

Table S2. Transcription factor binding motif positions identified with functionally variant ($p < 0.05$) positions in both targets with single primary inputs and multiple primary inputs (for Affymetrix data only).

Tf	position	Score (F)	pvalue
SUM1	8	37.7401	0
SUM1	7	25.6358	0.002
ABF1	8	7.54051	0.023