

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
V5BPJ2	V5BPJ2_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MNSIHALLELGVQRRNVERESRNAK KENGEAKKQRAMGIVSAIEAKQQML LRSLAQRQFVYPASEYPTAELAEISD LFTNLLRDVVEQKIFLLYNCSSDAC EARRCTCAKEGSGEVGDCKGRQR GEGNCRLCSTSGRGLLDVLRISTN VLHESHAGASSVDNIPLIEQWDAE LQAALHHVGDVVSRAHTEMLVVDL ALSDGRFPYLTSGSSTKESPLWTEI VRLFEDAFESVLTDDVRLLLWCIHQ HLLGRGGLAARPLFAVSKSHWSYAS SVVTCPLPSELHERLMRRWKELIW IVVPEFRLSQMTKRSPMLGRVKRP MMRTMFMEDSANNRLWEALHYAT SORCDELQETVRGLTNTYSFCFSHE EYYYSOLGAORLRELVNTRVSYGMN GVKHAGAGAFVVQTEVLQOVALSKL EPALKSALVTTVPATMRGDEGITVFR WKALMDVKDDASAPAGSLNRLNAF RAGRKVPNKYDKGLEPLAHTQRFLE EQRAEVKRMQRGRGTALEEGDKTQ AETROSNNGSTEDNKRSAVAASA AVSRRYINRFASSTISPPILNKLKQK LAANATAIGKKMGYETTPPARVLPS ASTTTQSTNEEKTPTGSTRHVSPLHP QSGSGRKGGRNFPSRHSSEVRLPTL GNYTPSSSQQFVRTRPASREIAIQN GREQSAPVMSCVDISKQSGDEQKR PASKEAIAENHRDQPTPVATSVSSV VERDVADDPVAAEPNPLTATSTLNS PPKTAVVNAAAAESAKRRILSLKKVC PFLPEVPFSGIFLEDLNLTDPRVEAA MNRMRRAVRGNASVTRDYRSIAMEAE DSLCAVVEIAEEMFNATSRMLSR PFLESVRGVLLGNIPELETDEDIQS LLMIRDAERARRSTASIDKCLRERA EALINDIPYVVEPERYGAHPFLAAP SASSATSHRSRDEKANSDGDDSRAT QRVGLQVPTPPGSDATSEVDKTAQA ETVLNDGEPHRAMDGREEQPRQSN GQEQPMGAVVKPVTTGGFTSQPLAA DPGQKVVENNGALSERPYESHLPQ HSSFAKARMGGGAPACSTVQATE EKQNFVIPRVFVADTPLPAGRTGRAG EQVQLLSSKPPVGPVNASRENVPV PFHRQGGKPFSLPLEEKQGANVAQ RAPHGIPTSNNIQSLTDFPIENQKMT PCGPTFGTVKRFLESERQELERLVI LGGVRRAGMGLKEGKGLPLDAAK QMPSEEVGKRNSAEVTRQAGKLO TGLPKKTSSTFTTPNVADGNGSA GLSTPQAAARREGAQLGLLFEQLPY MRRSNEGNELLRKLIEERRRVSEGH YTRELDLAALQSRQLKRSSHGPLPM HPNFLAEVAARSSLGAAPNGADGG QARLLPSLSQGPSHDSCATQQQEQQ QQQQQQKHSYGNORSTWGRYTIE RNPPTSVTVQRRLSGTPLLQQSNE DRMRFTISVVISADHISQAPDKTEY EKSSNNHIGMSHRYMLLCROAGIKP NSMLLRALPDTQGVSVSRVDTSGNY IGPKGLMPLLQVLRNTGLKYLNLS HNNLENDEIVLVSFLMNPASGASLQ SLDLSDNPISLAGGAAILRLAQARPS LLSVRLRGTLPQPVIRESIEVCDSN RASAS	None	None	None	None	None	None	None		