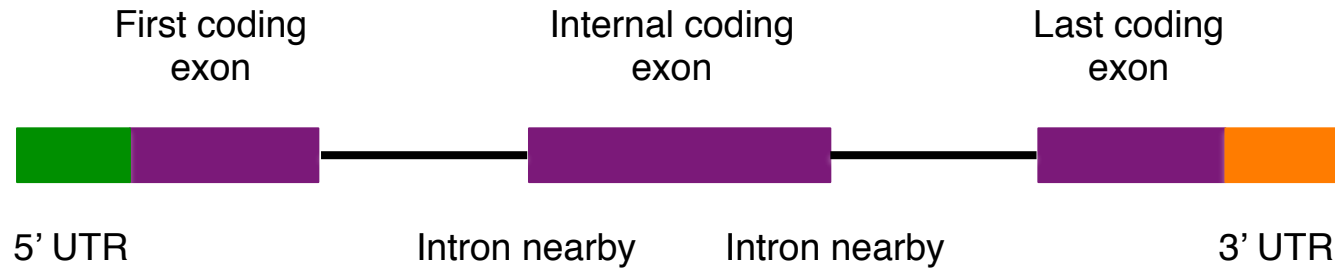


# RBP binding enrichment: SUB1 and RBFOX2



0.05%	0.29%	0.05%	0.42%	SUB1
0.13%	0.09%	0.14%	0.14%	RBFOX2
5' UTR	Coding exon	Intron nearby	3' UTR	

# RBP binding enrichment: all RBPs

