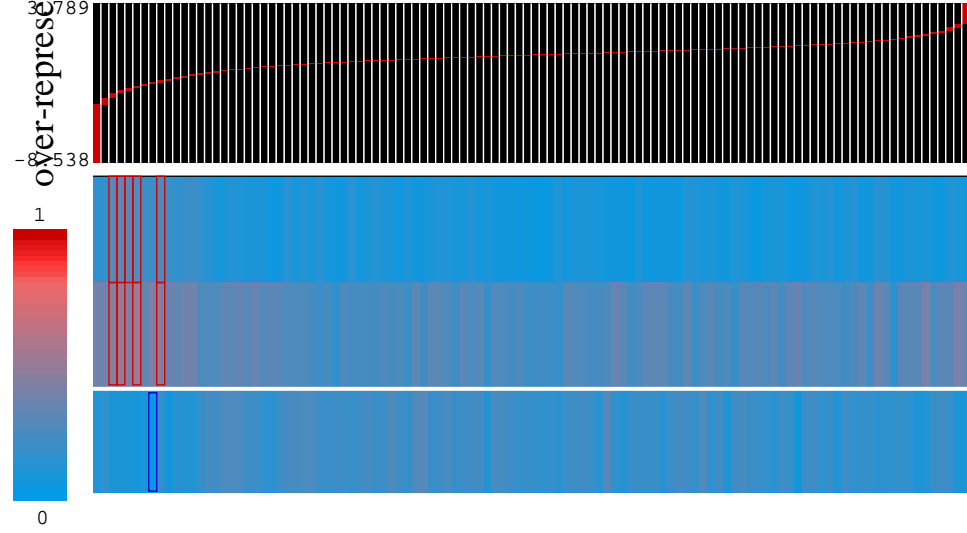
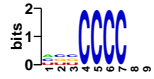
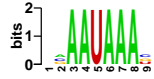
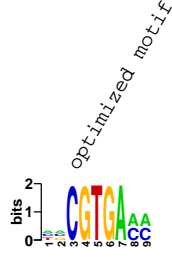


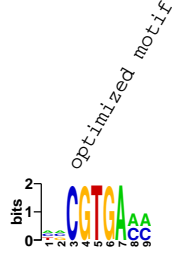
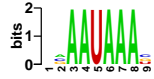
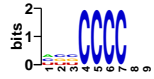
ation
over-representation

sugar binding, $p < 1e-16$
phosphate metabolism, $p < 1e-05$



structural constituent of cuticle, $p < 0.01$



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
	5'	0.025	18.9	10/10	Y	-	0.98	ACGTGAA	-
	3'UTR	0.017	10.1	10/10	Y	→	1.00	GAAUAAA	-
	3'UTR	0.016	8.7	8/10	-	→	1.00	UCCCCU	-