	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q5FFE6	EFTU_EHRRG	Ehrlichia ruminantium	Elongation factor Tu	25.830362	NaN	NaN	30930869	MVDGRKPHINVGTIGHVDHGKTTLT AALTTVLAKRLSGEGNKSVKYDEIDK APEEKARGITISTAHVEYETENRHYA HVDCPGHADYIKNMITGAAQMDAAI LVVSATDGAMPQTREHILLAKQVGV KDIVVWMNKCDVVDDEEMLSLVEM EIRELLTKYGYPGDDIDVVKGSAVKA LEEESADGVWSEKIMELMNALEKID LPIREKDKPFLMSIEDVFSIPGRGTV VTGRIERGVIKVGDKIDIVGLRDIQST VCTGVEMFHKALDAGEAGDNAGILL RGIKKEDVERGQVLSAPGQIHSYKGF KAEVYVLKKEEGGRHTPFFSNYQPQ FYVRTTDVTGNIKLPDGVEMVMPGD NISIEVNLDKPVAIDKGLRFAIREGG RTIGSGIITEILE	None	None	None	None	None	None	None	None	None