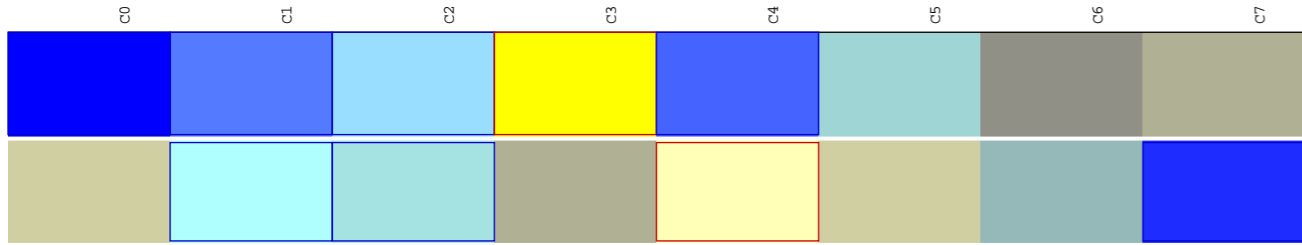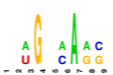over-representation

10

−10

	optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
		3'UTR	0.010	49.7	10/10	Y	↔	0.99	UUGAAAA	–	–
		3'UTR	0.004	17.5	10/10	–	→	0.99	UUCCCCC	–	–

C0 C1 C2 C3 C4 C5 C6 C7