

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
V5B6A1	V5B6A1_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MVKKQPGAENIRVVVRCRNLLPYEA ERGDKALVRLDLATNQVIVQHPIGD ADTFAFDVYNNNTYTQRDIFLQEVQ PLVEAVLQGYNATVFAYGQSGSGKT HTMTGKLSDPQMWMMPQVNNYL FNEVRKLSITTKSFKIKVSVVELYNG KSRDLLASKQVNLEIKQNMKNFYV KGAEMPEVTSFDEAIRWFNAGTDRR QTASTEINDTSSRSHSLFTLQVEQF DFEQDPSSPVLTSKINVVDLGSEK LSKTNATGETAKEGCNINLSLALAT VIDTIVKGGKHIPYRGSPMLTKDS LGGNAKTMVFANIGPSEKNISETIST LRFALRAKQIENKPIKNMDPKDARIO DLLDQIEELKSRLGNVDLNQEDQLR QRIEELIENADLRGGGVQNNIELEE DIRLQAKLEEANGMVAEKQKELEK GIDGRSLEWNLQSELGRGKELRSL AMNFIKQVCQEEQIQAIRSKMPQDG SYNTTDDGWDVKEIEFYLREFAAMY EAWRQSTYTREDLEKYAHLVSTGAL EQTORQLDEVARARDELVRLQEDS VKRAAENELVSQLRVDLNTLKDENT KLREKVERDQERMKAKIFKNKEEYK SLAEELERSKALVTEKERDIERLKRI SEETGNNAIASSTGGASWANADE RTAMLQQLNEEKNIKALESRIREV NVSLRRYGVCIAADAKEVTNESREVS EATDAFILAALEEKDTMDGDLCAKL QQQLRLKQRLTEL RHQHQRKLDMM LRKYELIKTGRTTANSGAGAENNML HPGLADEATAQIKELLHRKDEEIER VIKEKKTCDRIVKLNKCELRIKEL EATLEEERVQSTEERQELDRENREL QRYNQOLSVELETLRGQLSEMKEE MDFMERAKSAEIERWKEEAANGAG RLEEMRFKMAAYSSELQKSYDRLQE QLQRTVSLKEKSDNLENTROMVK WSNSLLATEKQKVEQMEMMLKQQ EVEMRRREEQLKSEMMNDINKQVA ANNRRMQDQAAQYQELISQEQDKQ KSIMKKVSSRKETQKAAQRYDEMI LENEALLSKLEELKVTSMRIFLEKQE AQRELEALRPTKHVANRRL	None	None	None	None	None	None	None		

0