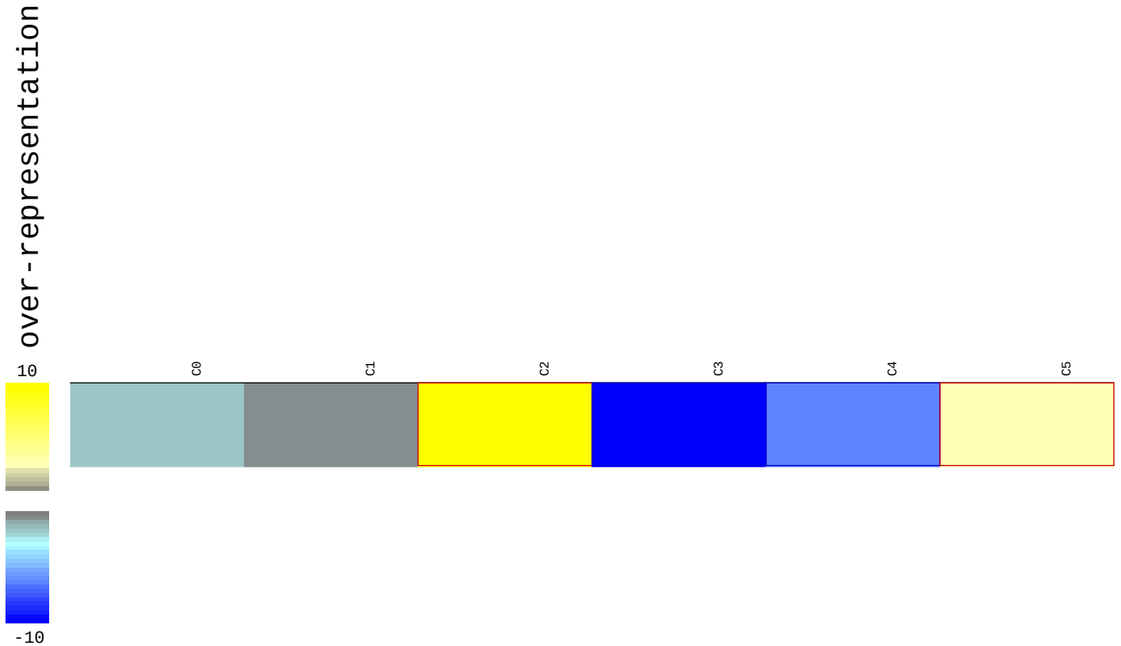

optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.011	64.8	10/10	Y	↔	0.98	UUUCAGA	-	-