

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
Q4DPB6	Q4DPB6_TRYCC	Trypanosoma cruzi	NaN	19.946178	NaN	NaN	30984116	MPTVLSPASRQAYAVVLLALMVSLLFISAADAEIIVIGVLDGSHKLTTEDIKNGFKENDKNFINSCGKFDVHGDKFVLIEMGKFADYLIPOKGVHLCDLLTSP TREQKYLAPQDPRDATWDGHTFPVTPSIYEDALGSSAEGYVNAERSTLSF WGDNGSSDNHTGGCCALMPGQNA AWYRYPYRLLFSPKEPAMYLAWGTKP TVAVPNVPVFEVNLWNMRSEPVLENVTFNITNGKKVTVTSPTGRYTHF SNASDPSAAKYTHIVDLVSGPDTVNT RQLKAVVDVFGKLPWDVPTAGGNI PALVRFKGAGTAIGPVDGIAALDEWD FASSIVGGESRIRPVAGHLNTPVEPYT LPPVSDKWENVTQADGVFRVQVPCK SVQYYAVAVYSAVQRIVLQYEFDTA AVMYFAGKRFPVPREEGSRRGSFAV DVMQGYHQVLVKLFSNSTATDP RKCTAVSSFSMRVVGTA VGYTHAVM PDIPEGAYEIWNNNYMFYLLHLGEQ REEYLYGALDEDFINVSTAIPKPEH MSGKVWKVVFYGGGRIPVSAESLPK DKEAAVSYVAIALYSGYEEKVPVGWTF TTGGKTD FYIDGKPVYSRPSDIGPL NESFTTPVSRGWHQLIVRIAATKGQ SWGLTFYIHF DNYRLGSAHITGSNLP FGVAPIRPNRPLLSLMELVDVKSRY GAIPFNPFNNTAVADEFTDLPIDASY MPGLLLNQSIWTLGQERQYQWVGR VSADGIWGHPRPFGGNSNOYWFVVL YATESMEATARVIFHGGYAMWLDG QLVGSSPLSSEKEVRETLPLNKGWN QLVMKLRANNSFHLIDTRNSRMSW TAANALCAETRHMELCPLSAVC PFG VGRPGVFPVPEGSSFVPVKNDENEW VQVFSGGGNGTSMCETWRDAHDG RAPIWGETSDNITERHLVPCCGDNS PPFIWLSITPSSTAEGELGYTYEIPVV VEAPVTVDDPAAVTQTVTITCATPG AIKYAVNNAEEMALYTHPFTISESS TIFAQAFHSRESRRLEVFVEIKS	None	None	None	None	None	None	None	None	

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