

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
V5BRU5	V5BRU5_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MVITRSQSRKEARSLICTSNGKKDD APGVVGGDDNQNSTQERLGRGCRIPR EPPASYDSMYVRKALLPSVMRRAEN EELQINSRYQPORDAAKRAVLRMSST LHSLGERTRRNFWPLEDNNGDGDD ESSNDGSEDGNASFQSLGRKYNGT RFTNLPKRKRVREPSINSFITDDSLRE GSNGRGKDADEELLVEDPSSLSSAE GCISEGEGRRVFRKAKESTVKHIPIN HYLDEVMKASERKTKKGDQKQKGT HRRRMEEMFLGLNEAHGGPVGAGK GGEGRSITETPAPLGDITPLHIDTSVS FEKVGGLPGHIVMLREMVLFPLLYP HLLQTMNLTTPRGVLFVGPPTGKT LMARALANEAASCAQRKIAFFMRKG ADILSKVWGESERQLSLLFEEARRR QPSIIFDEIDGLVPRHAKAEQSQA ALVSTLLALIDGLDRGQVIVIGATN RPDTIDPALRRPRLDRELYFPLPDA AARRHILSIVTQPILPLDHPDRNAIM EELATRCGEWSGAEIQALCTEAGLN RLRTSLPQLYVTSRRLQIPNALTIO KEDFFVAAQRFKTSVRRGTESIVKGL EEHLEHLLYSTRMKLVSHVASMWP SAATALANERRDCSNASDAVQELIG FPIPHTVKPMLLFLTSSVGRNVYGV SYEFALEKAALAFIKALPSIRVFTLNL PNLVVDRGSAPMMVTATGSAVPEG GAGNFEELGFPLNNSFTSGHIYSCI MAIQRSSGPSLVLLQGLDLWLAENG CPITNEEGNEGDGDEERAQHKKQ MESFCYYLALLRSDVLIVAPTLHCD VLSRFFGGLRLNTQIHKTVGNVPCA PSVDDLQRFLQYVFHVVELALLHCT HTVWEELHEDISPPPPSMEEVRAA RHGTLELWRRVEYRRRQLRYVLSK WLGQYLVSRKFSILCSADLDFTPSH DLYDEWRQHTQGSRIGLCDVMEKL ENEETSLSQYHDDIDMLVRNVRTF FRTRSSTDQYRSRALELKETTIVLNL YKINRNVVQFCEQYKDLAEPDVLSL SLSPNDDVKPRMEEPTESRAVQSTV KGRKPYRYYCRRRRRRRKRKTPK VRENDNLHKIEDENEETTREGRS ANRDELKTEPAAAE'TLRDEEEVDD HAVIVERNGINSSDFSSSIAQCCKAA EIRVSMEMQLGLSFLLELHGVFOAA MSLLEEVSRRRRNQQQTVSVSPFTG TSEVGSTSFFLWMLLEAVRAAVKER RTSD	None	None	None	None	None	None	None		

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