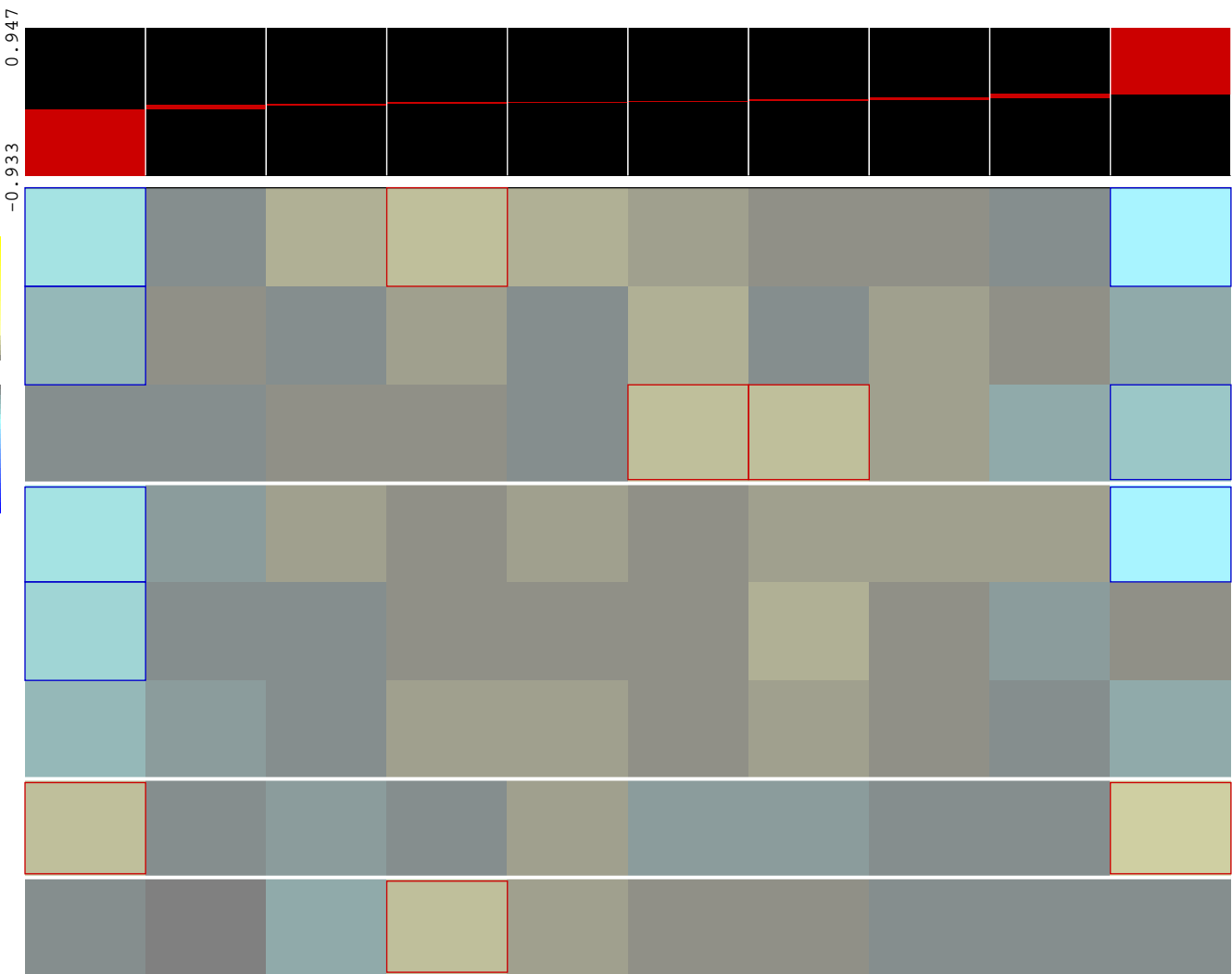

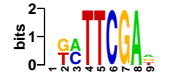
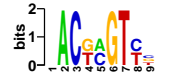
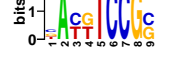
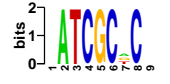
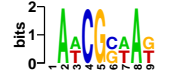




over-representation

under-representation



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.008	30.8	10/10	-	-	-	CCGCCGC	-	-
	5'	0.004	13.8	10/10	-	-	-	TCTTCGA	-	-
	5'	0.004	11.7	10/10	-	-	-	ACGAGTC	-	-
	5'	0.006	23.1	10/10	-	-	-	ACTTCCG	-	-
	5'	0.004	13.0	10/10	-	-	-	ATCGCTC	-	-
	5'	0.003	11.3	7/10	-	-	-	AACGGAA	-	-
	5'	0.004	11.5	9/10	-	-	-	TGCACAC	-	-
	5'	0.003	8.6	6/10	-	-	-	ACGCAAC	-	-