	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	Q8NDW8	TT21A_HUMAN	Homo sapiens	Tetratricopeptide repeat protein 21A	25.476563	T156;T162	NaN	36240223	MSSNDSSLMAGIIYYSQEKYFHHVQ QAAAVGLEKFSNDPVLKFFKAYGVL KEEHIQDAISDLESIRHHPDVSLCST MALIYAHKRCEIIDREAIQELEYSLKE IRKTVSGTALYYAGLFLWLIGRHDKA KEYIDRMLKISRGFREAYVLRGWVD LTSDKPHTAKKAIEYLEQGIQDTKDV LGLMGKAMYFMMQQNYSEALEVV NQITVTSGSFLPALVLKMQLFLARQ DWEQTVEMGHRILEKDESNIDACQI LTVHELAREGNMTTVSSLKTQKATN HVRNLIKALETREPENPSLHLKKIIV VSRLCGSHQVILGLVCSFIERTFMAT PSYVHVATELGYLFILKNQVKEALLW YSEAMKLDKDGMAGLTGIILCHILEG HLEEAEYRLEFLKEVQKSLGKSEVLI FLQALLMSRKHKGEEETTALLKEAV ELHFSSMQGIPLGSEYFEKLDPYFLV CIAKEYLLFCPKQPRLPGQIVSPLLK QVAVILNPVVKAAPALIDPLYLMAQV RYYSELENAQSILQRCLELDPASVDA HLLMCQIYLAQGNFGMCFHCLELG VSHNFQVRDHPLYHLIKARALNKAG DYPEAIKTLKMVIKLPALKKEEGRKF LRPSVQPSQRASILLELVEALRLNGE LHEATKVMQDTINEFGGTPEENRITI ANVDLVLSKGNVDVALNMLRNILPK QSCYMEAREKMANIYLQTLRDRRLY IRCYRELCEHLPGPHTSLLLGDALM SILEPEKALEVYDEAYRQNPHDASLA SRIGHAYVKAHQYTEAIEYYEAAQKI NGQDFLCCDLGKLLLKLKKVNKAEK VLKQALEHDIVQDIPSMMNDVKCLL LLAKVYKSHKKEAVIETLNKALDLQS RILKRVPLEQPEMIPSQKQLAASICIQ FAEHYLAEKEYDKAVQSYKDVFSYLP TDNKVMLELAQLYLLQGHLDLCEQ HCAILLQTEQNHETASVLMADLMFR KQKHEAAINLYHQVLEKAPDNFLVL HKLIDLLRRSGKLEDIPAFFELAKKV SSRVPLEPGFNYCRGIYCWHIGQPN EALKFLNKARKDSTWGQSAIYHMV QICLNPDNEVVGGEAFENQGAESNY MEKKELEQQGVSTAEKLLREFYPHS DSSQTQLRLLQGLCRLATREKANME AALGSFIQIAQAEKDSVPALLALAQA YVFLKQIPKARMQLKRLAKTPWVLS EAEDLEKSWLLLADIYCQGSKFDLA LELLRRCVQYNKSCYKAYEYMGFIM EKEQSYKDAVTNYKLAWKYSHHAN PAIGFKLAFNYLKDKKFVEAIEICND VLREHPDYPKIREEILEKARRSLRP	None	None	None	None	None	None	None	None	None