

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
Q4DU60	Q4DU60_TRYCC	Trypanosoma cruzi	AP-2 complex subunit alpha	19.946178	NaN	NaN	30984116	MDMRGLAHFIQDVRKATSSPADEK KRVDVELVKIRSKFRSTAEMTTYDR RKYVCKLMFIAMLGYPVEFGHMEG VKLLALKSPAELKIGYLSATVFLHEN HSLLLATHMIYKDLLSEQEFNINLA LTAIANAGGKEFAEVMSSRVKSILLS DRWNVHVRKKAVALTYLRIRKYPDV VDLGDVIPVVTDLLSPLLGMSCCAA VFLTGCLNKSNFHLFHSVPNRVIEL LGRRIILEKQTDPGYVYVGPAPWLQA MSMRLLRGFPVPTDLSLKEKILLVLR RIVQATDRVLRDAQTQKQKGTMN RVSAMNAVSEVLLVQLQWDLDTLRL HNECLDVLSWVFMKKDSNLRYLG FSLLSQLCSARSSYNDYRTRYCQYQP QVVVALHDPDVSIRTKALDVTVCMC DAETSREGIGALLSYLPIADGLFKEN LVLAISHLAEVYCVDYGWYVDTILSVI AHAGGLTPQFIYRVVNVINHKYVQ KRAATTLFSLKIRKTVPEVLLRVAAI VLGEFGYQIAVNAESTPLMQVTVLQ NHLKDASEETQSLLLSTFVKLYNYY DIAVREKILNTLCQYRSSFNVELQQR AMEYVGLIELGNDELLOKVLEPLPSF KEDDFSLVLPGRAQREANVAVSLN ISRDKREKLVAPRADKEPKLNSSDL LVINRDVPLNSGGESVDLEALFTV NSTMQLDRMAKENELRRRFIFLLE ETSGVLYSDEFVELHCEQQYRGADA RVTVTVRNKMAVGLKVVHETTGVD VGLLLQSRNERGDIASSGVLRFVFA RSWAPFREAPTIVRLSYKRPGECTTE MKILALPIVTRFLSPYDEKDEMAYD TLCKQMNSASVRRRLSAGCVTHN GAIEGVLKGMGFCVIHPVDSAEWKA MAAHATQSHGILSYEPVAVKVFVSHS PGAFTAQVITKSHLLSVAIDVLEQTL LSKGQTECEERFSVFGGNLTTLS	None	None	None	None	None	None	None		

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