	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
0	O95382	M3K6_HUMAN	Homo sapiens	Mitogen-activated protein kinase kinase kinase 6	31.708705	S1286	T806;S964;S984;S1129;S1149	30620550;22564745;37217939;30059200	MAGPCPRSGAERAGSCWQDPLAVA LSRGRQLAAPPGRGCARSRPLSVVY VLTREPQPGLEPREGTEAEPLPLRCL REACAQVPRPRPPPQLRSLPFGTLEL GDTAALDAFYNADVVVLEVSSSLVQ PSLFYHLGVRESFSMTNNVLLCSQA DLPDLQALREDVFQKNSDCVGSYTL IPYVVTATGRVLCGDAGLLRGLADGL VQAGVGTEALLTPLVGRLARLLEATP TDSCGYFRETIRRDIRQARERFSGPQ LRQELARLQRRLDSVELLSPDIIMNL LLSYRDVQDYSAIIELVETLQALPTCD VAEQHNVCFHYTFALNRRNRPGDR AKALSVLLPLVQLEGSVAPDLYCMC GRIYKDMFFSSGFQDAGHREQAYH WYRKAFDVEPSLHSGINAAVLLIAAG QHFEDSKELRLIGMKLGCLLARKGC VEKMQYYWDVGFYLGAQILANDPTQ VVLAAEQLYKLNAPIWYLVSVMETF LLYQHFRPTPEPPGGPPRRAHFWLH FLLQSCQPFKTACAQGDQCLVLVLE MNKVLLPAKLEVRGTDPVSTVTLSL LEPETQDIPSSWTFPVASICGVSASK RDERCCFLYALPPAQDVQLCFPSVG HCQWFCGLIQAWVTNPDSTAPAEE AEGAGEMLEFDYEYTETGERLVLGK GTYGVVYAGRDRHTRVRIAIKEIPER DSRFSQPLHEEIALHRRLRHKNIVRY LGSASQGGYLKIFMEEVPGGSLSSLL RSVWGPLKDNESTISFYTRQILQGLG YLHDNHIVHRDIKGDNVLINTFSGLL KISDFGTSKRLAGITPCTETFTGTLQY MAPEIIDQGPRGYGKAADIWSLGCT VIEMATGRPPFHELGSPQAAMFQVG MYKVHPPMPSSLSAEAQAFLLRTFE PDPRLRASAQTLLGDPFLQPGKRSR SPSSPRHAPRPSDAPSASPTPSANST TQSQTFPCPQAPSQHPPSPPKRCLSY GGTSQLRVPEEPAAEEPASPEESSGL SLLHQESKRRAMLAAVLEQELPALA ENLHQEQKQEQGARLGRNHVEELL RCLGAHIHTPNRRQLAQELRALQGR LRAQGLGPALLHRPLFAFPDAVKQIL RKRQIRPHWMFVLDSLLSRAVRAAL GVLGPEVEKEAVSPRSEELSNEGDS QQSPGQQSPLPVEPEQGPAPLMVQL SLLRAETDRLREILAGKEREYQALVQ RALQRLNEEARTYVLAPEPPTALSTD QGLVQWLQELNVDSGTIQMLLNHS FTLHTLLTYATRDDLIYTRIRGGMVC RIWRAILAQRAGSTPVTSGP	None	None	None	None	None	None	None	None