

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q58A65	JIP4_MOUSE	Mus musculus	C-Jun-amino-terminal kinase-interacting protein 4	11.096784	NaN	S109;S183;S185;S194;S203;T217;S238;S251;S265;S268;S272;T292;S311;S329;S332;S347;T348;T365;T418;T586;S588;T595;S705;S728;S730;S732;S733;S1188;T1264	28528544;28135057	MELEDGVVYQEEPPGGSGAVMSERV SGLAGSIYREFERLIGRYDEEVVKEL MPLVVAVLENLDSVFAQDQEHQVE LELLRADDNEQLITQYEREKALRKHA EEKFIEFEDSQEQEKKDLQTRVESL ESQTRQLELKAKNYADQISRLEERE AELKKEYNALHQRHTEMIHNYMEH LERTKLHQLSGSDQLEATAHSRIRK ERPISLGIFPLPAGDGLLTPDTQKGG ETPGSEQWKFQELSQPRSHTSLKVS HSPEPPKAVEQEDELSDISQGGSKA TTPASTANSDVSAIPDTPSKEDNEG FVKGTDTSNKSEISKHIEVQVAQETR NVSTESGENEEKSEVQAIIESTPELD MDKDLSGYKGSSTPTKGIENKAFDR NTESLFEELSSAGSGLIGDVDEGADL LGMGREVENLILENTQLLETKNALN VVKNDLIAKVDDELTCCKDVLQGELE AVKQAKLKLEDKNRELEEELRKARA EAEDARQKAKDDDDSDIPTAQRKRF TRVEMARVLMERNQYKERLMELQE AVRWTEMIRASRENPAEQEKKRSSI WQFFSRLFSSSNATKKPEPPVNLK YNAPTSHVTPSVKKRSSTLSQLPGD KSKAFDFLSEETEASLASRREQKRE QYRQVKAHVQKEDGRVQAFGWSLP QKYKQVANGQGETKMKNLPVPVYL RPLDEKDASMKLWCAVGVNLSGGK TRDGGSVVGASVIFYKDIAGLDTEGS KQRSASQSSLDKLDQELKEQQKEFK NQEELSSQVWICTSTHSTTKVIIDAV QPGNILDSFTVCNSHVLCIASVPGAR ETDYPAGEELSESGQVDKASLCGSM TSNSSAEMDSSLGGITVVGCSSTEGL TGAATSPSTNGASPVIEKPPMETEN SEVDENIPTAEEATEATEGNAGSTE DTVDISQPGVYTEHVFTDPLGVQIPE DLSPVFQSSNDSVYKQDISVLPNE QDLAREEAQMSSLLPTMWLGAQN GCLYVHSSVAQWRKCLHSIKLKDSIL SIVHVKGIVLVALADGTLAIFHRGVD GQWDLSNYHLLDLGRPHHSIRCMT VVHDKVWCGYRNKIYVVPKAMKIE KSFDAHPRKESQVRQLAWVGDGVW VSIRLDSTLRLYHAHTYQHLQDVDIE PYVSKMLGTGKLGFSFVRITALMVS CNRLWVGTGNGVIISIPLTETNKTSG TPGNRPGSVIRVYGDENS DKVTPGT FIPYCSMAHAQLCFHGRDAVKFFV

