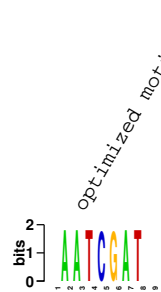
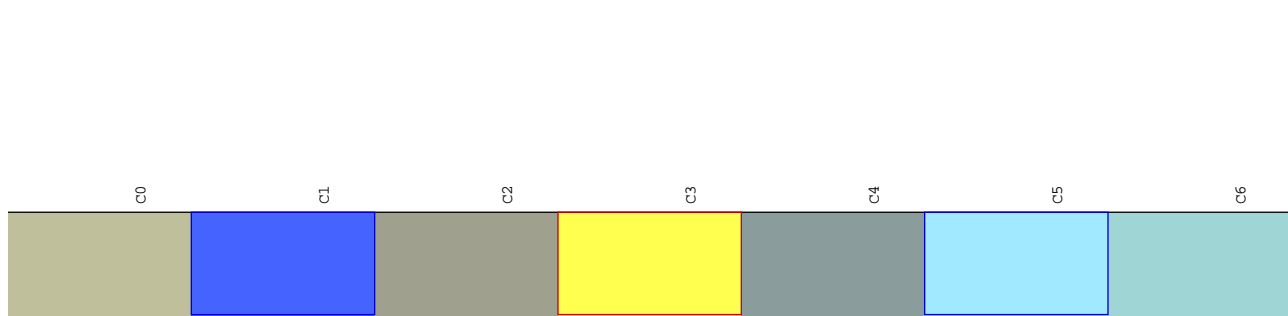


on
-10
10 over-representation



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
5'	0.004	19.7	10/10	-	↔	1.00	AATCGAT	CEH_28, CEH-28, CEH_38 ...	-