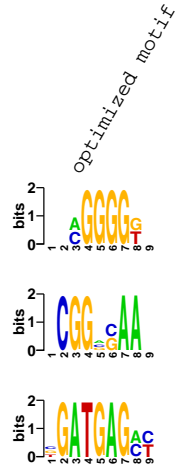


cytosolic part, p1e-04
cytosolic part, p1e-04



Location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
5'	0.026	14.4	10/10	Y	-	1.00	AAGGGGG	MSN24
5'	0.018	7.8	8/10	-	→	0.98	CGGGAA	-
5'	0.018	7.7	8/10	-	←	1.00	GATGAGC	PAC