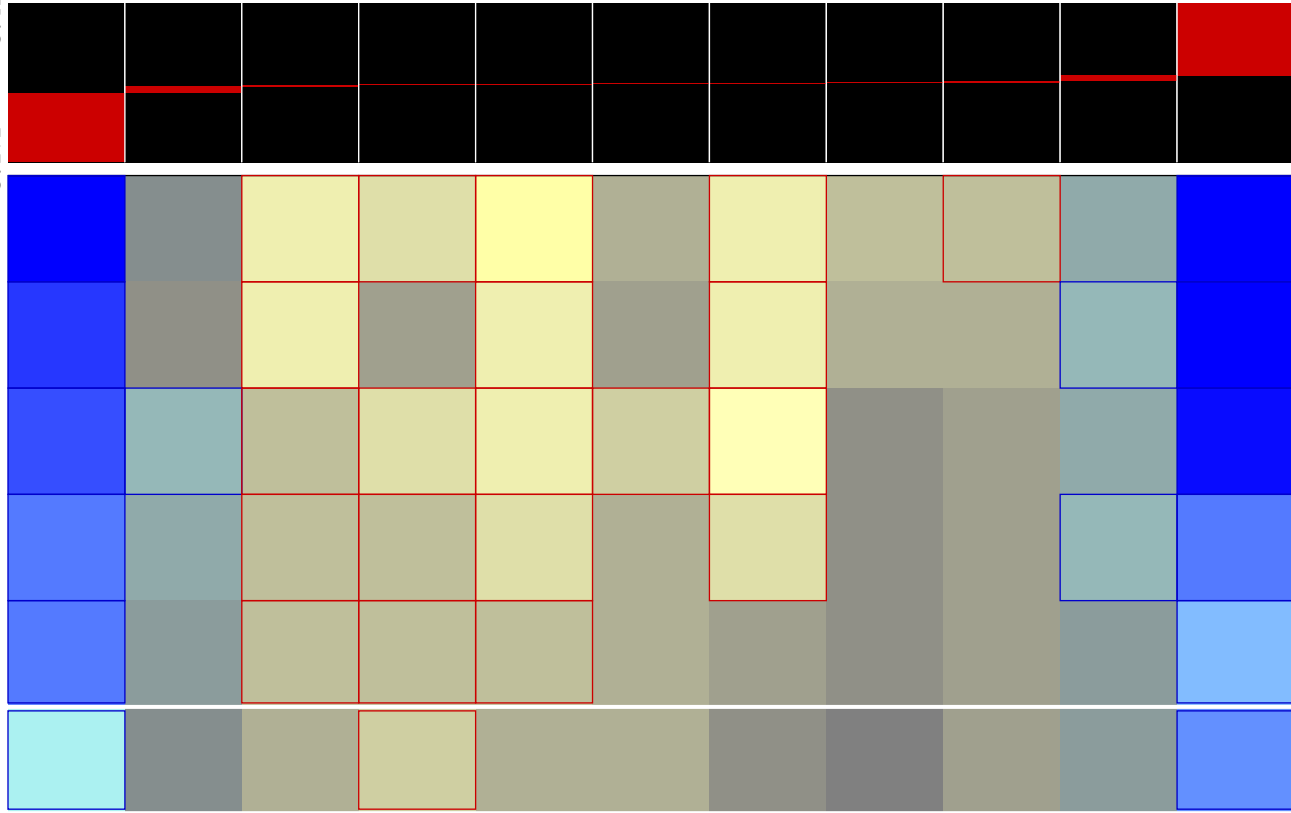
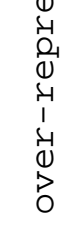
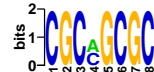

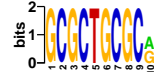
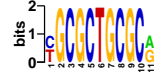




er-representation



over-representation



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.016	75.0	10/10	Y	-	-	-	-	-
	5'	0.011	49.9	10/10	-	-	-	-	-	-
	5'	0.011	48.8	10/10	-	-	-	-	-	-
	5'	0.008	36.7	10/10	-	-	-	-	-	-
	5'	0.006	27.7	10/10	-	-	-	-	-	-
	5'	0.005	21.0	10/10	-	-	-	-	-	-