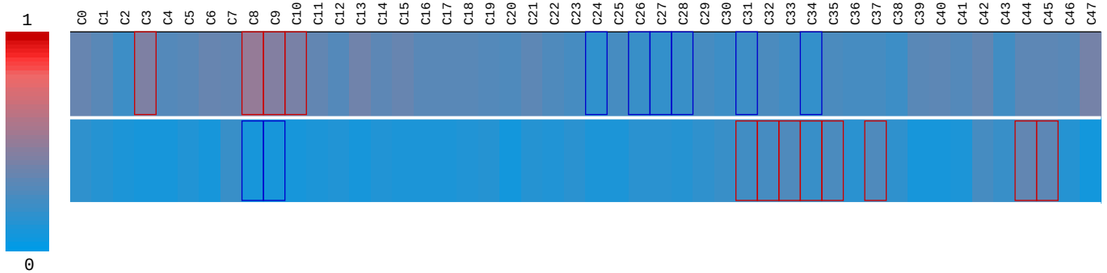
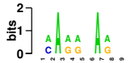
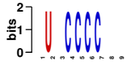


optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3’UTR	0.036	67.5	10/10	Y	↔	0.99	AAAAUAA	-	-
	3’UTR	0.017	29.3	10/10	-	↔	1.00	UUCCCCC	-	-