	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	extracellular region
0	Q8C5W4	MOR2B_MOUSE	Mus musculus	ATPase MORC2B	36.83566	S973	S615;S690;S724;S733;S737;S768;S770;T827;S846;S851	40885482	MAFTNYSTLNRAQLTFDYLHTNSTTHAFLFGALAELIDNARDADATRIDIYAEKREDLQGGFMLCFLDNGVGMDPNDVINVIQFGKSAKRTPESTQIGRYGNGLKSGSMRIGKDFILFTKKENTMSCLFLSRTFHEEEGIDEVIVPLPTWNSQTREPVTDNMEKFAIETELIYKYSPFHTEEEVMTQFTKISGTSGTLVVIFNLKLTDNGEPELDVTSNPKDIRMAEISQEGVKPERHSFCAYAAVLYIDPRMRIFIHGHKVQTKKLCCCLYKPRKYTFTSHRFKTRAEQEVKKADQVAQLAEEKAREAESKARTLEIHMGGDITRDSRVMLRQVQNTAITLRREADVKKRIRDAKQQALKEPKELTFVFGVNIEHRDHDGMFIYNCSRLIKMYEKVGPQLEKGMVCGGVVGVIDVPYLVLEPTHNKQDFADAKEYRHLLRAMGEHLAQYWKDIEIAQHGIIKFWDEFGYLSANWNRPPSDELHFKRKRAMQVPTTIQCDLCLKWRTLPFQLSAVEEGYPINWVCSMNPDPEQDQCEAFELKQKIPLGILKKAPKTQEERQKQLTEKIQQEQRKLKALKKIKPIHSQSDLKKLPLEVTSRPFSKYPAHIFQGPQSSFHVVKTNARRRPQSRHAPFRQLQRSSIICTNPKPPFLVDKTEAVLLQPPETPQKSVSLLVKTIPQPPPLVQSLSPSVVPKSNNPWKVETPQIMNTPVAEMPYVPVNPSLVICDHKRSPEVSDEIEDEDRRKRMCKRGRFTVKKEKIQASELSDSSGEENPVDLKTAQKDKGLYVEVRMMGECYKGHVTAVEVGDNVVWWKVKFEDMPKDSTPRDCWVEKGSENVWLVKPSPEYQSTDEQQEDRKGEEDTVVQQALALQQTSTSECFCTEPDTTASTANHKTIDLLVQILWNCLHYFMPLSFPISKKELGAMNSEELLSLPLKECFKQYEVGLQNLCRSYQRCADSQAKVSEESLRISQKKLQETEEKLQKLRTNIQTLLQMAQQGINIRADDELDAYIEDLVSSDD	True	False	0.188	4.428	0.234	0.188	False	2.062	False