

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
V5DH61	V5DH61_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MNYYQIYDEVGKGRFSIVYKGRKK TIEYYAISSIDKSRRQRVLTNVRFLRS AEHRCIIKFHNWYETNNHLVWVITEL CSGGDMRQVLNLDNRISRQAVQMF GTDIAEGLMYIHSRGVVYGDLPKPSNI LMDSTMSMRFYDFGLSCDFNTIRG DGTVGTPLYMAPELFTKDGIPSIASD LWSLGCLLYFEGTGKPPFNGSDLAS LIRNIMSEPFPCHEALSDELNSLES LLVKDPIRRATWGDIVTHDFWQGRLL QLPTAPFPQPAFERRQRMVAVAGAS GDWPMSGEELRRHVWAVESAKRN YTVSQSASEGGPSKCPAVTEIDLQD HSAVNDATTEGGVAASRVGQKDG RRLRAGPTGAALDATNSQFLRKES FRGEAANESHTAANCQNTTINAVS NAEVSIVMNGLENLHLDLMAHTS DAHIRPLAMNSRIERFVEQKYDSEK MGFEPPAKAEIKTFTEQQQSDFIKMI YRTLSSSSLSHEEKLNLVLCYFEDVCT DAAIANFIVNSSIMTLCLKMVSQRN APSNYRATAASIMSILVRHATFIHTD LAKANIFSSIIIEVYGEEDSPRVKRLT ACLGELLIYIAVQREDRNRVWGINA NATLQLYLSVLKDADDVLRHYAVKAI ENLASVSDHGVALDVFAPKEVVNAL LTVYALPPASIRSEHMSSAACAALK LAVLKEDLMPVVMESPHLKVKYYGE VIAEVTAPKTAQSLTFLNMAVLSL VAIRNPIMERWVKDNTNIFSSSLAQ EKGKKIIRISGVAEAVVSGLCGVCE RASVAIKGKMLLVLLGCMGDKLL RLSSNKLIGCIDGVLRDKDSYVQRCA ACLASYLGVFLEAQLVAVAKGAPSGT TSTILDVAVCNVLSARNLGMVVELSD SVFVSLGTCLEKVMSSPVVVTYEEEF NRVVGLLTQNSVSLKRRRAVVNEL FPLYIHMLGGSNIERRFSALRRLSAF VAPLVDSGLVREKIAEAEKLDHL MQTVAASPELLKESEPIPALRLLA TCGERRSKAFANIATVELVGDVLYQY MLRSKQSDLSVPLQLLLLLLALQTER AAEIMDYLAHQEFFTAVLLKTLTMAV AKELDHLEPCCELSVFFLKHAVNN TSSRMAQCQLLLPQHGMETLWLP LCDSPVRSAAAGNAAACVYHFVQLSP AAQQDLMTTESLTLVREIHSNTRDNP TAVLYVLRALRYIWERVRKRHVOKL SEWLLAALKAVREEDPDPRELYSEAV AFMEFIGG	None	None	None	None	None	None	None		

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