

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
E9PZW0	E9PZW0_MOUSE	Mus musculus	NaN	25.161187	NaN	NaN	34887587	MSCNGGSHPRINTLGRMTRAESGP DLRYEMTYSGGGGGGGGGGGT RTFYSHSRRCTVNDQNSDGYCQTGT MSRHQNQNTIQEMLQNCSDCLMR AELIAQPELKFGEQMQLAWNRELDE YFTQANDQMEIIDGLIREMRQMGQP CDAYQKRLLQLQEQMRALYKAISVP RVRRASSKGAGGYTCQSGSGWDEFT KRLTGECLGWMRQREEMDLMAW GVDAGSVEQHINSHRSIHNTIGDYR WQLDKIKADLREKSAIYQLEEEYENL LKASFERMMDHLRQLQNIQATSREI MWINDCEEEELLYDWSKNTNIAQ KQEAFSIRMSQLEVKEKELNKLKQE SDQLVLNQHPSADKIEAYMDTLQTO WSWILQITKCIDVHLKENAAYFQFFE EAQSTEAYLKGLQDSIRKKYPCDKN MPLQHLLEQIKELEKEREKIIEYKRQ VQNLVNSKSKIVQLKPRNPDYRSNK PILRALCDYKQDQKIVHKGDECILK DNNERSKWYVTGPGGVDMLVPSVG LIIPPPNPLAVDLSCKIEQYEAIAL WNQLYINMKSLVSWHYCMIDIEKIR AMTIAKLKTMRQEDYMKTIEDLELH YQDFIKNSQGSEMFGDDDKRRMQS QFTDAQKHYQTLVIQLPGHPQHQT TKTEITHLGTCCQDVNHNKVIETNRE NDKQETWLLMELQKIRRQMEHCEA RMTLKNLLAEQGSTHHITVKINEL KSVQNSQALAEVLNQLKDMLANF RGSEKYCYLQNEIFGLFQKLENING VSDGYLNSLCSVRALLQAILQTEDM LKVYEARLTEEETVCLDLKVEAYRC GLKKIKNDLNLKSLLATMKTLELQK AQQIHSQSSQYPLYDLDLGKFTEK VTQLTDRWQKIDKQIDFRLWDLEKQ IKQLRNYRDNYQSFCKWLYDAKRRQ DSLESMKFGDSNTVMRFLNEQKNL HSEISGKRDKSEEVHKAELCANSIK DYELQLASYTSGLETLNIPKRTMV QSPSGVILQEAADIHARYIELLTRSG DYRFLSEMLKSLEDLKLKNTKIEVL EEELRLARDANSENCKNKFLDQN LQKYQAECQFKAKLVSLLELKRQA ELDGSQAKQNLKCYGQIKELNEKI TRLTYEIEDEKRRRKTVEDRFDQK NDYDQLQKARQCEKENLSWQKLES EKAIKEKEYEIERLRLVLLQEEGARKR EYENELAKASNRIQESKQCTQVVQ ERESLLVKIKVLEQDKARLQRLDEL NRAKATLEAESRVKQRLCEKQIQ NDLNQWKTQYSRKEETIRKIESERE KSEREKNSLRSEIERLQAEIKRIEER CRRKLEDSSRETQSQLESERCRLQK EIEKLQRPYCSHRETQTEYEWTV SSKLVFDGLRKKVTAMQLYEQCLID KTTLDKLLKGGKSVEEVASEIQPFLR GAGAIAGASASPKEKYSVLEAKRKKF ITPESTMMLLEAQAATGGIIDPHRNE KLTVDNAVARDLIDFDDRQQIYTAEK AITGFDDPFSGKTVSVSEAIKKNLID RETGMRLLEAQLASGGVDPVNSVF LPKDVALARGLIDRDLRSNDPRDS QKNFVDPITKKKVSVMQLRERCRIE PHTGLLLSVQKRSMQFQIRQPV VTELVDGILRPSTVNELESGQISYD EVGERIKDFLQSSCIAGYNETTKQ KLGIEAMKIGLVRPGTALELLEAQA ATGFVDPVSNLRLPVEEAYKRGVLV IEFKEKLLSAERAVTGYNDPETGNIIS LFQAMNKELIEKGHIRLLEAQIATG GIIDPKESHRLPVD MAYKRGYFNEE LSEILSDPSDDTKGFFDPNTEENLTY LQLKERICDEETGLCLLPLKEKKKQ	None	None	None	None	None	None	None		

