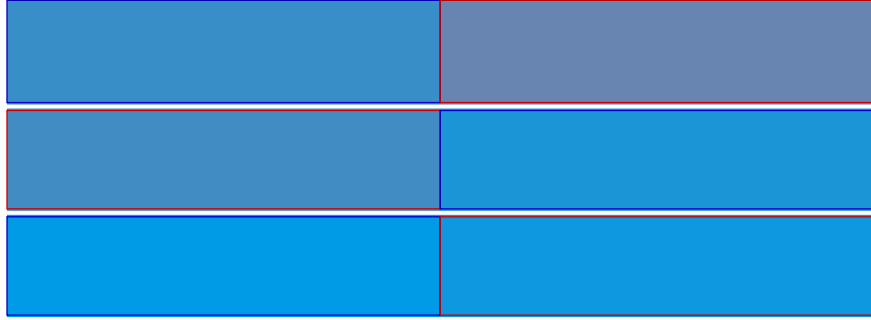







C0 cell organization and biogenesis,  $p < 0.001$

C1 mitochondrion,  $p < 1e-15$



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</p>	5'	0.017	31.4	10/10	Y	-	0.80	CACGATA	-
 <p>bits 2 1 0 1 2 3 4 5 6 7 8 9</p>	3'UTR	0.012	23.0	9/10	-	→	0.97	AAGAAAA	-
 <p>bits 2 1 0 1 2 3 4 5 6 7 8 9</p>	3'UTR	0.011	20.0	6/10	-	→	0.43	AUGAAAG	-