	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	K2M9C4	K2M9C4_TRYCR	Trypanosoma cruzi marinkellei	NaN	19.946178	NaN	NaN	30984116	MYTLNSTSMRRSQHLLHKQHEEKP LRQVAQASSPHGGIDGTRGLNIPSFV PEFICGNPSPVPLSTAHSTGVDIKWA PSPVAPKPPPPVGMNTYGVADPSGN SWETTKKAGTNAVAPGVLLRQQRVQ EELAQEKLRVQQEIALQQQRLKQQE LSLLHLSQQQGDERGLIQQLAKLKQ QPQQQPRTHDPVSSLHHVSRPQFN QQQQQQQPAQLHQQQRQQQSIMM QEGILPLGEQWVDSKAVAHSLPQPL QQQGGVRRGNRDSGPRRQQQIYKH QSMGRKWEPISEADFGHRTGSGDN RKDSPSVVNGAGTVHDILPSLSDEEI VHKFKSLYALQTDDEGGAEHLRAQA STALGVVGTLDVEKCRKLIRDHDVTI VVTDTGTGKSTLIPKAILDGQPDAKV VNAQPRRTPAIKLAQRVAALYGEKV GRKVGYWVRGEHVGTLGETPIMYVT NYTLFLHLIHNSPNDIGITHVIFDEF HERTTEVEVLLLMMKLVMLRNPGK LRLVLCSATAEVSKWNCFFENLSVA EYSKANAMYPIHDYYLEDVSKLIGM TFVEPSTCPTGVVQSIQLHTIILMLKE MLKFLANTAAPQHSILVFVPGRTVV EQISLWIRNNLGDQLDPIPWYRDIEL GFIQEALQRESVTKKKVYVATDIAEV SLTLPDVVFVIDSGTAKRPHIIETNPN SVAFPPLELVWESSVNMLQRRGRIG RVQQGFFFTLLSRKQAQQLSPNESR LGNAVLHEIVLHCLHLTPSPFIMFSL CREKPRKVSVELSLSTLCDGGYIVPE GDCGSLIEAIDNTIHHRVKDAWSALI ADAYASRKDDNDVLGLPLAQHRFKL SLRGLIVGGLPLGVEPGTIVFSGLIM GLTTLSIIAASCVSCNTPFYVPYNATD KLQRLHLLNEVEKFMRIFKGEYYND ALSAVGAVIAYMAMQREGISEEGQN AWCEERYLSRARITDILLVAKQTREQ LAAIVPFEDVTDVDELMEQYNENAL TLSFICSAAHMHHGIYVLFDAGVAQ KKGYAGSGVFVNINCCRDRSVPTTC PWERDTVCIPFTVQTVYGRLLGSFSS QLKPEVFNIMLLLLSSRFVYEEIHED AESVTFEVAQSGSCVYINSDRITALQ LLQLRRLMCARLCVLHFLMQHEAA PSDNDAIAEHLLRTGSDFGIPPGME QQPHLLSFMITSSLTKVVQRIGATIA TNCDRRVSGVSANSQPFCRVTERPA HFHPTRQSALLFSSNGCAPYECPAG AMVPVYRDPQANLIA	None	None	None	None	None	None	None	None	None