

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
V5BPJ4	V5BPJ4_TRYCR	Trypanosoma cruzi Dm28c	AP-4 complex subunit epsilon	19.946178	NaN	NaN	30984116	MSKLYRAQTQQGHSRGGFFEYIRSVG ESKSKQEEDAIVTRDLAELKKTLAN NIEKRLLKEYVVRIFYAEMLGVSAAF AHIHCVNLSSSPDLLSKRTGYLGTW LTVGPEHDLMYLIVSNLQRDMKSSS FLDIAAALTAASKLVRLELMNAINTE VGLLRHPNALVRKKAIVSTMHAFYR KSEGLIGDTKIFRQMLCADPVSVMG AALPLFADVICTDPTSQRELISIFLSIL KQIGEHRLSREYEHGIPAPWLQIKL LQMLPILIGDEPSLARKCEEALREVIT RADNGLVIGYAVMCEAIRVITLIPTIP TLVELAAEAISKFLSARKANLRYAGI QALSQIVRLDPKYAHEHQHVVMACL EEADDITIRRKTMMLLLAMCNEDNV EVIVTRLVKSLSRITTDKYFREEFTRRI CDVVERFSPGAVWYIETMKNKLLCA AEHVPIQITIQGILKLIVEGEGKDGK DAAFRTFCVETYFDLLEGSQKNLPE AFCRVAWVIGYEGFLAKRISRTMML DRLCDMLERAECADTRGWIIAMAMIK IVAHAGAMPDNVEDLITRFKDSRSV WIOQRCYEFSELVKMPVLMKKVLPL DGCCEEVDLDESMGFLDGIVQEALL AGARPYEKREILLGVKEEETLRDAY ELVRTGVVNEDELDPENFKTEEPHK LVIRPAVKRWGLKNLEEDITPTEGNP TGGPEASEATGSMDCFSEGAVLHHV DPHAPSSLSTPSAAVNLMPPREVK PSKNKKFLNDIFSVGNKKLNVRGH NGATESLVHAGKGSEENDAGSLLSI HMEYVRELATTGVVVRIVGKVGVEN VVISMRAFNCTLRVVSCSATSATV DGTAVTLNKLQANQTVVEVEMQVLP VGFPEGRVLVEITYRSVAVPGDPQCA SVSRALSARDILRPPAQMTTAEFGQT WVGLREECKLQMSATPLTPQSLQN LLMERASLRVVEAIRNELITAAKLIGT GKPVLCHVDVVGPNANLTVRADD KGFAAYVAHALSPP	None	None	None	None	None	None	None		

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