

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
V5BY63	V5BY63_TRYCR	Trypanosoma cruzi Dm28c	PPR long domain-containing protein	19.946178	NaN	NaN	30984116	MRRGGNILQTVAGRYAVWSQLRFM SSTPCQLQGSRVGHYRPPSSEDEVQ LAAVLTTKPRHVARYVRQSMHLSRR RITQFRNSVTFLMIALQSKLQKGEIA PDDALALAEMLMRECVDLROGDMA HLLFRASIRFRRYGMKIGFQLVKHLF DSYRTDNAKDLMKNMADDELERGEES LKMLAVVAYQFAGEYDEARELLQEIP QERLVTSDYTALIEAYGMTSRIDEIM KLVRLQFSTHKTNOGSLDLDAILSTG LVAIRGDNNGMEQLKEMALDHRVK LSEKAIGAIMRSRLTSAKSAAEVYNL EEALRHELAVQTLGMAAETAVIARC SELLSRTQKSGDDVMLRKVOHLESV VEECVVNDTVDDIEPSYLLSLIKGYG VLGRFEDMNRCFDSLKEAGVVRDH RLYDEMLRWYAHAYNLKEVIALKEE MOEKQIFHTAQTYQHVFRLDKYYP RMVEKYMNMERNKGIQIESFMYP LLRVFGELODFHMVEQLYREIKSKV AMGNTNVFSAVVVQLLKSQNDM ERCNSIIHEAENYGLLANELVQAEIV QFYSVHNRYDYLHALVSRLPYKSPD MYRVLLRDAAKRKDRRTFNALLRE MQENQVWINERIFGVVVLALGYFND LDGVKRYFHEALMNNAVRTPPLFFAL AASAFARLGDQAVDECWRDLVESK VTITMPVYNKFLDLYMANNVVEKVQ QILNTMMKLVPPNPVTATTVDMLG KMGRLEEMESVLEMSRSTNAVPT QVTYHQAMNAYAKTGDVAKMEAMR AKMKEEGFQENHVTFNLFEGYGRA KRYEHIKELVEERRSKQIPMEEFY MGLLNYSRARMVEETS KLVEEMVA SGVPFTSRMLS AVASSFSHIGDIPKM EHYISLLL SHPECRHRD VESIYLIYSK MRDITVKLQELLDTENLPKTQFIYNIC VAAFARSGEHTKVATLLTQMEEEKGF FLTRNTSITLSSLLKAGKLELAQTV LKWKGCAPHRGEEGDELAADESQ LLRADNGVSSSSSSSLIPETTREAS AVVAVGGGVSAS	None	None	None	None	None	None	None		

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