

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							
V5BNF6	V5BNF6_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MLPHAEVETKSLILDLLIQNARAPQT SITSWLCGYPLYGSSEELLAGYCLEP VIVGANSREVEVAYPHIAVKYVKLVY LLRANPSLSTPSLQRLMRRSGNDE MFYFLQTLRPDQCSPLVLNKYAFVM KLLALDVFISIGTKKPEELHVSlySTP NSSLLEVLFLLLHVSPPSMSLKSEY VSPGVIAGHDQSVTVDFQWPSFCL KSLPLFPTNCVAPGGAEGYIFRANCS TPQYSIPLLYEAFQKESQLNKRSL QKEIRERLSVFVRANECLAAAYAGGV QFIDGWCSLAKIAVTVLDKIQQEHIV YLAQC VLEGVQLTSSLTVAQQEQIVY PLSQTLSLMAQLKRKA AVTHDGS GFPKNGAQGTIAAYGSRFGDET DENVSFSRRARPDTTDLVQTSAVSQ TAKSSLVENFNRLSRAVKRDLTGAI ESKQEATDMFDRSLISLENLQLLKS LLVTTVQWGSRVTMARQHfYSAMM AFLGTPGVSPDDITLQRHYTALFDVL VQDITSSYGGNKL SALALLGOLLNL SPQLCEMLCISSTSDGTS PQLMTCII SAFQAIDESISALFAHGGTEVASILW LAQIVFDVLMVISQKHAI RL TQINAL SYCTAMRTWKTSARVLLGH T TSEND AAAYGLENIDAYREVVKSVLLATRW FNVVLSALDNSEAALLPVAEFVRSN QLLFQSVLSSPYLDFHRRHVDLLTL DLFSEIGGMLYRLSCSPLVDECRALA HLFALPDL L AVLTKSEFTESFIFITGN TTAPVSVSSQLTKQTGSLPSPTQRQ QIYVAVQNILQFLLRAENLGALFATW GENGTAVLNAEGAVVEEQPPRLCDE ARNRVLMTITEQVVIMEKFFHHGE VVQAVDFI PHLVALHSLVLF LHFALL LDRAAHVKDAKSQWGSIVHALEEAQ RVVNLWRAFHS DYLRGLRVTRVSGD RWSVAAPS VSNQSIFPAISAPPQDDT SCR TN IYHQHLLDAPMRQSLKQKTD TAMEQSFTNQLSTVTLPTAGSGAMQ LGT TVDAASLAFDAVTLPPAAGDASI GALTETRRNEVALIDEVRSVYPGRN DVLIMESLLRQLRV SISNALRLASRT	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None

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