

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q06033	ITIH3_HUMAN	Homo sapiens	Inter-alpha-trypsin inhibitor heavy chain H3	25.476563	T847	NaN	36240223	MAFAWWPCLILALLSSLAASGFPRS PFRLLGKRSLPEGVANIEVYSTKIN SKVTSRFAHNVVTRAVNRADTAK EVSFDELPKTAFITNFTLTIDGVTYP GNVKEKEVAKKQYEKAVSQKTAGL VKASGRKLEKFTVSVNVAAGSKVTF ELTYEELLKRHKGKYEMYLKVQPKQ LVKHFIEVDIFEPQGISMLDAEASFI TNDLLGSALTKSFSGKKGHVSKFSP LDQQRSCPTCTDSSLNGDFTITYDV NRESQGNVQIVNGYFVHFFAPQGLP VVPKNVAFVIDISGSMAGRKLEQTK EALLRILEDMEEDYLNFIKSGDVS TWKEHLVQATPENLQEARFVKSM EDKGMTNINDGLLRGISMLNKAREE HRIPERSTSIVIMLTDGDANVGERP EKIQENVRNAIGGKFPYLNLFVGN LNYNFLENMALENHGFARRIYEDSD ADLQIQGFYEEVANPLLTGVEMEYP ENAILDLTQNTYOHFYDGEIVVAGR LVDEDMNSFKADVKGHATNDLTF TEEVDMKEMEKALQERDYIFGNIE RLWAYLTIEQLLEKRNHAGEEKEN LTARALDLSKYHFVTPVPLTSMVVT EDNEDERAIAADKPGDAEATPVSPA MSYLSYQPPQNPYYYVDGDPHFIIQ IPEKDDALCFNIDEAPGTVLRLIQDA VTGLTVNGQITGDKRGSPPSKTRKT YFGKLGIANAQMDFQVEVTEKITL WNRVPSTFSWLDVTVTQDGLSM MINRKNMVSFSGDGVTFVVVHQQ WKKHPVHRDFLGFYVVDVSHRMSAQ THGLLGQFFQPFDFKVS DIRPGSDP TKPDATLVVKNHQLIVTRGSQKDYR KDasIGTKVVCWFVHNNGEGLIDGV HTDIYVPLNF	False	True	1.441	1.483	1.365	1.995	0.75	1.661	5.0