

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q4DM53	Q4DM53_TRYCC	Trypanosoma cruzi	TOG domain-containing protein	19.946178	NaN	NaN	30984116	MDRAHILRLLDNLRSDNTLRKSAE AEYESIIQGNVWMMCNLSELCAVT DSAPTMQMGLVLLKFLFSSKHNCF DVSDAQTQQAQVKGKGLMSQVLGKAAF GPQRGLAAACVSALVVKMHALGQE WGELWQSVFQILENAESDHQLKTIC CEIIATTGPSMASYFESHTGRLVTGI RNCLADPSVEARKSAFDALVNVAMC RSIPDFAQLVPLMLQVVQDSLNASN WDDAEQLTGKLADGVAHAPGLFAG HTSAVLHGLMEVASAPSVASGARH MAIETLLTYCESEPKTVRKVPNFSTS FLQLLFEYTLNPMVMPDDWDIKGVNL DDDLEEDDDDTVGSSGIDRLASSLG GRKLETVAOQLFVENIHSSDWKRRN AALLLITYLAEGMTTVLEKHLEQIVQ MVIPAVRDEVKYVRASALDCLTQMS SDFAPKMQEQLCHSVVPMVGCGLG DSVPAVATRAARCLDSFFDQCEESE NEDDTVFIKQFENYIEGLCVSLVTL KQTSHKFVREDCLGALSSVISTCKGL LKPFPVSHLVPVFQEV LAMPETPETI MMKCKAIECTLLACGVRGSEFAPY AHEMCNYLRDLLNHLARGSKEDDM RLRYVLRGWTMTDCLREEVTPYLA IVMPVLISMNVECDTEVENAEVGD DEEDEEEKDVTMRVVVPGVGVVKI KMHTGLIEEKDLAASVVSAMLSYVG KQLKPHLPQITESAVKLLSFQSDSSI RESGALIIDGVMDAYETAERAQVAVS VMSPLLNQFAEEDDLEASSAMSVVI SRCIDDAPTLVSIETVDSISEKILGVL RRAMESRTESLQSQEENDDDELD KLKEEEEEAEALIRDTCDLLDKMLE RAGAVFAPVFNKFIPLVLRMLQGN EKEFMVARGLALLCSLVEHAPDHVA GLISTIVQSVINFAQRHEDADVLSQSS FYLMNLLLQYFERHEYP AIRQFVQQ VYGIFSRYMAVAHKEEYEDTTCNAIS MAVTLLSLYYQLLPEHELAQSLDCV VNSLPAGGDKTEACRVHERVMMWV VQRHSLQGNEARVKAIVARLKS AKEDVTNESTRAQLAHM	None	None	None	None	None	None	None		