

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q99KG3	RBM10_MOUSE	Mus musculus	RNA-binding protein 10	38.223321	S515	S61;S89;S718;S723;S733;S736;S738;S781;S797;S845	40997131	MEYERRGGRGDRGTGRYGATDRSQD DSGENRSRDHYDRMDYRSYPREY GSQEGKHEYDDSSSEEQSAEDSYEAS PGSETQRRRRRRHRHSPTGPPGFPR DGDYRDQDYRTEQGEEEEEEEEDEE EEEKASNIVMLRMLPQAATEDDIRG QLQSHGVQAREVRLMRNKSSGQSR GFAFVEFSLQDATRWMEANQHSL NILGQKVMHYSDPKPKINEDWLCN KCGVQNFKRREKCFKCGVPKSEAEQ KLPLGTRLDQQALPLGGRELSQGLL PLPQPYQAQGVLTSQLSQGSEPPSS ENANDTIILRNLNPHSTMDLSILGALA PYAVLSSSNVRVIKDKQTQLNRGFA FIQLSTIVEAAQLLOILQALHPPLTID GKTINVEFAKGSKRDMSNEGSRIN AASVASTAIAAAQWASQASQGGESA WAAPEEPPVDYSYYQQDEGYGSSQG TDSL YAHGYLKNKSGPGMTGKGD AGTGP EASLEAGADSVSLQAFSRAQ PGAAPGLYQQSAEGSSGQSTATNSQ SYTIHSPAVLKAELQSPTOPSSSAFPP ATSPTAPEAYSQYPVPDVSTYQYDET SGYYDPQTGLYD PNSQYYNAQS QQYLYWDGERRTYIPALEQSADGHK DTGASSKEGKEKKEKHTKTAQQIA KDMERWARSLNKQKENFKNSFQPI SALRDDRRESATADAGYAILEKKG LAERQHTSMDLPKASDDRPSPPRG LVAAYSGESDSEEEQERGPEREEK LTDWQKLACLLCRRQFPSKEALIRH QQLSGLHKQNL EIHRRRAHLSENELE ALEKNDMEQMKYRDRAAERREKYG IPEPPEPKRRKYGGISTASVDFEQPT RDGLGSDNIGSRMLQAMGWKEGSG LGRKKQGIPTIEAQTRVRSGLGAR GSSYGVSTESYKETHKTMVTRFN EAQ	True	False	1.212	4.18	1.205	1.033	False	1.103	1.143