

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
A0A0G2JYI7	A0A0G2JYI7_RAT	Rattus norvegicus	NaN	36.054081	NaN	NaN	18683930;38843836	MIVAEVLGKNHRCQOHTQHSRHNK WYSCTDLLTLELSLATAEQMRLAQ VIFDKNDSDFEAKVKQLMEVTKNQ DECIVALHDCNGDVNKAINTLLEGS SDTTSWETVGGKKKNGFGRESSENK ENREKRTEREASRGRGNRRKGRG GSRGREFKGEENGVDSCSQGDKPSE RGKRTRGRGFRGRGRGAGRLSAQ SMGTFNPADYSESMSTDGCGTKLV VWEEAQNGTDEGPEGLAKPHSISQE LPSKSSYGLKGAWKNSVEEWTAED WTEDLSETKVFTASSAPADNHVPPG QSIDLVALLHKPAPPTQAAEVNSFET SQQQGFGQALVFTNSQHNNPTAPG TASSTPASSYSPQSLSSVLGSGFGEL AQSKMVNISNSQILDKLPPLSPFP AASSTQQNDNASPPATTAAWDLKPS APQPSVLSHLDFKSQPEPSPVLSQLS QRQQQNQAVSVPPPGLESFPPSQTK HRESAPGDGPSTVNRLLQLPTMAVE NIVSAHQQPQKHILPKRRIPPASKV PVSAVEMPGSADITGLNVQFGALEF GSEPSLSEFGSTASASENSSQLPISL YPKSLSEPLNASFPMTSAVQNPTYTT SVITSTLTSSVSSTSPVTTCSYDQ SSVHTRIAYQSPASPADAAPGSVANG HGGGRSQHTVDTTSSVPAPKTDPS ALPSVSSLPSSTCTALLPSSQHTA TLPALTPGELSSPLSQLSSLSHQ SSLSTHATRSTSTPHTHASVESTPS SAAFSAATSAPSAPSSGVALPGSMST VSSLCLGGTTVSAPSSSTRATALVTS GKAPPNLPQGVPLLHNQYLVGPGG LLPAYPIGYDELQMLQSRPMDYY GIPFAAPTALASRDGNLANNPYSGD VTKFGRGDSASPAPPTTQAQQSQ SQTHHTAQQPFLNPPGYSYTG PYTGVPGAFQYGPTMFVPPPSAKQ HGVTLPPTPFQOASGYGQHAYSA GYDDLQGTAAAGDYTKGGYGGSSQA PNKSAGSGPSKGVSVSSGTGLPDMT GSVYNTQTDFDKQGFHTGPPPFSL PSALGSTGLAPGAAPGYAPAPFLHI MPAHQQPHSLLHHLPQDAPSGS GQRSQPSSLQPKSQASKPTYGNSPY WTN	True	False	1.938	4.233	0.504	False	False	False	False