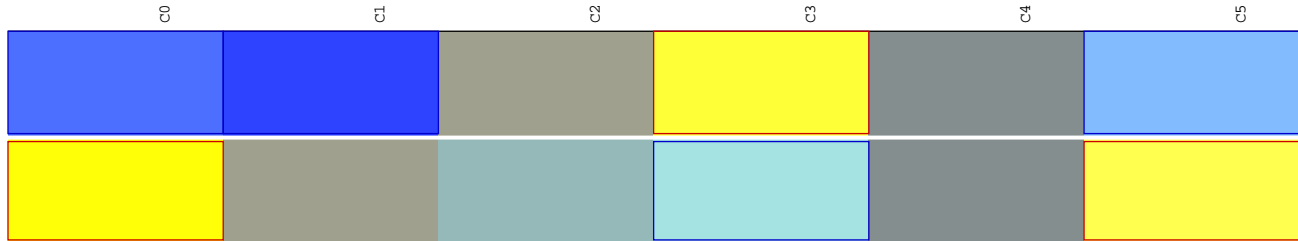
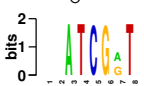
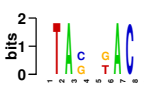


over-representation
on

10
-10



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.005	32.7	10/10	-	↔	0.99	AATCGAT	CEH_28, CEH-28, CEH_48 ...	-
	5'	0.004	21.5	10/10	-	↔	0.88	TACATAC	-	-