	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	Q4D6Z3	Q4D6Z3_TRYCC	Trypanosoma cruzi	Guanine nucleotide-binding protein subunit beta-like protein	19.946178	NaN	NaN	30984116	MSSRRARLSPRSQPQQSSARSSRAG TPSRCPCSPPTSATKRGKKNHRVSS VGPQKFRSRALDVTPLFTVEGAVRYI DVINERTAWTVECDGSISVRSIPKGN EMKQISGRKGVFCTCLLFVGKTGKV WAAFSDGFLQVYDVNTFLLENTFVQ HTGAVECIVEMEDRIYTGGRDWKIY QWIPEMYCYERQFAGHGNTVRCLC PYTGSTGAVLFTGSDDGTVKAWNPY PSVKQSKENNPCVHTFVGHTRGVLS LEIVAPNNQLWSGGEDTTIRVWDLQ TLDCVAVLEEHNSPIACLSVVEHRV WSGDKHGRILLWDLRTLSPLQELSR QMRVEHRSVFAIRKMLQTVVWKVW TTGSAGHIHCWNAESMPLVFDSDTP GEMLPHTDEENMRALEVDNEALRK EVARLTRQLEKESAHAISEKFNVMQ NQQLIMDTNDGLRRQLRETQRLGA VGNGGNTEHGIRSPYTELGYPQAPR LDLSMPFSDASRNGGEIGGNERGEA PFWEPITMSIYRNDVTPIITKIEQFFP GSAWSEILQDYENELRQTFIREAAQ ALGVPEEKLQHVSMEPNPQGLFATV KVPHSSDTSAEELKRPLKAHPFVPVL NLYGAALANKNERLKEALIQAHNEN QSASRSMQDNPKTKTDGPSAIMEK NEKIAHLSNELDSVKRQNVRLQKEL KKINEQLAAREAPNRINAQEEYDAL KSFVDNSLKPLISRLKRTNEEKELDL RANEERIRQLLHDKEQLQKQLATRE APSRINAQEEYDALKSFVDNSLKPLI SRLKRTNEEKELDLRANEERIRQLL HDKEQLQKQLATREAPSRINAQEEY DALKSFVDNSLKPLISRLERTNEEKE LDLRANEERIRQLLHDKEQLQKQLA AREAPSRINAQEEYDALKSFVDNSL KPLISRLERTNEEKELDLRANEERIR QLLHDKEQLQKQLAAREAPSRINAQ EEYDALKSFVDNSLKPLISRLKRTNG EKELDLRANEERIRQLLHDKEQLQK QLAAREAPSRINAQEEYDALKSFVD NSLKPLISRLERTNGEKELDLRANE EHIRQLLSKCGRSWSLCALNTGAFFI HSEGMESDRHVLLDALVEAEESARV RESDLLALVDSLQNVVDRYVVEKEK LVARLESLNGELSEALKERDMAMS DGKLLEEECAALRDQLRGELLAQIPV GEQKLLAAYLDEVQRRRELEKEMYR GLSKRVKASW	None	None	None	None	None	None	None	None	None