	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	K2MNZ2	K2MNZ2_TRYCR	Trypanosoma cruzi marinkellei	Calcium-transporting ATPase	19.946178	NaN	NaN	30984116	MAPFSLPANPSTMDASAVTKSLCVD AKRGLSADEVEERRRQFGSNELPTK PSTPFWKLILAQFEDTLVRILLFAAM TSFVMALFEKNAGDFVEPFIILLILVL NATVGVWQENRAESAIEALKSFVPK TAVVLREGKLVTVGAEDLVPGDIVEV SVGNRVPADMRVLELHSTTLRADQA ILNGESVEAIKEADAVTGHHDRFPSS MVYSGTSIVYGKAQCVVVRTGALTEI GSIERDVREQEEVKTPLQIKLDEFG MLLSKVIGYICLAVFAINMVRWYSV HIPTPDEPWYERFIAPAIHCLKVAIAL AVAAIPEGLPAVVTTCLALGTRRMAR HNALVRDLPSVETLGRCTVICSDKT GTLTTNMMSVMEIFTLGVDGKLKEY ELKDSRFNVMPNVVTCEGKPVTSAL ETDGALSMLSNIAVLCNDASLHYNT AKGQVEKIGEATEAALLVMSEKLAH ATDPTAICVFRKLAEEKWKKNITLEF TRQRKSMSVHATSTAGGKLNSLFVK GAPEEVLRRSTHVMQVDGVVIPLND ALRSRIMAKLDAMSGSEHALRCIGF AFKSTQPVRELKLSDPSTFEKIESDL TFVGACGMLDPPRAEVREAIDNCRT AGIRVVVITGDRKETAEAICRKLGLLL KTETSGLSYTGAEFDAMNPTEKRKA VMSAVLFSRTDPSHKMQLVKLLQE QKLICAMTGDGVNDAPALKKADIGIA MGSGTQVAKAASKMVLADDNFATV VKAVGEGRAIFNNTKQFIRYLISSNI GEVACILLTGLCGLPEALSPVQLLWV NLVTDGLPATALGFNAPDEDIMQQP PRHVDEPIVNGWMFLRYMVIGVYVG LATIGGFLWWFLSHGFNWKDLTTYA ACTDMQDAKCAILADPETARAIALSI LVLVEMLNALNALSENASLITARPSS NIWLLLAIFSSLALHLMIMYVPFLAA LFNIAPLGVDPLIVKEAQPFSVLVPS NFDDWKAVIVFSVPVIFIDELLKYITR HMQKSRNKKNN	None	None	None	None	None	None	None	None	None