	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	K2MNF2	K2MNF2_TRYCR	Trypanosoma cruzi marinkellei	NaN	19.946178	NaN	NaN	30984116	MPDTLFIKGEDGGPELQTRFDHVFD RDAAQSDVFDIIGSEVLNTLFSGYNA SVFAYGQTGSGKTYTMEGIRSKPEG LGLTPRLIGALFERFKANKDITDSVC EVSLVQIYQEKIQDLLAGQRQLEIHM DRTGQYIARDATWARVKSLEESMKL YTKASEMRATSATEMNLVSSRSHMI MMVKLQWDEPSLPGSHAQLNLIDL AGSERLAQSGAKGDSMWEAIHINKS LSALGTVVSKLVEQAKHKGRRIHVPY KDSKLTYLLQSSLGGSNPIHFILAISC SALWSGETKSTIEFGKRALQLVLRPV RNAIDYTRLAEMEEMIERMRSHIAS LEEELRNKRSNEAAEFLKLKQIPQQ GDENNDRFAERRHQSRQKKKLKVQ TELARIMANLPETFDDLTSHCVLFPE SKASFRELGGLQRLVYFVDRSASTFY RSNAAQTIASVLDDAGREMFQDIGG LDALTRLLQVKEERCKEAACVALEA ACRGCLKNKMKLTPVVYTELVELIY NYPNQQVQEAACTAVAAIVDLYPDA RRCFERLEVVPKLLETIRNAPEEVVN LTKAATNCVGRLAHGDSGMQQIIAS FGGIDLLIDVLFSPAGTRDHQVPILAS YALVNLCCSNKHNLDIARDNLRYGE VKFRLLEGLARAFGINAAREGYGRAT AQETESPFPYYGVTIVDKWSVTSSGG RPIFSTFMDNPQFYLYVTETTDIAFM IQDVLYESRMLKKKKNNTVYMGLAL FEGDPELSKAGLKQLDFHGRMVEIG KYTSNCENVLHCTLQPSETPYVVVP FTSQRGRQTEFALSAFGDKPIQLTAV PEQVGWVRTVLEGSWTELTGRGGS GFDWRCNSQIRLQPKENCRVVFVLS YLSVDRQRAQSREEEGEEQNMRPR LHGRLFTTFAPEKRYLKALVPLPQKS TFVASNNFASNSYITTSANLKAGESY VYIPFTETPYEDQWRLSVYCDTDDVT IAPIEGNKAEWYCTSSAGTWAGKPIS VQVETKGKMVAVASSPGVFLRIRLL NAKGKKLEGIDAYWNAEASVEYKSQ GVVIVQVEGMMRTDKGQEAAREMK FDVFVFTENKCTVQHVDFAAVPSTL PHPTKSTPADLLQYPVEVVDADVPF DTIDQSDDEEEEHNDVGNTEEHER ELQKAITLRNEENARLMTRIVEQQR EIMQLRQVHDSTSPSNSVGKSPANG KRQKKSRSITPLEPLPSMRKQTLKKP DGGVSAKGDAEETAAGVETWGGPE DSRDVSGAPAKALQEAVEYSLLTLN SIERHTRAPTEAEWADMQEDLRELQ RRLFFALEKISLT	None	None	None	None	None	None	None	None	None