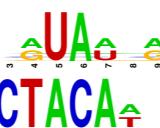
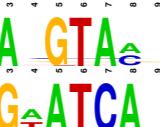
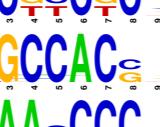
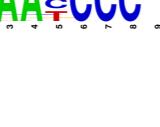


motif	location	MI (bits)	z-score	seed	motif name
	3'UTR	0.014	75.5	AAAUAAA	-
	5'	0.006	32.5	ACTACAA	NHR_28_NHR_53_NHR_70, UNC_39, NHR_256_NHR_79 ...
	5'	0.005	27.2	TATGTAC	CES-2_CES_2.2, CES_2, ZIP_7
	5'	0.003	17.4	TGAATCA	PIE_1
	5'	0.002	11.7	ACAGGTG	SNAI_1, HLH_1.4
	3'UTR	0.006	31.1	UUCCCCC	-
	5'	0.006	29.9	AATCGAT	PIE_1, CEH_28, CEH-28 ...
	5'	0.005	25.8	TCTTCTC	PIE_1
	5'	0.004	20.1	GGCCACC	CRH_2
	5'	0.002	11.1	TAATCCC	EOR_1, PIE_1

under-representation -10  10 over-representation