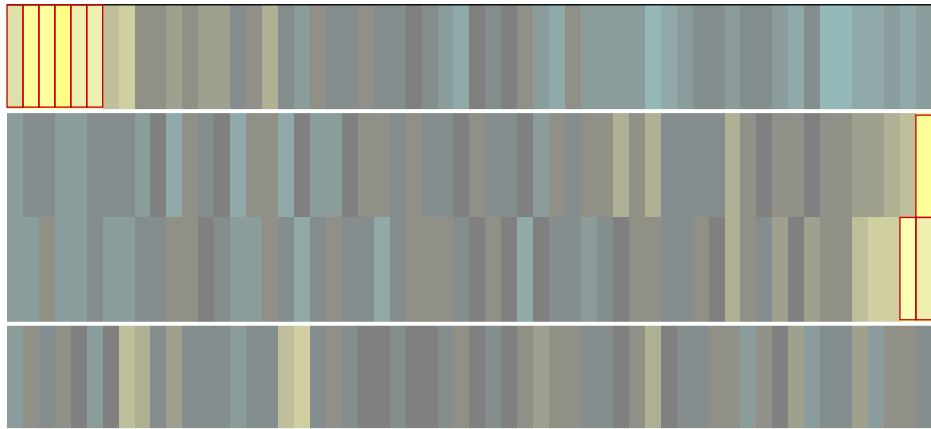


-20

cytosolic ribosome (sensu Eukaryota), p<1e-30  
 cytosolic ribosome (sensu Eukaryota), p<1e-27  
 cytosolic part, p<1e-07  
 cytoplasm organization and biogenesis, p<1e-09  
 'de novo' IMP biosynthesis, p<0.01  
 translation initiation factor activity, p<0.001

generation of precursor metabolites and energy, p<  
 energy derivation by oxidation of organic compound



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name
	5'	0.037	22.8	10/10	Y	-	1.00	AAAATTT	RRPE
	5'	0.029	16.1	10/10	Y	-	1.00	CCCTTAA	RRPE
	5'	0.027	14.8	10/10	Y	-	1.00	AGGGGAG	MSN24
	3'UTR	0.016	6.0	6/10	-	→	0.90	GUGGUUG	-