

<b>-100-0nt</b>		<b>0-+100nt</b>	
Motif	Z-score	Motif	Z-score
AATAAA	5.1	TCTGTG	5.2
TTTTGT	4.6	CTGTGG	4.4
CTTTTG	4.3	TTCTGT	4.2
TGTATT	4.3	CTGTGT	3.9
TTGTAA	4.3		
ATAAAC	4.2		
ATAAAG	4.2		
TTAAAG	4.1		
TAAAGT	3.8		
TTTGTA	3.7		

**Figure S22. Comparison of the proximal PAS of Fip1 APA target vs. non-target.** Motif analyses of the proximal PASs of Fip1 APA target using the proximal PAS of non-target genes as background. Two regions (-100nt to 0nt and 0-+100nt, 0nt being the cleavage site) are analyzed separately. Enriched hexamer motifs and the corresponding Z-scores are shown. Z-scores above 3 are considered significant enrichment.