

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
V5B755	V5B755_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MDVFLTSAREADDMKLVNFVAMHD AIAAARTSEEGRELLAALMDAYYAAD RPHAVYQPALMKAIGCCLRHAGPSS SDNDSMNRWQLTADDYILLALFFN ELSTVDDNKLFCIFTEAVTLFFQQT PIVYGAVGYDLSDASTVESLCSLEQ TLRVMMAKRLAAPPADSSTCEGSVG MQLGIIDCLRTLLLSLMAAATRADT KDTLLEAPGKEGENVNLAVRQAVEG ILSPLRTAVDAVRSSASGGSSSEKASA SVRLHAGTTVSPVLVLSALRVLEEIV WRCSCHVTPHDVLEKTVAEETPLIAA ASLAVTRVLGALQQYL VQQQHS ESAAGSIEKSSLAVDVLRTAGVQWA VRRLLSVVRLHLHLFSPMGNAVVA LNAFGGNMEMEDPFQLMRMRREC ETALNDENNNDATNNNGSGSSSN GSIKDVAVRWKAWEVAAMSSSSSDA FSLRLRECEQILEAPHEETTANDGVT AELGDLDDTTLLQGDGGGGREAL LQSTVGPVSVNALVEMVLLSVRHM DIMTEQAIQALHTOGLERMOYEA RAQREIILRORRIEQQGVGIPVGR LEHLKESTSLYAKGTHLLESESALA MVLRSAFFSLLDAYDAFLEETESR EAQALIALALVQLPPSMIDAADVAV VRLRKEFSRRAHDKKAPLPLYSAYQ AMQVLFSLQAPVLERGTTAFAL TATNSMTEAELQLPDSMMLRSHAA FTIDTENPVWELQSTTHDAAGLNN RKRLRDEETTQMKDEETGEEGVE NGVSHFFREDTTQSPCMYSYFLCRV VELVTEAEQPALLLDVLLQPCPCITRY VWHYIHKQYCLSTEKARCLLGMWL LKSLALKRTVYRRYAVNSLLHLAAAR NEYQRRLLATKLGELLSAKGADGRR VIDTNAETLLVRYAKRQMAAIPVTKL PSSTTSIAAAAANPVAKMEGANTSHD EGGTASLRDKEMRKLADALDRHLG PFLMLCARQPKELFPALLAVFKECV ERQNEAMMHLLAEHVDVRRMCQR LFRMDAMSFMSNVMPYLRRHSSDA TVLVQKILWAVCSSELRAMGQATESS ASQLEQTALALLGHARVMYENSEIP QMYHPSSSAGGLGDATVPLHDVRFI SSFFSLISAEEELRRTYLRSLHFVQL QLQFQLQQQQQLGSDAGMNATAEV TRMNSEELDAFIRDVVKEVLVRSVP HFSDGIARGLSRDLDLVYLSHSHES NAADSQVAAGKNSTLSASAGREGLR GILGAGLSSTAATEEAAGTKPIAEYLT ISPFTTKRVIALLKLTTRTFDGTTFE LYGPEEIKSAVRQVMQQLGGVGGSG SGRSGIISNVSSVPPQLMVTLLILACKI HAQRPHADLVRFVHHVLLPLAKDA TWEKDTKLWRGVLLFAEEHYRECS FLVNLPEKVLIALRSQPQLREYFRE EHGNNAAFGHILGSL	None	None	None	None	None	None	None		