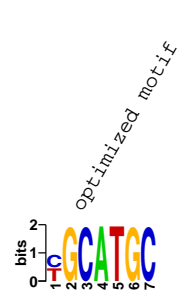
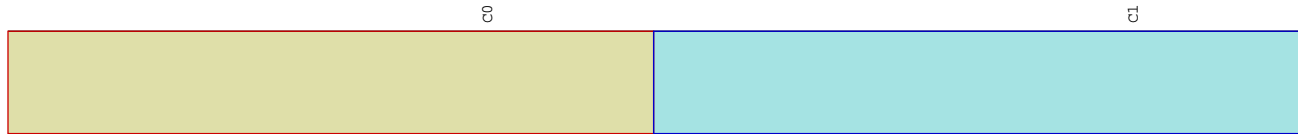


over-representation
on



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
5'	0.000	21.7	6/10	-	-	0.83		-

protein array
PA_343, PA_48, PA_62 ...