

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
F8VQM5	F8VQM5_MOUSE	Mus musculus	NaN	25.875327	S135	NaN	30059200	MAKKKFSGLEISLIVLFIIVTHIAIALVV VLATKVPAAVEEVKSPSTSPSPGRCPPEQGEPLNERINCIPEQHPTKAKCEE RGCCWRPWNNTIIPWCFFADNHGY TAASVTNDNSGLKATLSRIPSPTLFG EDIKSVLLTTQSQTRNRFRFKLTDPN NKRYEVPHQFVKDGNPAAADTLYD VKVSENPFSIKVIRKSNNKVLFDTSI GPLVYSNQYLQISTRLPSEYIYGFG EHIHKKRFRHDLYWKTWPIFTRDEIPG DNNHNLYGHQTFMGIEDNSGKSY GVFLMNSNAMEVFIQPTPIITYRVTG GVLDIFYIFLGDTPQVQVQYQELIGR PAMPAYWNLGFQLSRWNYVSLDKV KEVVRNREAGIPYDAQVTDIDYME DKKDFTYDEVAFKGLPEFAQDLHNNH GQKYIILLDPAISINKRANGAEYQTYV RGNEQNVVWKESDGTTSLIGEVWP GLTVYDFTNPRTWEEWANEENLNF HQQVEYDGLWIDMNEVSSFIHGSQ KGCAPNLLNYPFTPGILDKIMYSKT LCMDAVQHWGNQYDVHSLYGYSM AIATEKAVEKVFVNKRSPILTRSTFA GSGHAAHWLGDNTASWEQMEWS ITGMLEFGMFGMPLVGADICGFLAN TTEELCRRWMLGAFYFPSRNHNA EGYAEQDPAFWGADSLVNTSRHYL TIRYTLPLFLYTLFYRAHAFGETVARP FLHEFYEDPNSWIEDTQFLWGPALL ITPVLRPETKYVSAYIPDATWYDYETG EKRPWRKQRVDMYLPEDKIGLHLR GGYIIPQPDVTTTASRKNPLGLIVA LDENQAAKGELEFWDGKSDTIEKK IYLYEFSVSNLIVNCTHSSYPEG NTLVFKTIKVLGLSATVAVTVGEND QQMNPFLAFTFDANLKSITDLTF NLGKTFIVRWTTQSFSDNEKFTCYP DVGATEKTCVERGCIWEPVSGLAN VPPCYFNSNHNPLYLLTSTQKLATGIT AELQLNPASARIKLPNPISTLRVEV KYHKNDMLQFKIYDAHKKRYEVPVP LNIPDTPTSSEENRLYDVEIKENPFG IQVRRRSTGKLIWDSCLPGFAFNDQ FIQISTRPLPSQYLYGFGEAHTAFKR NLNWHWTGWMFTRDQPPGYKLSY GFHPYMALEDEGNAHGVLLQNSN GMDVTFQPTPALTVRTIGGILDFYMF LGPTPEGATKQYHEVIGFPVMPYVW ALGFQLCRYGYRNTSEIEQLYNDMK AAQIPYDVQYTDINMYERQLDFTIGE RFKTLQPVEKIRKEGMKYIVILDPAI SGNETQYPYPAFERGIQKDVFKWPN TNDICWAKVWPDLPNITIDEITTEDE AVNASRAHVAFPPDFRNSTSEWWT REIYDFYNEKMKFDGLWIDMNEPSS FVNGTVTNKCRNDTLNPPYPPELT KRNEGLHFRMCMETEHLSDGSS VLHYDVHNLYGWSQVKPTLDALRN TTGLRGIVISRSTYPTAGRWGGHWL GDNYANWENLEKSLIGMLEFNLFGI PYVGADICGFNDSEYHLCARWMQ VGAFYPYSRNHNQFTRRQDPVSWN ETFAQMSKKVLEIRYTLPPYFYTQM HEAHIHGGTVIRPLMHEFFDDKETW EYKQFLWGPAPMVPVIGPFQTAV NGYVPKARWFDYHTGEDIKVRGKLO TFSAPFDTINLHVRRGGYILPCQEPAQ NTYYSRQKYMKLIVAADDNQTAKGT LFWDDGESIDTYEKNQYTLIEFNLN QKNLTSTVLKDGYNKSEMKLGSY VWGKGTTSISQVNLTYGGNEQQLSF TQDEAKEILTIDLKNVTVTLDEPIQIS WS	None	None	None	None	None	None	None		