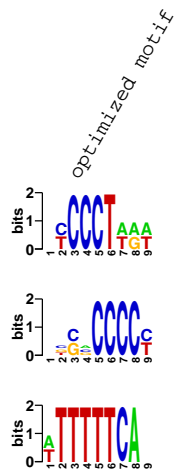
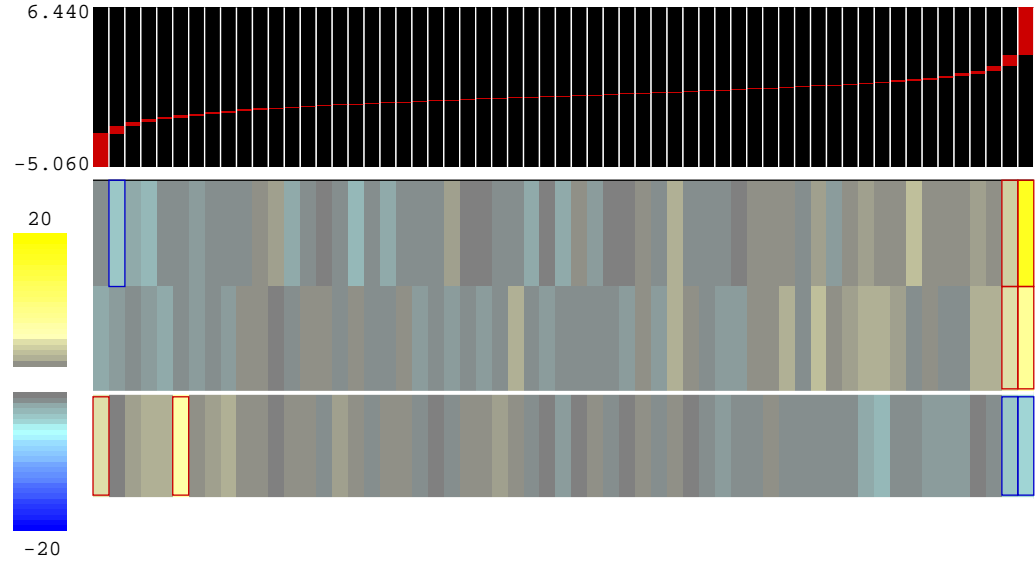


cell wall, $p < 1e-13$
translation, $p < 1e-09$
eukaryotic 43S preinitiation complex, $p < 0.0001$
cytosolic part, $p < 0.001$
cytoplasm organization and biogenesis, $p < 0.0001$
structural molecule activity, $p < 0.01$



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
5'	0.024	12.9	10/10	Y	-	1.00	CCCCTTA	-
5'	0.021	9.9	9/10	-	-	1.00	CCACCCC	MSN24
5'	0.017	7.4	7/10	Y	-	0.99	TTTTTCA	RRPE