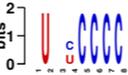


Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	Motif name	Protein array
 A A A A A C U A G G G G U C	3'UTR	0.024	119.5	10/10	Y	↔	0.99	AAAAUAA	-	-
 U U C C C C U U C C C C	3'UTR	0.007	34.4	10/10	-	↔	0.99	UUUCCCC	-	-