

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
E9QJU8	E9QJU8_MOUSE	Mus musculus	NaN	29.185005	NaN	NaN	22645316	MRFGSKMMPMFLTVLYLSNSEQHFT EVPVTPETICRDVVDLCKEPGENDC HLAEVWCGSERPVADNERMFDVLQ RFGSORNEVRFRLRHERPPNRDIVS GPRSQDPSVKRNGVKVPGEHRRKE NGVNSPRLDLTLAELQEMASRQQQ QIEAQQQMLATKEQRLKFLKQODQ RQQQAAAEQEKLKRLRELAESQEAK LKKVRALKGHVEQKRLSNGKLVVEI EQMNSLFQQKQRELVLAVSKVEELT RQLEMLKNGRIDGHHDNQSAAVEL DRLYKELQLRNKLNQEQNAKLQQQ RECLNKRNSEVAVMDKRVSELRDR LWKKKAALQKQENLPVSPDGNLPQ QAVSAPSRVAAVGPYIQSSTMPRMP SRPELLVKPALPDGSLLMQSAEGPM KIOTLPNMRSAAASQSKGSKAHPAS PDWNPNSADLLPSQGSSVPQSAGTA LDQVDDGEIAVREKEKKVRPFSMFD TVDQCAAPPSPFGTLRKNQSSSEDILR DAQAVNKNVAKVPPVPTKPKQIHL PYFGQTAQSPSDMKPDGNAQQLPIA ATSVGAKLKPAGPQARMLLSPGAPS GGQDQVLSPASKQESPPAAAVRPFT PQPSKDTFFPAFRKPQTVAASSIYSM YTQQQAPGKNFQAVQSALTKTQPR GPHFSSVYGKPVIAAAQNPQQHPEN IYSCSQGKPGSPEPETETVSSVHESH ENERIPRPLSPTKLLPFLSNPYRNQS DADLEALRKKLSNAPRPLKKRSSITE PEGPNGPNIQKLLYQRTTIAAMETIS VPSHPSKSPGSVTVNPESSVEIPNPY LHVEPEKEVGSLVPEPLSPEDMGSA STENSDVPAPSAGLEYVSEGVTDSSST NLQNNVEETNPEAPHLLEVYLEEYP PYPPPPYPSGEPVSEEDSARMRPPE ITGQVSLPPGKRTNLRKTGSERIAHG MRVKFNPLALLDSSLEGEFDLVQR IIEVDDPSLPNDEGITALHNAVCAG HTEIVKFLVQFGVNVNAADSDGWTP LHCAASCNNVQVCKFLVESGAAVFA MTYSDMQTAADKCEEMEEGYTQCS QFLYGVQEKMGIMNKGVIYALWDYE PQHDDPELLMKEGDCMTVIRREDEE EIEWWWARLNDKEGYVPRNLLGLY PRIKPRQRSLA	True	False	4.096	4.422	1.591	1.106	0.862	1.574	1.326