

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
Q6ZN16	M3K15_HUMAN	Homo sapiens	Mitogen-activated protein kinase kinase kinase 15	19.994096	NaN	S994	37217939	MESGGGNAPAGALGAASESPQCPPP PGVEGAAGPAEPDGAEGAAGSSGE GESGGPRRALRAVYVRSESSQGG AGGPEAGARQCCLRACEAEGAHLTS VPFGELDFGETAVLDAFYDADVAVV DMSDVSROPSLFYHLGVRESFDMA NNVILYHDTADDTALSLKDMVTQKN TASSGNYFIPYIVTPCADYFCESDA QRRASEYMOPNWDNILGPLCMPLV DRFISLLKDIHVTSCVYKETLLNDR KAREKYQGEELAKELARIKLRMDNT EVLTSDIINLLLRYRDIQDYDAMVKL VETLEMLPTCDLADQHNKIFHYAFA LNRRNSTGDREKALQIMLQVLQSCD HPGPD MFCLCGRIYKDFLSDCKD DTSRDSAIEWYRKGFELQSSLYSGIN LAVLLIVAGQQFETSLELRKIGVRLN SLLGRKGSLEKMNNYWDVGGFFSV SMLAHDVGVKAVQAAERLFLKPPV WYLRSLVQNLLLRFFKKTIEHSPR QERLNFWLDIIFEATNEVTNGLRFP VLVIEPTKVYQPSYVSINNEAERTV SLWHVSPTEMKQMHEWNFTASSIK GISLSKFDERCCFLYVHDNSDDFIY FSTEEQCSRFFSLVKEMITNTAGSTV ELEGETDGDITLEYEDHDANGERVV LGKGTYGIVYAGRDLNSQVRIAKEIP ERDSRYSQPLHEEIALHKYLKHRNIV QYLGVSSENGYKIFMEQVPGGSLSA LLRSKWGPMKEPTIKFYTKQILEGLK YLHENQIVHRDIKGDNVLVNTYSGV VKISDFGTSKRLAGVNPCTETFTGTL QYMAPEIIDQGPRGYGAPADIWSLG CTHEMATS KPPFHELGEQAAMFKV GMFKIHPEIPEALSAEARAFILSCFEP DPHKRATTAELLREGFLRQVNGGKK NRIAFKPSEGRGVVLLALPTQGEPM ATSSSEHGVSVPDSDAQDALFERT RAPRHHLGHLSPDESSALEDRGL ASSPEDRDQGLFLLRKDSERRAILYK ILWEEQNQVASNLQECVAQSSEELH LSVGHKQIIGILRDFIRSPHEVRMAT TISKLVLDLDFDSSSISQIHLVLFQFQ DAVNKILRNHLIRPHWMFAMDNIIR RAVQAAVTILIPELRAHFTEPTEG VDKDMDEAEEGYPPATGPGQEAQP HQQHLSLQLGELRQETNRLLEHLVE KEREYQNLLRQTLQKTOELYHLQL KLKSNICITENPAGPYGQRTDKELID WLRLOGADAKTIEKIVEEGYTLSDIL NEITKEDLRYLRRLRGGLLRLWSAV SQYRRAQEASETKDKA	False	False	2.513	2.626	0.869	0.774	0.538	0.98	1.13