

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
V5ATR8	V5ATR8_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MKSEDLADIEGVFDRFTRKSTQSTFDN SNGVQDEEWWKRDEYRWDYKHKG LKGGRWTSVVEGGSIRAPEFEIQMR EEWYRIQCWWKTDQRRNWLAFRD DEWWKEELYITDWLENGENGLMW NAADELSGYNRRGNRRQAVKIELDR RVQWYKANGPKGVVKTWCALVEGS KEHCTVEEKFERDDYYKNGDWWKS EIHVNSLLNPNPECCALRFAAAAAM DQEWVKQPPFRADFEKGGVLWKAL NEELAVRGVEDFAAEVEEVERRRNW FKNYWWWGGEKYIQDYEKNSAGKL WK AIDAGSTA IADAVLLRSREEWYLA CRETEWWWGGEKYIQDWADYGEKGF LWTAAWRNAALRGDGKNKASSLAL KKREYYEKNFVKQNKYVRDYIAFG EKGVLWKSNOVSAAKDSDDWWKSTS FFCQWLLANHAFPGFVKDPDVIS DYWKNKDTGKKWTASNILAALEKG NVDKASSELLIRERFYEKYWWRQR EYWNDFCARGTKWSALTPGNGKC SAVELKERENYFQPISLSRVVLGYOR LIESDAIAPMAGLCVSFEELCLRDQY YQWSWWKSQAGLTDYVFHGGDSLY IRAASYDDARALEGINFSSPIERATR MLFFDNGFLEYSADYEYTGFEVASA DWWQEPAIISDFLEKGDGKTKWKAC NSASGCGFAEKDGSSEEQRYRKD FYAANFWRSLLECLQDFRKNKGIKGL WTSSQPLGKGERVDEKHLFRAQVL SGYPEDSWIEEMETNWWKASEVRA DFYANGKSGCMVSAVNAYAVERL GTKIEYKADEEEIKRVEFYRTHPLS LEDGEKCHYIYLIEPFWITPESVEDY WMNGSRGKRWSASNAAFSCSLGD KFAATPAELKRREVLQRNFWRISA YKEDFLTHGKEGKLWNKSEYGGMG DVISHEEQIMRISYYLRLSEKYWADC RYAYDNWWKAPEVRNDYYINGSRG SLVGAATAAVASMGLYRNEQYGVSE EEKEKE	None	None	None	None	None	None	None		

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