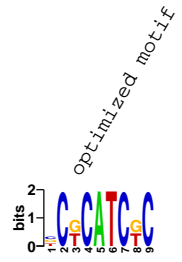


rRNA metabolism, $p < 0.001$
 ribosome biogenesis, $p < 1e-07$
 cytoplasm organization and biogenesis, $p < 1e-07$
 nucleolus, $p < 1e-04$
 ribosome biogenesis, $p < 1e-07$
 protein folding, $p < 0.001$



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
5'	0.036	20.9	10/10	Y	←	1.00	CTCATCG	PAC