

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
V5B2J6	V5B2J6_TRYCR	Trypanosoma cruzi Dm28c	CSC1/OSCA1-like 7TM region domain-containing protein	19.946178	NaN	NaN	30984116	MDLSLKPRSTWDDQGLWFTLTLNSI VLVLSGLSIFCLTAPLLWLRRLYRT NHTITVSVGPSGFPAATTPLTASSP SASPTPPRQRVGRVLPYRCSLFSLS LRDWLSIEEEVANLESVVQPLSTTAS PKDDEDDMKRQISGEMEMSKVNRK ENTAVEDTFLVSATKAKVFPDQVA LYLAFLLKLFATVFLGGVFNFWICVL SGTDNYLEKYMVRRDIHGCQSLGD NETACNDRAPFCYVETDCLPVRLR GLYDLSAQNISPHSWRLWLVGILD FFSAIYVAVLFFLYRFDAYIKSVMH LQMENALGHRVAIVRGLSADMMSE KRFHQGYLVEEAYFRPRGGKGRVR NGDSGNDNNGNAEEDELVDVYYOYH CTGLSCLFSICGFYKTRRDAIFT RDCSVRQLLFPKSPGMYESMERT ETAMKELQEAADQRAYSKKQRERM PVAHERGEDVEETLMMCAPFPSCC SKVPKVEYRKKVFAEAKRLNQFIDA TPARDAKGSAFIIFSDALEAYEFVNL FESRRGGVSGAQATIAGPPDGIHFN ITTDRTGWLRVTVFFVYILLFFW AVPVGFVSSIDNLVYIPYIGKFLWY MKVPETLRGAVTSFLPVVMLELNL ILPFMIRCFVIAMGVVNRDSLGGEL FLQYLFMVLTCVIFQAALQGGLSQL ADVLDKPSGEAIGFFLAIVSPNGGY WYAKVITAACISTWLNLDPLPLIRA FLRRKEVSVQRVYDKLFKPCSFDP QFYSFDLTILAMGLLFRMTVPLLALL VGIYFFVRYMTQRCRLYDRYRPHIP LHDCTVVSASAQVMRAASWLYCLG GIGGVLVMSLRKXVGGVLCISLAV GILLARVHVSRRWVNSLANARRF LKHRAVPQTRYGAMAKRKATQNR QRTDKPSAPPQSTNVLSEGCSMEQL ATELVSQTFETDGDVDNRAATDRSD SSTDDNNGVVDGNINAREGATSF QVPIPAATELLPHNPNVISRYQPKHQR LLHIDAEQEIQRMEKTVFHVESYWN SSFELIDDVDMTPLKKKRGVCRNEIS SIDQCVAHEENKKLFLFLFLFIFL LKNENSSEATFIYVS	None	None	None	None	None	None	None		