

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O94979	SC31A_HUMAN	Homo sapiens	Protein transport protein Sec31A	32.778005	T160;S451;T658;S666;T674;S762;T774;T903;S904;T910;S914;S915;S917;S938;S948;S963;S964;S965;S1041;S1047;S1048;S1050;S1051;T1073;T1187;T1190;S1202	S527;S532;S799;T1161;S1163	27655845;21740066;34725712;38665916;34931806;34142147;35008409;34019948;29913562;29351928;30397120;29161034;20305658;28510447;32119511;32870666;32885474;33214551;34229054;23301498;38253038;35138101	MKLKEVDRTAMQAWSPAQNHPHYL ATGTSAQQLDATPFSTNASLEIFELDL SDPSLDMKSCATFSSSHRYHKLWIG PYKMDSKGDVSGVLIAGGENGNILY DPSKIIAGDKEVIAQNDKHTGPVRA LDVNIFQTNLVASGANESEIYIWDLN NFATPMTPGAKTQPPEDISCIAWN QVQHILASASPSGRATVWDLRKNEP IIKVSDHSNRMHCSGLAWHPDVAT QMVLA SEDDRLPVIQMWDLRFASSP LRVLENHARGILAIAWSMADPELLL SCGKDAKILCSNPNTGEVLYELPTNT QWCFDIQWCPRNPAVLSAASFDGRI SVYSIMGGSTDGLRQKQVDKLSSTF GNLDPFGTGQPLPPLQIPQQAQHSI VLPLKPKPKWIRRPVGFASFSFGGKLV TFENVRMPHQGAEQQQQQHHVFI SQVVTEKEFLSRSDQLQAVQSQGF INYCQKKIDASQTEFEKNVWSFLKV NFEDDSRGKYLELLGYRKEDLGKKI ALALNKVDGANVALKDSQVAQSD GEESPAEEQLLGEHIKEEKEESEFL PSSGGTFNISVSGDIDGLITQALLTG NFESAVDCLHDNRMADAILAIAG GQELLARTQKKYFAKSQSKITRLITA VVMKNWKEIVESCDLKNWREALAA VLTyakPDEFSALCDLLGTRLENEG DSLLQTQACL CYCAGNVEKLVACW TKAQDGSHP LSLQDLIEKVILRKAV QLTQAMDTSTVGVLLAAKMSQYANL LAAQGSIAAALAF LPDNTNQP NIMQ LRDRLCRAQGE PVAGHESPKIPYEK QQLPKGRPGPVAGHHQMPRVQTQQ YYPHGENPPPPGFIMHGNVNPNA GQLPTSPGHMHTQVPPYPQPYPYQP AQPYPFGTGG SAMYRPOQPVAPPTS NAYPNTPYISSASSYTGQS QLYAAQH QASSPTSPATSFPPPPSSGASFQHG GPGAPPSSAYALPPGT TGLPAASE LPASQRTGPQNGWNDPPALNRVPK KKKMPENFMPPVPITSPIMNPLGDP QSQMLQQQPSAPVPLSSQSSFPQPH LPGGQPFHGVQQPLGQTGMPPSFS KPNIEGAPGAPIGNTFQHVQSLPTKK ITKKPIPDEHLLIKTTFEDLIQRCLSS ATDPQTKRKLDDASKRLEFLYDKLR EQT LSP TITSGLHNIARSIETRN YSEG LTMHTHIVSTSNFSETSAFM PVLKV VLTQANKLG V

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