

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
V5AV92	V5AV92_TRYCR	Trypanosoma cruzi Dm28c	NaN	19.946178	NaN	NaN	30984116	MVVGLSSSKGILALLNETDADVIQFA LKRLLLELVDITYWYEISPDLPPIEELH ESCDLSEETRMLAALVASNVHFHLG AYNDSVHFALAEAGTAFNYMERSLFT DTILSCCIDQFVQYQEQPEETRGKLD PRLDLSLFTALTNKWILQEGTPVRELV GFTIRARCLDFLEKVLQRDIALTKSA AILNYTFDLASVMLRDIQFRGKILRL LADLYAQGGLSTIDYFSRAQCCLFLG DSESTRLITDLMQANDKATAYQLA FDLYEYGNQEFLSAVVAALERESAEI PHPGLTSSPPENEGETSTKEVDTS NLSDVHQKLLAVLTGETSASFYKFL YAQCNADIHILNHIKQSIDAKKSVPH NATVIAHALMYSGTTIDAFLRGNME WLARANYWAKFVVTASIGAIHRGHI DEGLAVVDPYLPKEGVGLLPYQEQAG ALYALGLIHAPVGVTRNRKTIDYLN VLRKYSTSEQMIHGASLGIGLAAMG LKDEELFDSLFCVSGCDAVASEGA ALGIGLLMMGSGNRGAIQSMRALAS EDNQKEKTIRGISMAMALMMLGRE DECWAVANELDDSDPWVRLGGCF MLGLGYAGAENTKVLERLLSVAVKD TSDDVRRNATTMIGFLTIKDPALCLE LTRVLVDSYNPHIRYGVGMALAVAA AGTGDSSVIEVLWTLKDDLVDVFRQ GAYIALAIVMVQLTEAEPRVKELRE IFSKKIAGRKEDMCSKFGSIATGLLD AGGRNCTFSLHRQRHRLDKAVAGM FVFLQHWYWFPYLLMITLAMRPTCF IGLNDNLDVPEYTFKSNAPASRFALP KSVLTEKKEAKAASVKAVVLSSTTKKE EQFHRLKRMSSGSGRAHDAAEA SKNAAAAEPALTPPGAETEKKEPEPS FELLKNPTRVTSRQFAVISHDVG VPLKPYPTGICLLRDTKPSDGKQKIV SEVGWNSADDAPPEPFTWP	None	None	None	None	None	None	None	None	

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