over-representation

structural molecule activity, p<1e-12
structural molecule activity, p<1e-08
structural constituent of cuticle, p<0.001

cytoplasmic part, p<0.01
non-membrane-bound organelle, p<0.001
larval development (sensu Nematoda), p<0.01
transferase activity, transferring hexosyl groups,

under-representation

optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
	3'UTR	0.022	15.8	10/10	–	→	1.00	UUUCCCC	–
	3'UTR	0.017	10.8	10/10	Y	→	1.00	AAUAAAU	–
	3'UTR	0.016	9.3	10/10	–	→	0.98	UUCUCUC	–
	5'	0.015	8.9	8/10	–	→	0.94	CTCTCTC	–
	5'	0.015	8.0	6/10	–	←	1.00	GCGACGC	–
	3'UTR	0.015	8.1	8/10	–	→	0.97	AAUAAAA	–