	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	Q8N5H7	SH2D3_HUMAN	Homo sapiens	SH2 domain-containing protein 3C	34.455997	NaN	S22;T278;T283;S359;S440;T793	40914422	MTEGTKKTSKKFKFFKFKGFGSLSN LPRSFTLRRSSASISRQSHLEPDTFE ATQDDMVTVPKSPPAYARSSDMYSH MGTMPRPSIKKAQNSQAARQAQEA GPKPNLVPGGVPDPPGLEAAKEVMV KATGPLEDTPAMEPNPSAVEVDPIR KPEVPTGDVEEERPPRDVHSERAAG EPEAGSDYVKFSKEKYILDSSPEKLH KELEEELKLSSTDLRSHAWYHGRIP REVSETLVQRNGDFLIRDSLTSLGDY VLTCRWRNQALHFKINKVVVKAGES YTHIQYLFEQESFDHVPALVRYHVG SRKAVSEQSGAIIYCPVNRTFPLRYL EASYGLGQGSSKPASPVSPSGPKGS HMKRRSVTMTDGLTADKVTRSDGC PTSTSLPRPRDSIRSCALSMDQIPDL HSPMSPISESPSSPAYSTVTRVHAAP AAPSATALPASPVARRSSEPQLCPGS APKTHGESDKGPHTSPSHTLGKASP SPSLSSYSDPDSGHYCQLQPPVRGS REWAATETSSQQARSYGERLKELSE NGAPEGDWGKTFTVPIVEVTSSFNP ATFQSLLIPRDNRPLEVGLLRKVKEL LAEVDARTLARHVTKVDCLVARILGV TKEMQTLMGVRWGMELLTLPHGR QLRLDLLERFHTMSIMLAVDILGCT GSAEERAALLHKTIQLAAELRGTMG NMFSFAAVMGALDMAQISRLEQTW VTLRQRHTEGAILYEKKLKPFLKSLN EGKEGPPLSNTTFPHVLPLITLLECD SAPPEGPEPWGSTEHGVEVVLAHLE AARTVAHHGGLYHTNAEVKLQGFQ ARPELLEVFSTEFQMRLLWGSQGAS SSQARRYEKFDKVLTALSHKLEPAV RSSEL	True	False	4.711	2.64	3.068	2.411	2.007	3.908	1.9