	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	Q5FFZ1	CH60_EHRRG	Ehrlichia ruminantium	Chaperonin GroEL	25.830362	NaN	NaN	30930869	MANMVVTGEQLDKSIREVVRILEDA VGCTAGPKGLTVAISKPYGAPEVTKD GYKVMKSIKPEDPLALAIANIIAQSAS QCNDKVGDGTTTCSILTAKVIEEVSK VKAAGADIICVREGVLKAKEAVLEAL KCMKREVLSEEEIAQVATISANGDK NIGTKIAQCVKEVGKDGVITVEESKG FKELDVEKTDGMQFDRGYLSPYFVT NSEKMLVEFENPYILLTEKKLNIIQP LLPILENIARSGRPLLIIAEDVEGEAL STLVLNKLRGGLHVAAVKAPGFGDR RKDMLGDIAILTGAKHVINDELAIKM EDLTLCDLGTAKNIRITKDTTTIIGSV DNSCAHVQSRICQIRMQIDNSTSDY DKEKLQERLAKLSGGVAVLKVGGSS EVEVKERKDRVEDALHATRAAVEEG VVPGGGAALLYTLSALDNLKSKNDD EQLGINIVKRALQAPIKRIIKNAGSEN APCVIAHLLKQNDKELIFNVDVMNF ANAFTSGVIDPLKVVRIAFDFAVSLA AVFMTLNAIVVDIPSKDDNSAAGGA GMGGMGGMGGF	None	None	None	None	None	None	None	None	None