

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
V5DN77	V5DN77_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MNLKYPRFRISVDRGGTFTDVGDV TFEDGTTTHWTKLLKLLSCDPQHY PDA PSEGIRRMRCYIPDVVPPARGLIENEY LDEVRMGTTVATNALLEHNGERSV MVLTEGFADILDIRDQARPSIFALHI KKNPVLAECTVEAKERVHISLKEAR STIPWPKEWIKCGESYIADVVRPLEK DDLKQKLOKAYNDGIRSVGVSLLHS YAYSKHEEEIKEIAKEIGFTNISLSSE LMALMKYVPRSTTASVDAYLTPLIFN YIRSFRAFKNMLGTRLVYFMQSD GGLTSADDDFFGFRSVLSGPAGGVMG CARCAVQDLGEHVQVVGIDMGGTS TDVSRCDGENVSHIVESEISGIQLHA PQVRVHTVAAGGGSLLKWENGIFCV GPGSAGSHPGPACYGKGGPLTVTDA NLVLGHLNTEFFPRVFGHHADEPLH VNAARALFETLTKIAKDTGKPWTV EEVAHAFVVVANETMCRPIRNITEA NGYSCPSHTLAVFGGAGGQHACAIA RSLGMGRIYFHFPLASLLSAVGVSM DIVQERISSQIYINQKNAEAAIQEQF RSIATSAKEELQRLGFDHIVLEHY LSMRYEGTSTSIMVLLDGTCEGQTV ESGSDGAGNLMHRVKANFTHCYE QQFGFLLMKREIMVDAVVRGRGV QIGKEARARIAQQKQNSAVHQPYS TRVKLAPRPISMKTYSNGWETVN VYVNVADNGPVEGPAVLVGEGETSILL ESNSIAYTNDGSLIHTSOLIEKIST ALHPLHLSIFSHRFMSIAEQMGNAL QRTSISTNIKERLDFSCAIFDKNGNL VANAPHIPVHLGAMGAIRWQREYY GDQWREGEMVLSNHPCGGSHLP DITVMSPPFFRGEVIFYVASRGHHAD VGGTTPGSMPPFSKTLQEEGAAIKTV KLIEDGIFQEEIRIRELLKPGKARM SGCRTLEDSISDLHAQVAANKRGMQ LLKELMETYGYEVVQAYMGYIQDFA EASARDALKRVAKFYGTTLPSVDYM DDGSEIRLKEIDPETGSSCFDFTGTS SQVLNSTNCPTAVVYSAIHCIRCIVD ADIPLNQGCMRPIRVVLERGSILSPD DELPVAVGNVLTQRVTDVIFTALRA VACSHGCMNFTMGSSDVAYYETI CGGSGAGDGFNGTSAVHTHMTNTR MTDPEILETRYVILRFRIRRGSGG NGKYKGGDGVRSFLFLQDLTAVLL TERRVLEPKGLFGGGNGMKGLNMV YVPNETAARVSVSPQEMLHDVRR GEWTREEETRSDGFVYRARNAG GKNVLAVRCGDILTIHTGGGGCYP EEAKQENKRGTIHD	None	None	None	None	None	None	None		