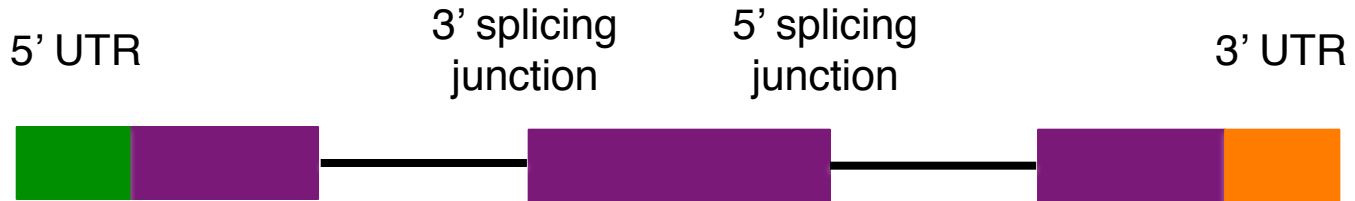


RBP binding enrichment: SUB1 and RBFOX2



	5' UTR	3' splicing junction	5' splicing junction	3' UTR	
$\frac{\text{Total length of the peaks in 5' UTRs}}{\text{Total length of 5' UTRs in the genome}}$	0.23%	0.23%	0.27%	1.06%	SUB1
	0.19%	0.17%	0.12%	0.27%	RBFOX2

RBP binding enrichment: all RBPs

