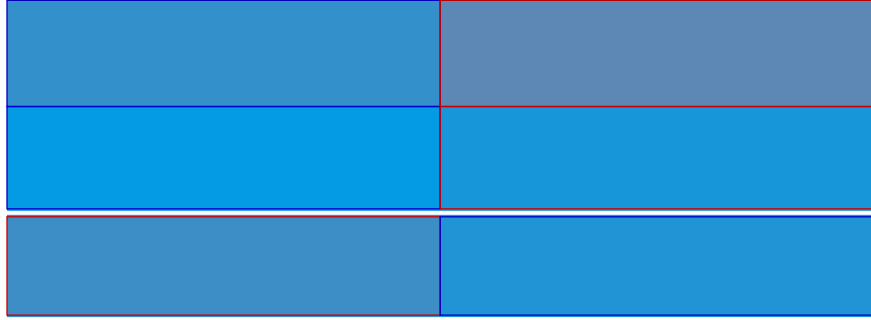




C0 oxidoreductase activity, p<1e-04

C1 nucleus, p<1e-22



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
<p>bits 2 1 0 1 2 3 4 5 6 7 8 9 C C A C C C C</p>	5'	0.015	28.5	10/10	-	-	0.99	CCACCCC	-
<p>bits 2 1 0 1 2 3 4 5 6 7 8 9 G G C G U G G</p>	3'UTR	0.011	19.7	9/10	-	→	0.96	GCGUGG	-
<p>bits 2 1 0 1 2 3 4 5 6 7 8 9 T C G A C T C</p>	5'	0.010	18.7	10/10	-	-	0.30	TCGACTC	-