

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
Q4DSI2	Q4DSI2_TRYCC	Trypanosoma cruzi	NaN	19.946178	NaN	NaN	30984116	MSGRPEEGLYGNVESQSSNVSQGG RRRTRSEFEGDIDDYSSITRIRLEGMY RPQWTMSSSVKDILLEGSTLRDMDK LNEFIRSNLGEWVVERNNGNVIMEA FVRRPNAYVQDQQLLEEILNLTAYQ ALKMELEERKILLEAAYKLHHEGVYF LRQWGEHEGKDTVSSHARGKLNQV LTGIEIEEKERRDAEGEKARREEKAR RHAEKARREEQINFNLTLSEIKDAI FQGRARVYKMKLNRFKMELDGRG VADTNRSVLLKEFFKDPTRYVVDAG VLNEIQASDAYLGAERAVREEMDME KDLRKLHYKHLRLLGWSTATAEIR ASVCDITKRFLDAALQEARNPPTTSA VMNLEGLYESVYNARWHHAVEVVG GERTQTGTGMEAKEGKPKKSWTYR AVGDTLEKDDAVQQSGAAPPVLMVL TSDRGWPYTLNGPHGCGNDLCVNC EVERVWQIVLDDLTKWLSNDDLTLN PSPPLRVLIGTPGIGKSMNAGSYLLY QLLHYDVEKLVQVVIHCFGSAMYVFD KTTKIVTKYMGNKTSKSVLGGWLQ RGMKGYHYDVAKKGTTPPDFAPST GWGMIVVSSPNLDNYDEWKTQLKA SRIIMNCPDEVVKAMCAWMKRGL EQDKQAGYWEKVEKRMKVGPIPR HIFDENEYGRRLAAVDDALSAIKDTD VEEYFSMGGEKKWYSKNPCHKLV KIVRERKVEGTEVFLNASICTDIEFRI ADRLAKKKGAKDLLLILASRGTLAS QLVEQLGLRAFMYGEFVIALKELKE LPPSPSKPRSSVLNVNHQGYPTRA VGLKELQGGVTRIPMEYGVLYIPKVE NFPLVDGFFVDSRKTLLVGLQITTA GEHHTIPSTVRLFKNNMATYFNGW EELSRDMSWEIVYVQHENSTMITN WQRCGPVNTENLSLAENEIVAFWN GKVHQYQFALTTDFVNKIREM	None	None	None	None	None	None	None	None	