phosphate transport, p<1e-05

regulation of cell cycle, p<0.01

extracellular matrix, p<0.01
plasma membrane part, p<0.01

structural constituent of cuticle, p<1e-43
phosphate transport, p<0.07

over-representation

6858

-4831

1

0

under-representation

optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
	5'	0.019	14.1	10/10	–	–	0.99	AAGGCGC	–
	5'	0.018	12.3	10/10	–	–	1.00	AATCGAT	–
	5'	0.017	10.8	10/10	–	–	0.99	TTTCGCA	–
	3'UTR	0.017	11.1	10/10	–	→	1.00	UUCUUCU	–
	5'	0.016	10.6	9/10	–	←	0.96	TCGACGA	–
	5'	0.015	9.1	10/10	–	→	0.78	ATATGTA	–
	5'	0.013	7.1	9/10	–	–	0.93	AACGATG	–