| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatus | plasma membrane | ext reg |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | Q69ZX6 | MOR2A_MOUSE | Mus musculus | ATPase MORC2A | 37.55213 | T811 | T582;S614;S703;S728;T731;S737;S741;S775;S777;T834;S854;S859 | 41018975;40885482 | MAFTNYSSLNRAQLTFEYLHTNSTTHEFLFGALAELVDNARDADATRIDIYAERREDLRGGFMLCFLDDGAGMDPSDAASVIQFGKSAKRTPESTQIGQYGNGLKSGSMRIGKDFILFTKKEDTMTCLFLSRTFHEEEGIDEVIVPLPTWNARTREPITDNVEKFAIETELVYKYSPFHTEEQVMNQFMKIPGNSGTLVIIFNLKLMDNGEPELDIISNPKDIQMAETSPEGTKPERRSFRAYAAVLYIDPRMRIFIHGHKVQTKRLSCCLYKPRMYKYTSSRFKTRAEQEVKKAEHVARIAEEKAREAESKARTLEVRMGGDLTRDSRVMLRQVQNTAITLRREADVKKRIKDAKQRALKEPKELNFVFGVNIEHRDLDGMFIYNCSRLIKMYEKVGPQLEGGMACGGVVGVVDVPYLVLEPTHNKQDFADAKEYRHLLRAMGEHLAQYWKDIAIAQRGIIKFWDEFGYLSANWNQPPSSELRFKRRRAMEIPTTIQCDLCLKWRTLPFQLSSVETDYPDTWVCSMNPDPEQDRCEASEQKQKVPLGTLKKDPKTQEEKQKQLTEKIRQQQEKLEALQKTTPIRSQADLKKLPLEVTTRPIEEPVRRPQRPRSPPLPAVIKNAPSRPPSIQTPRPSTQLRKTSVISLPKPPTTAARGETSTSRLLQPTEAPRKPANPPIKTVPRPTPPVHTPPLSLIPSSKSLREVPAQKAIKTPVVKKPEPPVKQSVATSGRKRSLAVSDEEEAEEEAEKRRERCKRGKLAVKEEKKEANELSDSAGEDHPAELRKAQKDKGLHVEVRVNREWYTGRVTAVEVGKNAVRWKVKFDYVPTDTTPRDRWVEKGSEDVRLMKPPSPEHQSPDTQQEGGEEEEAMVARQAVALPEPSTSDGLPIEPDTTATSPSHETIDLLVQILRNCLRYFLPPSFPISKKELSVMNSEELISFPLKEYFKQYEVGLQNLCHSYQSRADSRAKASEESLRTSEKKLRETEEKLQKLRTNIVALLQKVQEDIDINTDDELDAYIEDLITKGD | True | False | 3.15 | 4.499 | 0.719 | 0.281 | False | 1.688 | Fal |