	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	Q9NVR5	KTU_HUMAN	Homo sapiens	Protein kintoun	18.419428	NaN	S461;S467;S640;S641;S773	30379171;23301498	MAKAAASSSLEDLDLSGEEVQRLTS AFQDPEFRRMFSQYAEELTDPENRR RYEAEITALERERGVEVRFVHPEPG HVLRTSLDGARRCFVNVCSNALVGA PSSRPGSGGDRGAAPGSHWSLPYSL APGREYAGRSSSRYMVYDVVFHPDA LALARRHEGFRQMLDATALEAVEKQ FGVKLDRRNAKTLKAKYKGTPEAAV LRTPLPGVIPARPDGEPKGPLPDFPY PYQYPAAPGPRAPSPPEAALQPAPTE PRYSVVQRHHVDLQDYRCSRDSAPS PVPHELVITIELPLLRSAEQAALEVTR KLLCLDSRKPDYRLRLSLPYPVDDGR GKAQFNKARRQLVVTLPVVLPAARR EPAVAVAAAAPEESADRSGTDGQAC ASAREGEAGPARSRAEDGGHDTCVA GAAGSGVTTLGDPEVAPPPAAAGEE RVPKPGEQDLSRHAGSPPGSVEEPS PGGENSPGGGGSPCLSSRSLAWGSS AGRESARGDSSVETREESEGTGGQR SACAMGGPGTKSGEPLCPPLLCNQD KETLTLLIQVPRIQPQSLQGDLNPLW YKLRFSAQDLVYSFFLQFAPENKLST TEPVISISSNNAVIELAKSPESHGHW REWYYGVNNDSLEERLFVNEENVN EFLEEVLSSPFKQSMSLTPPLIEVLQ VTDNKIQINAKLQECSNSDQLQGKE ERVNEESHLTEKEYIEHCNTPTTDS DSSIAVKALQIDSFGLVTCFQQESLD VSQMILGKSQQPESKMQSEFIKEKS ATCSNEEKDNLNESVITEEKETDGD HLSSLLNKTTVHNIPGFDSIKETNM QDGSVQVIKDHVTNCAFSFQNSLLY DLD	True	False	4.508	3.13	1.414	1.384	0.988	1.407	2.842