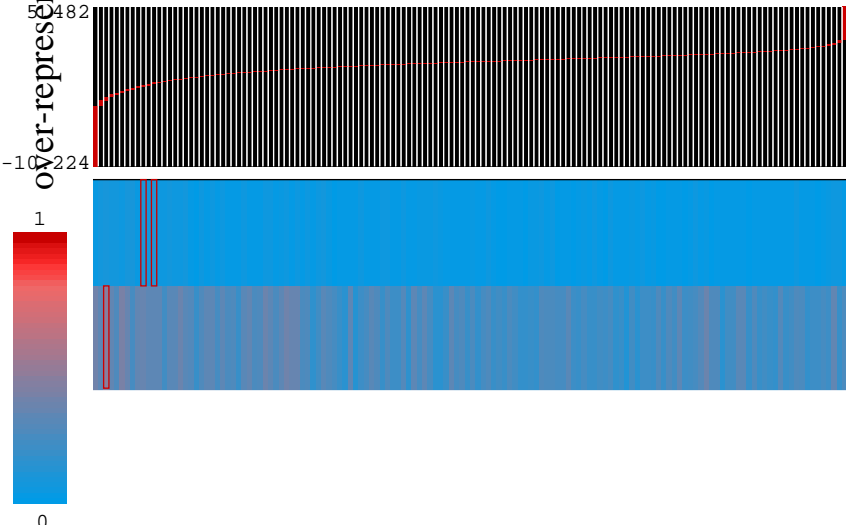
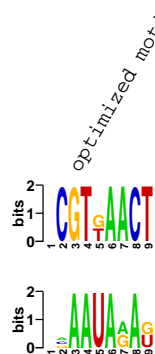


over-representation



macromolecule biosynthesis, p<1e-04  
phosphate transport, p<1e-05  
genitalia development, p<1e-04

macromolecule biosynthesis, p<1e-04  
genitalia development, p<0.01



Location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name
5'	0.015	8.2	9/10	-	→	0.89	CGTGAAC	-
3'UTR	0.015	9.2	7/10	-	→	1.00	AAAUAAA	-