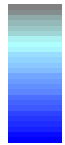


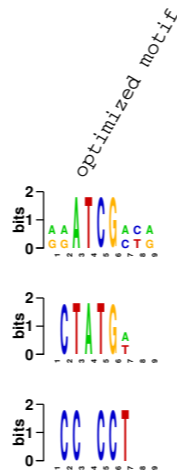
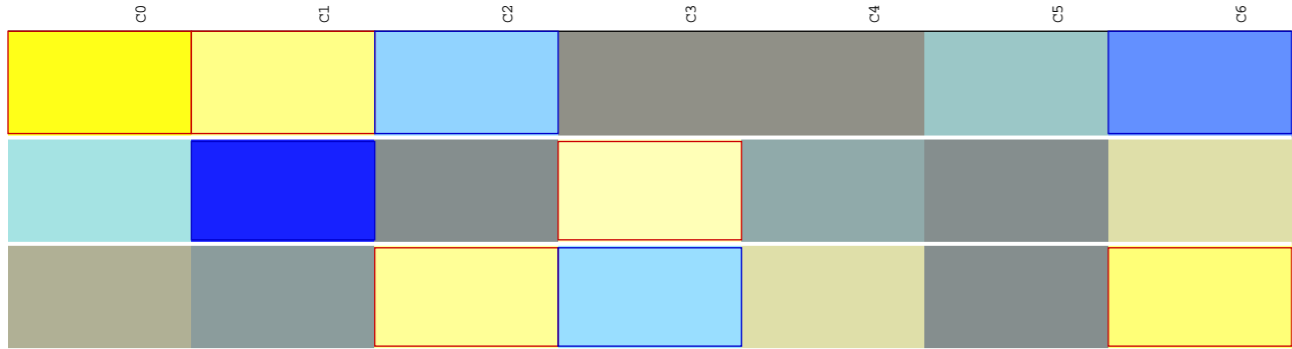
over-representation  
-10  
ation

10



-10

ation



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
AAAGACAC	5'	0.005	24.9	10/10	-	↔	1.00	AATCGAT	PIE_1
CTATGAA	5'	0.003	14.9	9/10	-	←	0.55	CTATGAA	-
CCACCTC	5'	0.003	12.9	9/10	Y	→	0.98	CCACCTC	ZTF_14

protein array