	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	K2MTF7	K2MTF7_TRYCR	Trypanosoma cruzi marinkellei	glutamate dehydrogenase	19.946178	NaN	NaN	30984116	MMRRVVIGGLVARTAFCCSGMNGF RFMTSDINASCIINTVKEIGMFDEAS IRKAVQLFTGGLSDSSYGLCYTEDEV TNHVKGFLCAKAKSTMGDPFEYVH ESDDCAFYICRKDHESQLRAVRRLS RFVSREREGNQCISMRGYTTDDGTT SVYTAKWQPFVNPHPKPGETDIKQL ASHVFLEQRSPELQNRYQDLLTRFQ GSVGPVFTLVDGEDGVTCFSMAVAA DRNYYLASLMAIIHEIPGAVVIRSFSE TFSNGIHIYTLYIDGATGTQLQDRVS MVCLLPNRPPNSITRLHEDLIFNVE QTVFTDAAIVFAFYFTPNPTSDDYHH LRALLLKEPNGVNRLNSLRTSLSLE MMSERYIGTLISLYPEYMIDIYEDFR RGTTAESRQVIQNKIAARFREDQRTT HDLEIFTSFLRFNEVIVKHNFFKKEK VALCFRLNPLFLKSLEYPLVPHGLFL LAGGQWRGFHVRFTDIARGGVRMII SRENAYRHNKRTVFQENYNLAHTQ LLKNKDIPEGGSKGTILVSSRFLNRF DHARCQRVFLQYADALLDTILPGES GIVDNLKETEIIFLGPDENTAGTFPS DGALLGKRRGYQAWKSFTTGKTPS MGGIPHDVYGMTTRSLRTFLRGVYE KLGLNEGDMTKFQTGGPDGDLGSN EILQSNEKMLAISDISASLHDPKGID REELKRLALGRLQLRHFDNPTLSEG FLVLTEEKNVTLPDGTFVSDGASFR DEFHFTKYTEADVFVPCGGRPRSVT MANVGRFLKVPDADGESMLAGRFE STEALKYKIIVEGANLFISQDARLALE RCGVVLFKDASTNKGGVTSSSLEVY SGLALSDEEHGKYMCVTSKENAPEF YKNYVKDIIHRIEDNARREFEAIWRD HESHPGMSKTIISDTLSEKNVKVRA NILASDVFKNKKLVRYILFHYTPKTL LEVVPVDELMNRVPIAYQHAICAMW LASEYVYSTGIDSNEFDFFTFMSTHI ERASAMAKEQAR	None	None	None	None	None	None	None	None	None