

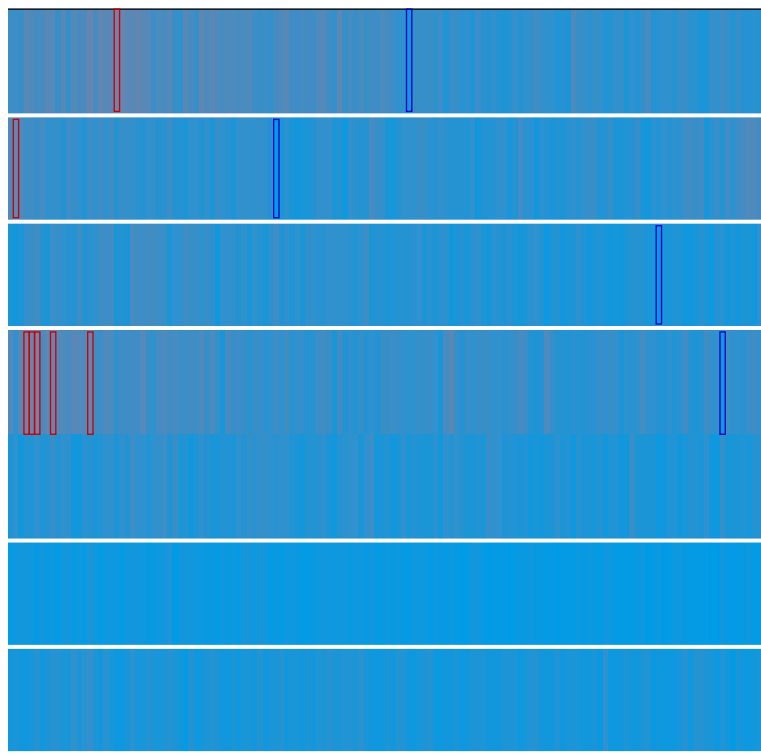
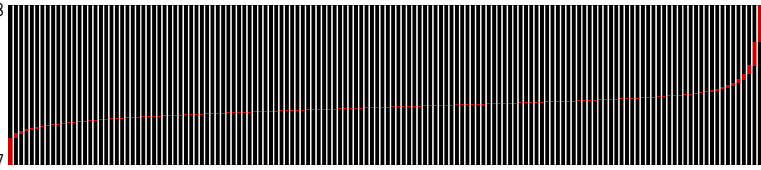
over-representation

under-representation



structural constituent of ribosome, p<le=07
 biological process, p<le=07
 ribonucleoprotein complex, p<le=01
 RNA processing, p<0.01

oxidoreductase activity, p<0.01
 phosphate transport, p<le=04



bits	Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name
1.8	1 A 2 G 3 U 4 CCC 5 C	3'UTR	0.019	13.9	10/10	-	→	1.00	UCCCCU	-
1.8	1 T 2 CC 3 T 4 TATC 5 A	5'	0.019	13.3	10/10	Y	-	1.00	TCTTATC	-
1.8	1 AA 2 G 3 G 4 CCCC 5 C	5'	0.018	12.0	10/10	-	-	1.00	AACGCGC	-
1.8	1 U 2 G 3 GUU 4 AA 5 AA	3'UTR	0.017	10.9	10/10	Y	→	1.00	UUGUUAU	-
1.8	1 C 2 G 3 G 4 G 5 AA 6 AA	5'	0.014	8.0	10/10	-	←	0.80	CAGCGAA	-
1.8	1 C 2 AC 3 T 4 CCG 5 G	5'	0.014	7.7	6/10	-	-	0.99	CACTCCG	-
1.8	1 G 2 A 3 G 4 GG 5 G 6 G	5'	0.013	7.1	6/10	-	→	0.83	ACAGCGT	-