

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | ei re |
|--------------|-------------|--------------|------------------------|--------------|---|-------------------------------|---|--|---------------|---------------|---------|---------|---------------|-------|
| Q9NTZ6 | RBM12_HUMAN | Homo sapiens | RNA-binding protein 12 | 13.643945 | S67;S103;S111;T121;T122;S124;S130;S132;T135;T137;T138;S139;S143;T149;T178 | S352;S375;S420;S422;S424;S525 | 31637018;35132862;35254053;34725712;23301498;32119511;34019948;30059200;32574038;30379171;37340703;22661428;38665916;36240223;28510447;39302247;35289036;29351928;27655845;39531497;33214551;31492838;34229054;35138101 | MAVVIRLQGLPIVAGTMDIRHFFSGL TIPDGGVHIVGGELGEAFIVFATDDED ARLGMMRTGGTIKGSKVTLLSSKT EMQNMIELSRRRFETANLDIPPANA SRSGPPSSGMSSRVNLPPTVSNFN NPSPSVVTATTSVHESNKNIQTFSTA SVGTAPPNMGASFGSPTFSSTVPST ASPMNTVPPPIPPIPAMPPLPPMPS IPPIPVPPVPTLPPVPPVPIPPVPSV PPMTPLPPMSGMPPLNPPVAPLPA GMNGSGAPMNLNNLNPMFLGPL NPVNPQMNSQSSVKPLPINPDDL VSVHGMPPFSAMENDVRDFFHGLRV DAVHLLKDHVGRNNGGLVKFLSP QDTFEALKRNRMLMIQRYVEVSPAT ERQVVAAGGHITFKQNMGPSGQTH PPPOTLPRSKSPSGQKRSRSPHEA GFCVYLKGLPFEAENKHVIDFFKKL DIVEDSIYIAYGPNKATGEGFVEFR NEADYKAALCRHKQYMGNRFIQVH PITKKGMLEKIDMIRKRLQNFSDQ REMILNPEGDVNSAKVCAHITNIPFS ITKMDVLQFLEGIPVDENAVHVLVD NNGQGLGQALVQFKNEDDARKSER LHRKKLNGREAFVHVVTLEDMREIE KNPPAQGKGLKMPVPGNPAVPGM PNAGLPGVGLPSAGLPGAGLPSTGLP GSAITSAGLPGAGMPSAGIPAGGEE HAFLT VGSKEANNGPPFNFPGNFC GSNAFGPPIPPGLGGGAFGDARPG MPSVGNLGLGLDVPFGGGGPN NLSGPGSGGGPQNFNGNGPSLGG PPGFGSGPPGLGSAFGHGGPPAFG PGPGPGPGPIHIGGPPGFASSGK PGPTVIKVNMPFTVSDIEILDFYF YQVIPGSVCLKYNEKGMPTGEAMVA FESRDEATAAVIDLNDRPIGSRKVKL VLG | True | False | 1.629 | 4.823 | 0.706 | Fi |