

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q6P5E3	Q6P5E3_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MSRRKQAKPQHFQSDPEVASLPRR DGDTEKGGQPSRPTKSKDAHVCGR CAEFFELSDLLLHKKSCTKNQLVLIV NESPASPAKAFPPGPSLNDPDDQMK DAANKADQEDCSDLSEPKGLDREES MEVEVPVATTTTTTGGSGGGGGST LSGVTNITTPSCHSGSSTGTSAITTSL PQLGDLTTLGNFVSNVNIENLQS TKVAVAFQSEARCGGASGGKLLIST LMEQLLALQQQIHQLQIEQIRHQI LLLASQSADLPAAPSIPSOGLTRTSA NPLTTLSSHLSQQLAVAAGLAQSLAS QSANISGVKQLPHVQLPQSSSGTSIV PPSGGTPNMSIVTAAVTPSSEKVA SNAGASHVSSPAVSASSPAFAISSL LSPASNLLPQPTPANAVFPTPLPNI ATTAEDLNSLSALAQQRKSKPPNVT AFEAKSTSDAEFFKHKCRFCAKVFG SDSALQIHLRSHTERPFKCNICGN RFSTKGNLKVHFRHKEKYPHIQM NPYPVPEHLDNVPTSTGIPYGMSIPP EKPVTSWLDTKPVLPPLTTSVGLPLP PTLPSLTPFIKTEEPAPIPIHSAASPQ GSVKS DSGAPDLATRNPSGVPEEVE GSAVPPFGGKGEESNMASAVPTAG NSTLNSPVADGGPGGTTFTNPLLPL MSEQFKAKFPFGLLDSAQASETSK LQLLVENIDKKATDPNECHHRVLS CQSALKMHYRTHTERPFKCKICGR AFTTKGNLKTHYSVHRAMPPLRVQH SCPICQKFTNAVVLQQHIRMHMG GQIPNTPVPDNYPEMESDTGSFDE KNFDDLDFSDENMEECPEGSIPDT PKSADASQDSLSSSPLLEMSSIAAL ENQMKMINAGLAEQLQASLKSVEN GSMEGDVLTDSSSVGGDMESQSA GSPAISESTSSMQALSPSNSTQEFH KSPGMEEKPQRVGPEFANGLSPTP VNGGALDLTSSHAEKIKEDSLGILF PPRDRGKFKNTACDICGKTFACQSA LDIHYRSHTKERPFICTVCNRFSTK GNLKQHMLTHQMRDLPSQLFEPSS NLGPNQNSAVIPANLSSSLIKTEVNG FVHVSPQDSKDAPTSHPQGPLSSS ATSPVLLPALPRRTPKQHYCNTCGK TFSSSALQIHERTHTGEKPFAC TIC GRAFTTKGNLKVHMGTHMWNSTP ARRGRRLSVDGPMTFLLGPNVFKFPE MFQKDLAARSGSGDPSSFVNQYTA ALSNGLAMKANEISVIQNGGPPPIG SLGSGSSSPISGLTGNVEKLGNSPEPS APLAGLEKMASSENGTNRFRTRFVE DSKEIVTS	True	False	1.41	5.0	1.657	1.065	0.76	1.743	1.893