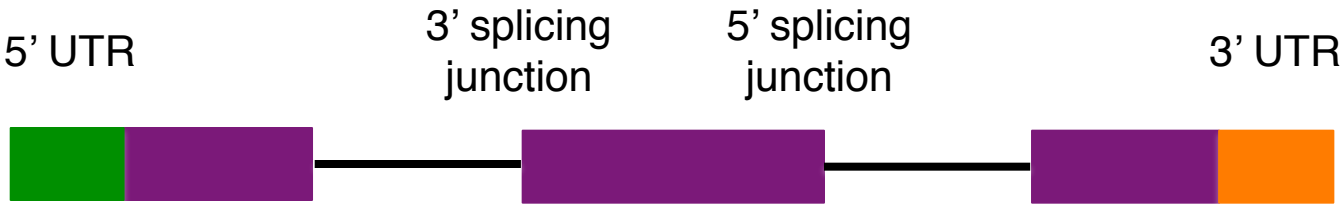
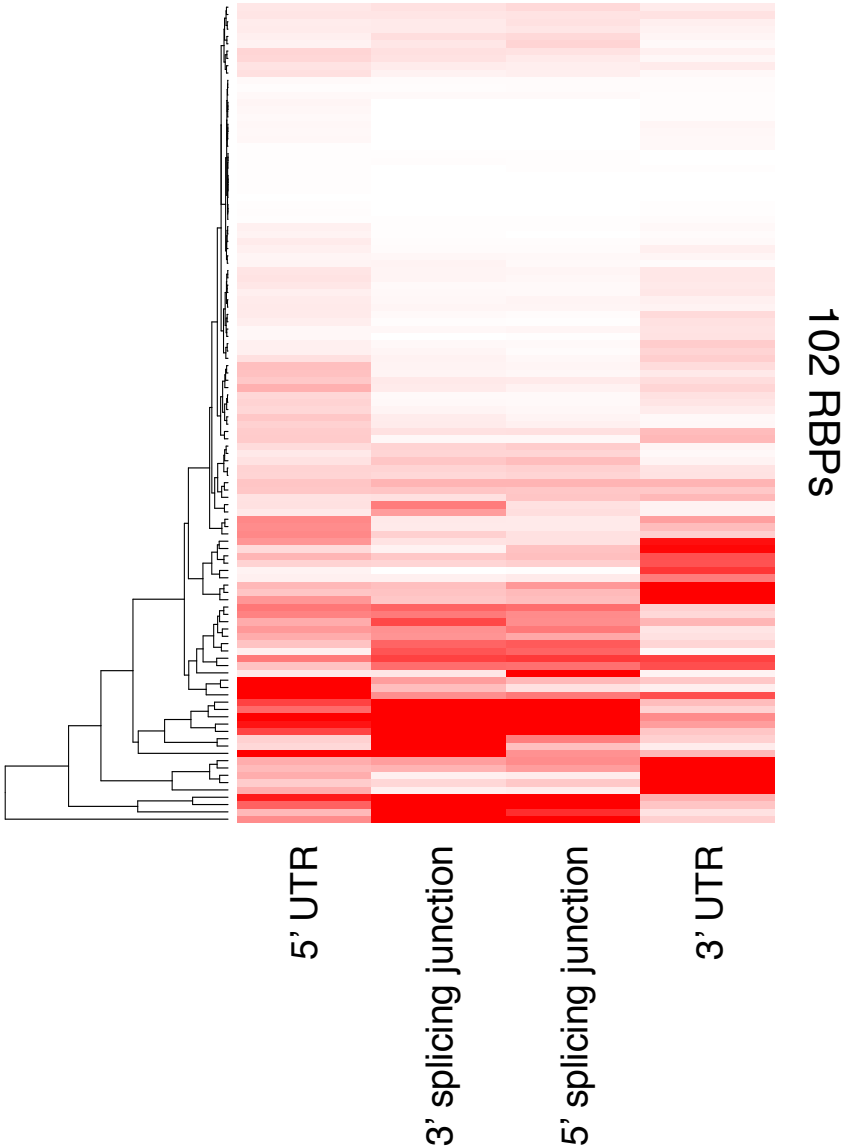
RBP binding enrichment: SUB1 and RBFOX2



$$\frac{\text{Total length of the peaks in 5’ UTRs}}{\text{Total length of 5’ UTRs in the genome}}$$

	5’ UTR	3’ splicing junction	5’ splicing junction	3’ UTR
SUB1	0.23%	0.23%	0.27%	1.06%
RBFOX2	0.19%	0.17%	0.12%	0.27%

RBP binding enrichment: all RBPs