

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
V5BB69	V5BB69_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MYLRICNNEGGMKSVSSLSQLHNE AEKLFGRRLFLAVDEEFVCSLHCFP PTETQRAGGSVFSVLYRDLVQRTCF RKSSVSTGIPRLLFSYVAEAVLCSR GEFMGPLCSPLRKSHTAMLRSNQY FLTEKSDGVRVILVSVFCVHAFPCWV LRDAEGPTCNILRLDDVAALEEMRQ QFFQGATSQSNRSRLQISIGSFYIEA EGSKTYNCEMETFTLTKSCPESNSGV FSNGNIIMAERTQGTRHMAYCFDRS MEQAYLLEEYAAPTLSAFVAVDGE MLALRDLKTRERTRPLYSCFDVFRYTP MKDAVGTGVCLTRWPMSKRYDVLR DHIIASIHASENLRTQKPLLHVFAKE MFPVKNFGEVCVARLRRADTSGNTG GVMYYDGPWGWTLSDFGIFTPEKF DLVHGASKEQLKWKWPSMLSDVWS LAAVEGRTDEYVVDLFFRKRFRGHR PDSVGRSRLSSAMRLNPFSLRPTG RSVVAECVDFDRHKSWSIERLRITDK SEPNSVVTVISVLESLENITLDVLV GMLGVEDTVSNRDLKSLQEATDLEE EGKIVSAPESPIEKKTCQLTLRATQS QMQGEHEIHLYWTVRLLSEKQHIPC IHCKVSECSGLGFACPMDDERSKLT ENLYVELANAGGSYAWSDFTVSALF DGDTGRWNIVLSHSGDNRKSTSV GVIEHLQWLLVHGGERVLDVMPVV NTKKPSGMMGEPMGELVEKINSHY ACKTRELLSGKNRSVLRKFNNWVK NVLISNAVSYAQRSRDKDSECGELA VADVCCGRGGDLFKWRAHNPRYLF MVDSCLEAVAEAAAGRYSVSKGLSLRI VPQDKSSIGVIAYFTVCDVFDEAGAF TVKLDEFFNQHLKGRRLDIVSCQFS LHYGCSTEKRMSCFLRAISKALRPG GVFFGTTVSDVELLRQVAEHGPPVFG NSLYSVRFPSSETPPDPSFGVQYFISV EESVSELPEYLPWNRLVALCSSVG LLPVESLGFMEYSDMHYNSPIGQKL RDAVLRGGKRDSGDGHALLQLSSEER EAASLFRITFFFVKNLVENEACFVNN	None	None	None	None	None	None	None		