	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	Q4CZF0	Q4CZF0_TRYCC	Trypanosoma cruzi	Glycine cleavage system P protein	19.946178	NaN	NaN	30984116	MYRCFHHFSTLPLAFVRRTATNSYV NRHIGPSAAESISMLRIIGKESLSDL MTAVIPENILRSPLKEFPAMSEQDAL SLLHSLGSRNKVLKSMIGQGYYESIT PPVILRNVIENPAWYTPYTPYQAEISQ GRLESLLNFQSMIIDLTAMNLANAS LLDQATACAEAMYLAFHHGRKERM TFFVSRDVFPSCVEMAKTRAEPLKIK VFVGDPNLIDWSDSSLCGILVQTPDA MGMLHDFTTLFEKAKQHGVVSCCG TDLMASVLLKPPGEMGADVVLGSAQ RFGAPLGFGGPHAAFFAVKEEFKRLI PGRVIGISKDSTGCPAIRMALQTREQ HIKKERATSNICTSQALLANVAAFYA IYHGSEGLKEIASEMHNKAKILSVGL ESVGHTVVNGTFFDTITVNLKGITPE DYVTCCVEKGINIFVDYSHGTVSISV DEATTEGHVVSLLEAAGLKLPVIGVL SKLAEQKRAMPLQMLRKSVFLGHSI FQKYKSESELMRYIHRLHRKDYGLT HGCVPLGSCTMKLNPAAAMLSLSW PEFTNIHPLAPKEQTRGYSALCLDLE QKIRDITALDAVSLQPNSGAQGEYAG LRVIRSYHNSKKESHRNICLIPESAH GTNFASALLAGMVIVKIKCLADGRID MKDLENSCQKHTKNLSCIMITYPST YGLFDREILAITSMVHYHGGQCYIDG ANMNAMVGYTAPGCIGGDVCQVNL HKTFSIPHGGGGPGMGPIAVRQHLA SFLPDSVFIQNVGGSQPFGQVSQAA YGSASILPVSYLLMLMLGSRGLKTCT EYAILNANYLKKRLDGHYPVLFLGE NDFCAHEFIIDLRPFKKTAQIEAEDV AKRLMDYGLHSPTLAFPVAGTLMIE PTESESKRELDRLADALISIRTEIASIE KGEESTTNNVLKNAPHTAKCVTSDD WDRPYTRKTAAFPSSHSYTEKFWPS VGRIDGTYGDRNLMCSCALTNFCE	None	None	None	None	None	None	None	None	None