	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9Z2Q1	SC31A_RAT	Rattus norvegicus	Protein transport protein Sec31A	36.506183	NaN	S526;S531;S799;T1190;S1192	18683930;34502162	MKLKEIDRTAMQAWSPAQNHPIYLA TGTSAQQLDATFSTNASLEIFELDLS DPSLDMKSCATFSSSHRYHKLIWGP HKMDSKGDVSGVLIAGGENGNIILY DPSKIIAGDKEVVIAQKDKHTGPVRA LDVNIFQTNLVASGANESEIYIWDLN NFATPMTPGAKTQPPEDISCIAWNR QVQHILASASPSGRATVWDLRKNEP IIKVSDHSNRMHCSGLAWHPDVAT QMVLASEDDRLPVVQMWDLRFASS PLRVLENHARGILAIAWSMADPELL LSCGKDAKILCSNPNTGEVLYELPTN TQWCFDIQWCPRNPAVLSAASFDGR IRVYSIMGGSIDGLRQKQVDKLSSSF GNLDPFGTGQPLPPLQIPQQTSQHSI VLPLKKPPKWIRRPVGASFSFGGKLV TFENVTGQPQQGAEQPRRQPVFISQ VVTEKDFLSRSEQLQHVVQSQGFIS YCQKKIDASQTDFEKNVWSFLKVNF EEDSRGKYLELLGYRREDLGEKIALA LNRVDGSDVALKDSDRVAQSDGEES PAEEGQLLGERIKEEKQECDFLPSA GGGTFNISVSGDIDGLITRALLTGNF ESAVDLCLHDNRMADAIILAIAGGQ ELLAQTQKKYFAKSQSKITRLITAVV MKNWKEIVESCDLKNWREALAAVL TYAKPDEFSALCDLLGARLESEGDSL LRTQACLCYICAGNVERLVACWTKA QDGSNPLSLQDLIEKVVILRKAVQLT QALDTNTVGALLAEKMSQYANLLAA QGSIAAALAFLPDNTNQPDIVQLRDR LCRAQGRSVPGQESSRSSYEGQPLP KGGPGPLAGHPQVSRVQSQQYYPQV RIAPTVTTWSDRTPTALPSHPPAACP SDTQGGNPPPPGFIMHGNVVPNSPA PLPTSPGHMHSQPPPYPQPQPYQPA QQYSLGTGGSAVYRPQQPVAPPASK RYPNAPYVSPVASYSGQPHMYTAQP ASSPTSSSAPLPPPPSSGASFQHGGP GAPPSSSAYALPPGTTGTPPAASELP ASQRTGPQNGWNDPPALNRVPKKK KLPENFMPPVPITSPIMNPGGDPQP QGLQQQPSASGPRSSHASFPQPHLA GGQPFHGIQQPLAQTGMPPSFSKPN TEGAPGAPIGNTIQHVQALPTEKITK KPIPDEHLILKTTFEDLIQRCLSSATD PQTKRKLDDASKRLECLYDKLRDQT LSPTIISGLHSIARSIETRNYSEGLTV HTHIVSTSNFSETSAFMPVLKVVLS QASKLGV	None	None	None	None	None	None	None	None	None