	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	V5DPI3	V5DPI3_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MTVLTKSRNQHNMSGDSPSMSSSG AFSFESVVPKPPTTWSEHMRSVLLA AGRMLSLFLFFFFTWPILLLFHGLLT LLDRHNRRVVLKHQQRWPFESVPH VRPVRVAVSREFPIENWHLRCEDGR QRWCYGELLNEEEGNTLGKAQAAG LTILPSEDVPMVGEHYEWAAGNAVT PTKEPDMRAIKAKQRRFLEQYQLGL HNTTEIRRRASPEDAMRDGVEFLLR LQDPYSGHWPNDYSGPLFLTPGFIF TKFIVAGGDIRKMFPPHRDHQHKND EPCRCGEAERVEMIRYLRNYMNKD GGFGQHTEGHSTMLGTALNYVALR FMGVPADDADATRARAWIRSHGGA VSVPTWGKVWLCIVGLYSWDGINPI PPELSLLPAWFPLSQGRLWCHSRVI SVPFSYLYGIRWAPEPHPLLEALRQE LYTEPYDQIQWDQHQSNICNLDCYT PISSTYKLVAVLLKLYEKWHIKSLRR HALEVAWSHVAYDDEDTKFICLGPV NKALDMLLTWIREGENSGRYQNHV SRLADYIYMGPEGMRVCGYNGSQL WDTSFAVQAVCACNMELLYPQQMS LAHHYVDVAQVQKDPKSAAHFYRH RTKGAWNFSTASQSWQVSDCTAEG LRVVLLLRHNPFSVSRIRDAVDEILS LRNAKGGWASYEPTRAPLYVELFNS SDVFRDVMTDYGYAECSSSCIHTLA LFREHYPGYRRAEINAAIREGLKFVL SLQRPDGSFYGSWGVCFTYAAWIVA SALCISREIPDMANHPSCVRLIDFLL SHQNADGGWGEDVTASVRSLWVD NPSGSQVVNTAWAVMAIMSAAGEA ARSLPRWREQISVAVERGIRLIMSRQ LVTGDWAQERISGVFNGNNPIHYPG YKNTMPVWALGMYNKWKKTYTAPT APLMHPSGGI	None	None	None	None	None	None	None	None	None