001513	Wild type	LF
		Skipping exon 5	SF
ROBO1	NM_002941	Wild type	LF
		Skipping exon 24	SF
EWSR1	NM_005243	Wild type	LF
		Skipping exon 8+9	SF
		Skipping exon 8	SF
PRKDC	NM_006904	Wild type	LF
		Skipping of exon 80	SF
CDC37	NM_007065	Wild type	SF
		intron 3 retention	LF
MGC8721	NM_016127	Wild type	LF
		Skipping exon 2	SF
MCL1	NM_021960	Wild type	LF
		Skipping exon 2	SF
hnRNPAB	NM_031266	Wild type	LF
		Skipping of exon 7	SF
hnRNPD	NM_031370	Wild type	LF
		Skippng of exon2	SF

