

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q80U28	MADD_MOUSE	Mus musculus	MAP kinase-activating death domain protein	45.57388	S699;S701;T703;T704;S828	S155;S688;S691;S812;S817;S819;S857;S861;S915;S920;S929;S1058;T1060;T1065;S1109;T1235;S1237;S1266	33300544;22645316;40885482	MVQKKFCPRLLDYLIVGARHPSSD SVAQTPELLRRYPLEDHPEFPLPPDV VFFCQPEGCLSVRQRMSLRDDTSF VFTLTDKDTGVTRYGICVNFYRSFQK RMPKEKVEGGAGPRGKEGAHTSGA SEEAATGSSESGSTLQPPSADSTPDI NQSPWVKRRAKAGSRSRNSTLTSLC VLSHYFFSTFRECLYTLKRLVDCCS ERLLGKKLGIPRGVQRDTMWRIFTG SLLVEEKSSALLQDLREIEAWIYRLL RSPVPVSGQKRVDIEVLPQELQALT FALPDPSPRFTLVDFPLHLPLELLGVD ACLQVLTCLLEHKVVLQSRDYNALS MSVMAFVAMIYPLEYMFVPIPLPTC MASAEQLLLAPTPIIGVPASFFLYKL DFKMPDDVWLVDLDSNRVIAPTNA EVLPIPEPESELELKKHLKQALASMS LNTQPILNLEKFHEGQEIPLLLGRPS NDLQSTPSTEFNPLIYGNVDVSDV ATRVAMVRFNFNSANVLQGFQMHTR TLRLFRPVVAFQAGSFLASRPQTP FAEKLARTQAVEYFGEWILNPSNYA FORIHNTTFDPALIGDKPKWYAHQL QPIHYRVYDGNSQLAEALSVPPEERD SDSDPTEDSGSDSQDYDDSSSSYSS LGDVFSEMMKCDINGDTPNVDP HAALGDASEVEIDELQPKKEGEEPG PDENSQENPPLRSSSTTASSPST VVHSAHSEADSTEMGDKATAGISK PLPPVPSICKSTVDRRQTETGEGSV CQRTYDNPYFEPQYGFPEEDEEEQ GESYTPRFSQHVSQRAQKLLRPNS LKLASDSDAESDRASSPNSTVSN STEGFGGIMSFASLYRNHSTSFSL NLTLPTKGAREKTFPPLKVFGLNT LMEIVTEAGPGSGEGNRRALVDQKS SVIKHSPTVKREPSPQGRSSNSSEN QQFLKEVVHSLDGGQGVGLNMKK VRRLLESEQLRVFVLSKLNRAVQSE DDARQDVIQDVEISRKYKGMILL KCTVLSLEQSYAHAGLGGMASIFGL LEIAQTHYYSKEPDKRKRSPTEVNVT PVGKDPGLAGRDPKAMAQLRVPO LGPRAPSATGKGPKELDTRSLKEEN FVASVELWNKHQEVKKQKALEKQR PEGIKPVFDLGETEEKKSQMSADSG VSLTSASQRTDQDSVIGVSPAVMIRS SSQDSEVSTVSNSSGETLGADSDLS SNAGDGPGEKSAHLASSRATLSDS EITNSATSAIFGKAHSLKPKKAPAG SPIRSEDVSRVYLYEGLLGRDKGS MWDQLEDAAMETFSLSKERSTLWD QMQFWEDAFLDVAVMLEREGMGMD QGPQEMIDRYLSLGEHDKRLEDE DRLLATLLHNLISYMLLMKVNKNDI RKKVRRLMGKSHIGLVYSQVNEVL DQLNSLNGRDL SIRSSGRHMKKQT FVVHAGTDTNGDIFFMEVCDDCVVL RSNIGTYERWWYKELINMTYCPKT KVLCLWRRNGSETQLNKFYTKKCRE LYYCVKDSMERAARQQSIKPGPEL GGEFPVQDMKTGEGGLLQVTLLEGIN LKFMMHNQFLKLLKWW	False	False	2.987	1.915	0.848	0.91