

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
V5B6G9	V5B6G9_TRYCR	Trypanosoma cruzi Dm28c	Aconitase hydratase	19.946178	NaN	NaN	30984116	MFRSTASRTGNPQLNPFIKYVATM AADGGEAKYFKLHEIDPRYETLPFSI RVLLESAVRNCDEFDITSKMVDNIF NWKETCHRNIEIPFKPARVVLQDFT GVPCVVDLAAMREATKRLGGDLNKI NPQIPVELVVDHVSQVDKAGIPTAVK ENQDMEMQRNRERFEFLRWGSKA FDNLLIVPPGSGIVHQVNLEYLARVV FNNKGLLYPDSVVGTDSTTMVNG VGVVWGVGGIEAEAGMLGQSSLM VLPQVLGYRFTGKLTGCTATDLVLT VAKNLRKFGVVGKFEVYFGPGVDNL SLPDRATLANMAPEYGATTGFFPIDR ETINYLRCTNRSVEQLERIEAYAKAV KMFRTGDEKIEYSHHLELDLSTVEP CVAGPKRPQDHVPLKNMKEDFAAC LQAKSGFKGFGIPAKDVNKTKNYMV DGQEAVMRHGSSVIAAITSCTNTSN PHVLVAAGLVAKKALEKGLKVPPGIK TSLSPGSHVVTRYLEAAGLQSSLDAL GFNTTGYGCMTCIGNSGDIHAEVSK CISENNFVAAAALSGNRFEARIHP LTAANYLASPLLVVAYALSGRVDIDF NEEPIAKGVFLRDIWPRNEEVQEIVS RYVTPELFKSVSNITTINEQWNALQ VNEGKLYEQPNSTYIHHPPYFESM TMEPTPTSVIKDAAACLALFGDSITTD HISPAGTIAKDSPAKFLQDHGVERK DFNTYGSRRGNDLVMVRGTFANTR LGNRLVGEQGTGPFITYFPTNEKMYI FDAAMKYQENIPLVIIAGKEYGSGS SRDWAAGPFPMQGKIVIAESFERI HRSNLVGMGIVPLQFKPGESAQSLG LTGKERYSFDFSGGLRPGQETTIVHK GDGSSFSTILRIDTEMEVKYVENGGI LQYVLRKIRGAXXXXKPKNEKERR RNGNNSYIYIKSILHTCEQWCSWG AQLQLYLHLFILLALCIALC	None	None	None	None	None	None	None		

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