

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
000418	EF2K_HUMAN	Homo sapiens	Eukaryotic elongation factor 2 kinase	21.731224	NaN	S18;S27;S61;S66;S70;S71;S72;S74;S78;S243;T348;T353;S359;S366;S392;S398;S435;S445;S470;S474;S477;S491;S500	23301498	MADEDLIFRLEGVDGGQSPRAGHD GSDSDGSDDEEGYFICPITDDPSSN QNVNSKVNKYYSNLTKSERYSSTSGS PANSFHFKEAWKHAIQKAKHMPPD WAEFHLEDIATERATRHRYNAVTE WLDDEVLIKMASQPFGRGAMRECF RTKKLSNFLHAQQWKGASNYVAKR YIEPVD RDVYFEDVRLQMEAKLWGE EYNRHKPPKQVDIMQMCIELKDRP GKPLFHLEHYIEGKYIKYNSNSGFVR DDNIRLTPQAFSHFTFERSGHQLIVV DIQGVGDLYTDPQIHTEGTDFGDG NLGVRGMALFFYSHACNRICESMG LAPFDLSPREDAVNQNTKLLQSAK TILRGTEEKCGSPQVRTLSGSRPPLL RPLSENSGDENMSDVTFDLSPSSPS SATPHSQKLDHLHWPVFSDDLNDMA SRDHDHLDNHRESENSGDSGYPSE KRGELDDPEPREHGHYSYNSRKYESD EDSLGSSGRVCVEKWNLLNSSRLHL PRASAVALEVQRLNALDLEKKIGKSI LGKVHLAMVRYHEGGRFCEKGEW DOESAVFHLEHAANLGELEAIVGLG LMYSQLPHHILADVSLKETENKTK GFDYLLKAAEAGDRQSMILVARAFD SGQNLSPDRCQDWLEALHWYNTAL EMTDCDEGGYDGMQDEPRYMML AREAEMLFTGGYGLEKDPQRSGDLY TQAAEAAMEAMKGRLANQYYQKAE EAWAQMEE	True	False	4.581	2.889	2.006	1.951	1.188	1.921	2.008