

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q13469	NFAC2_HUMAN	Homo sapiens	Nuclear factor of activated T-cells, cytoplasmic 2	17.391142	T224;S225;S314;S333;S633;T733;S850	S23;S99;S107;S110;S148;S168;S171;S172;S174;S175;S177;S180;S213;S217;S221;S236;S243;S255;S268;S274;S276;S280;S326;S330;S363;S755;S757;S759;S856;S859	29351928;35289036;35132862	MNAPERQPPDGGDAPGHEPGGSP QDELDFSILFDYEYLNPNNEEPPNAH KVASPPSGPAYPDDVLDYGLKPYSP ASLSGEPGGRFGEPPDRVGPQKFLSA AKPAGASGLSPRIEITPSHELIQAVGP LRMRDAGLLVEQPPLAGVAASPRFT LPVPGFEGYREPLCLSPASSGSSASF ISDTFSPYTSPCVSPNNGPDDLCPQ FQNIHAHYSVRTSPIMSPRTSLAEDS CLGRHSPVPRPASRSSSPGAKRRHS CAEALVALPPGASPPQRSRSPSPQSS HVAPQDHGSPAGYPPVAGSAVIMDA LNSLATDSPCGIPPKMWKTSPPDSP VSAAPSKAGLPRHIYPAVEFLGPCEQ GERRNSAPESILLVPPTWPKPLVPAI PICSIPVTASLPPLEWPLSSQSGSYEL RIEVQPKPHHRAHYETEGSRGAVKA PTGGHPVVQLHGYMENKPLGLQIFI GTADERILKPHAFYQVHRITGKTVTT TSYEKIVGNTKVLEIPLEPKNNMRAT IDCAGILKLRNADIELRKGETDIGRK NTRVRLVFRVHIPESSGRIVSLQTAS NPIECQSRAHELPMVERQDTSCL VYGGQQMILTQGNFTSESKVVVTEK TTDGOQIWEWMEATVDKDKSQPNML FVEIPEYRNKHIRTPVKVNFYVINGK RKRSQPQHFTYHPVPAIKTEPTDEYD PTLICSPTHGGLGSQPYYPQHMPVA ESPSCLVATMAPCQQFRTGLSSPDA RYQQQNPAAVLYQRSKLSPSLLGY QQPALMAAPLSLADAHRSVLVHAGS QGQSSALLHPSPTNQQASPVIIHYSPT NQLRCGSHQEFQHIMYCENFAPG TTRPGPPPVSQQRSLSPGSYPTVIOQ QNATSQRAAKNGPPVSDQKEVLPAG VTIKQEQLDQTYLDDVNEIIRKEFS GPPARNQT

0