	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	K2MM88	K2MM88_TRYCR	Trypanosoma cruzi marinkellei	NaN	19.946178	NaN	NaN	30984116	MNAKEPRAKQQQEIMEETGTGIAPE MPPRTIEEVLAGAMWVASLTDPTLC LERLHETFNEVIFSCVDEEAEDEEE RVGRLKTAADRLRDQRAKEDLRRAL MELQDNTYVDEAEGEGEDEGDGVD EDDNDGEFSDESQLMNDEEFLLWI HRNDTQRHEDRAEVMGVDVSAADS AVNGVELQLHRVGLGEEDRLVAEDP LPERLKAIEMRERRPALGSRILPHGY SYKIPLEYEAFWILRQLRGAGQPLAH NTRICAIATRPPAEVADEPVVVSIAYA VQKMTQEYMEPAHLMLYHQTALHP LLCALLEASPFTEAAESTMPVASYAY SIKTGRRQRPFISDSYVGFSALADRSI GEGDPAWNRIGSNEADFAQRKPCFL ELGRLLVRILELDVLCHRLESQRQQL LLKLKDHPESSVQEQRVALESLVFE SGIEADLWQTFSGLLTMGQGRGPTQ VIKGLGLEDAFEEYTITGLQYQENLL ANAPIHVCQSPQTMPLMDWARQVG ATARSPRSAESVLFLFNQAIVEQFSK LPILRKRLFDVFCAEGDLIVTYKLCE RRDDVYPLATFFVEHPRHYLDLLER ERQGTVTLYYVLSEELVCREMNLRQ LYDRSPAADEWFLERERAMEILLVRI IDGLIPKVKTELCQFAQQFVQTQAVE RLGLMCAQGPYEPTRLCLEVYDDDA LTWEQEWNVVEALPLKQMHVDGYK PLRRVCAAYRGDSGITHISFIDESGS FIGSTRWENSAWSSETSRAADLMQ QNQLEKLCARCSPSVIVVGASNMAS LSLMRSLLRFLRGRVYAAFHVHIPVV WAPVEVARLYSSTTFAELEIPGADSL ARTSLALARYVQDPLHAICVLFDKER TALRLSLGTTAYITGEQEDQLYARLC WEMSLWVTACGVWVEDCVTRPTSV ASLQFLAGFGSSRARKLHQLLTLRR PNSREECGQLITECFGAYVSRNAISS LRFGPPSVASTGSGGSFSRWHLLDQ TLIPREWYSAAAFVARAALDNVPARL LALVEFMGLEASEKRDRIDRNAHQQ KMITAIREARRPEWDSLLGEREVEFL QEEMITAGQSFLRRPYRRLSDRELM SFVTGIVYCTQSSTSSDTRSVDTRSL VICEGDYVTGTVQGVRGGGAVPGIRL STSFGLGAFIAAEDIDDETMRQDVLL YEQSVRESAGMPQPPSNLTSPTWLR RGVPIQGIVVGCNWERCELRLRWCR PRDAISNGDRSNAEQGGKDDVDSD NHYLSYATCVNDAMSSANSLRTVR MNIDARVFATKISRHPLFRDTTSATA QTYLHDKRIGEVILRPATGRRNKAIA VVKIGERSTCNWLISEERRPNGAIYY RLKDKLTGREVEFGDVDEFLNNFIA PMVALVRSIRSHRRFVESARDVQDA LDAQRQSGLGFTYVFAEVSQQSRPP FYRVFTRGGVTERNFYLHIDESSIYV RVPIRRSGGRGGIDLRWVSCRNAEH VSEVVKGLARPS	None	None	None	None	None	None	None	None	None