

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	ext reg
008796	EF2K_MOUSE	Mus musculus	Eukaryotic elongation factor 2 kinase	33.906399	NaN	S27;S61;S70;S73;S77;S242;T347;T352;S358;S365;S391;S397;S434;S444;S469;S473;S476;S499	21540332;22645316	MADEDLIFCLEGVDGGRC SRAGHN ADSDTDSDDDEGYFICPITDDHMSN QNVSSKVQSYYSNLTKECGSTGSP ASSFHFKEAWKHAIEKAKHMPDPW AEFHLEDIATEHATRHRYNVAVTGEW LKDEVLIKMASQPFGRGAMRECFRT KKLSNFLHAQQWKASNYVAKRYIE PVDRSVYFEDVQLQMEAKLWGEDY NRHKPPKQVDIMQMCHELKDRFGQ PLFHLEHYIEGKYIKYNSNSGFRVDD NIRLTPQAFSHFTFERSGHQLIVVDI QGVGDLYTDPQIHTEKGTDFGDGNL GVRGMALFFYSHACNRICQSMGLTP FDLSPREQDAVNQSTRLLQSAKTILR GTEEKCGSPRIRTLSSSRPPLLRLS ENSGDENMSDVTFDLSPSSSATP HSQKLDHLHWPVFGDLNMGPRD HDRMDNHRDSENSGDSGYPSEKRS DLDDPEPREHGHNSNGNRRHSEDED SLGSSGRVVCVETWNLLNPSRLHLPR PSAVALEVQRLNALDLGRKIGKSVLG KVHLAMVRYHEGGRFCEKDEEWD ESAIFHLEHAADLGELEAIVGLGLMY SQLPHHILADVSLKETEENKTGFD YLLKAAEAGDRHSMILVARAFDTGL NLSPDRCQDWSEALHWYNTALET DCDEGGEYDGIQDEPOYALLAREAE MLLTGGFGLDKNPFQSRSGDLYTQAAE AAMEAMKGRLANQYIEKAEAWAQ MEE	False	False	2.9	2.971	2.044	1.826	0.993	1.824	1.94