

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
K2NHB5	K2NHB5_TRYCR	Trypanosoma cruzi marinkellei	Post-transcriptional regulator MKT1 N-terminal domain-containing protein	19.946178	NaN	NaN	30984116	MRKVGGNRRNERQGRGQQQGS MAPPVSSMGYQQHQHQHHQYHHH HHHQHHHSSSYNVYEGSRPPPHS MQQMSPHGYGPGGGGAPAAAGG NGWNTPQQQHQQMPMYNQGYG HMGPGAGGGSGGGYDMGGVYQD MHPGQYPPSPSPSHMGGGSGMPYM RNPPMHGGVPPAGPIPQYGGPLGQ GAPLPPMGYGRGGHPPMDPMQQG GMYGGPMVRHDGYPDNRISPPQV GYGPNVGAAPPPPPGIPYGGGNPMP VARSPAGVPTAAGGYDPRSKGMLN NPMPMPNAPRHPRVGTPEPGIMGG GIPPGRFNHQALAPPGLPPANYM PMGSVPPNQPPIMPAPTPNGDPLHV ERFSLGGPTENLYDFLYEHDLSVD NLSSFFPEGYDPKRDPPLKIAVDGNF CLNSLRDELKRKRDPLWFLHSTLPEE LLDSVCHHVEWMRSLNLEPIWVFN GLSVSGDVFETFLTSEGLRARDIVV SRLEEGEIPDEADIQTAFDQPLGEDV QMAVSRYLKELNVMAVTAPFLNW AQMVAFHKEGLADLLMGPPEVLLLP YDNMMLVQIDVSNVSYFDRERV LRVLFPHYVTESTRGAGDRLMDLG LITATHPALSSTSVGLNLTMEEVYEE LSSETPKFRSIKDFINAHAYVDANK KKAGLIKHSKGRGYLRYSGVFSSKL QDSPLVYLIRVLEPGLTNADMPTNLI GVLGNLVLPLSLFYLFVGLLSVRIMT VITQSYIRDECPVSDTKDYHEKLEVL MTMRSQIIAQVFKKISKDRHPNNT CLSWVRWFKPILAPMQRPRDLINLD EWDISSSEEVQLDEEHLERYSIASV LTFSAEAARPIRDQNICSPRNTPIRY HGKRETFFTVLLKAFDFLYGFSHST VSPMGEDEMMAGGVEEGVEAVG HGINAEGAPILITDSNSNHYHELHD SLPEEGINEYPTVYFPTYLMNAIKHN PRSFQNAFVVLTEILRVGIIGSLPCRY IDPADPQLAISMENQESNC DPRVLL ASRIGCLINLPYRRKTEDLPFIWAPVY SRHLCAFTVMVRAM	None	None	None	None	None	None	None		

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