	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	ext reg
0	Q80U28	MADD_MOUSE	Mus musculus	MAP kinase-activating death domain protein	33.64034	NaN	S155;S688;S691;S812;S817;S819;S857;S861;S915;S920;S929;S1058;T1060;T1065;S1109;T1235;S1237;S1266	33300544;22645316	MVQKKFCPRLLDYLVIVGARHPSSD SVAQTPELLRRYPLEDHPEFPLPPDV VFFCQPEGCLSVRQRRMSLRDDTSF VFTLTDKDTGVTRYGICVNFYRSFQK RMPKEKVEGGAGPRGKEGAHTSGA SEEAATGSSESGSTLQPPSADSTPDI NQSPWGKRRAKAGSRSRNSTLTSLC VLSHYPFFSTFRECLYTLKRLVDCCS ERLLGKKLGIPRGVQRDTMWRIFTG SLLVEEKSSALLQDLREIEAWIYRLL RSPVPVSGQKRVDIEVLPQELQQALT FALPDPSRFTLVDFPLHLPLELLGVD ACLQVLTCILLEHKVVLQSRDYNALS MSVMAFVAMIYPLEYMFPVIPLLPTC MASAEQLLLAPTPYIIGVPASFFLYKL DFKMPDDVWLVDLDSNRVIAPTNA EVLPILPEPESLELKKHLKQALASMS LNTQPILNLEKFHEGQEIPLLLGRPS NDLQSTPSTEFNPLIYGNDVDSVDV ATRVAMVRFFNSANVLQGFQMHTR TLRLFPRPVVAFQAGSFLASRPRQTP FAEKLARTQAVEYFGEWILNPSNYA FQRIHNNTFDPALIGDKPKWYAHQL QPIHYRVYDGNSQLAEALSVPPERD SDSDPTEDSGSDSQDYDDSSSSYSS LGDFVSEMMKCDINGDTPNVDPLT HAALGDASEVEIDELQPQKEGEEPG PDSENSQENPPLRSSSSTTASSSPST VVHSAHSEAADSTEMGDKATAGISK PLPPVPPSICKSTVDRRQTETGEGSV CQRTYDNPYFEPQYGFPPEEDEEEQ GESYTPRFSQHVSGSRAQKLLRPNS LKLASDSDAESDSRASSPNSTVSNN STEGFGGIMSFASSLYRNHSTSFSLS NLTLPTKGAREKTTPFPSLKVFGLNT LMEIVTEAGPGSGEGNRRALVDQKS SVIKHSPTVKREPSSPQGRSSNSSEN QQFLKEVVHSVLDGQGVGWLNMKK VRRLLESEQLRVFVLSKLNRAVQSE DDARQDVIQDVEISRKVYKGMLDLL KCTVLSLEQSYAHAGLGGMASIFGL LEIAQTHYYSKEPDKRKRSPTENVNT PVGKDPGLAGRGDPKAMAQLRVPQ LGPRAPSATGKGPKELDTRSLKEEN FVASVELWNKHQEVKKQKALEKQR PEGIKPVFDLGETEEKKSQMSADSG VSLTSASQRTDQDSVIGVSPAVMIRS SSQDSEVSTVSNSSGETLGADSDLS SNAGDGPGGEGSAHLASSRATLSDS EIETNSATSAIFGKAHSLKPKEKPAG SPIRSSEDVSQRVYLYEGLLGRDKGS MWDQLEDAAMETFSLSKERSTLWD QMQFWEDAFLDAVMLEREGMGMD QGPQEMIDRYLSLGEHDRKRLEDDE DRLLATLLHNLISYMLLMKVNKNDI RKKVRRLMGKSHIGLVYSQQVNEVL DQLNSLNGRDLSIRSSGSRHMKKQT FVVHAGTDTNGDIFFMEVCDDCVVL RSNIGTVYERWWYEKLINMTYCPKT KVLCLWRRNGSETQLNKFYTKKCRE LYYCVKDSMERAAARQQSIKPGPEL GGEFPVQDMKTGEGGLLQVTLEGIN LKFMHNQFLKLKKW	None	None	None	None	None	None	None	None	Nor