

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q60591	NFAC2_MOUSE	Mus musculus	Nuclear factor of activated T-cells, cytoplasmic 2	15.818779	S335	S23;S53;S54;S56;S99;S107;S110;S136;S150;S170;S173;S174;S176;S177;S179;S182;S215;S219;S223;S238;S245;S257;S270;S276;S278;S282;S328;S365;S757;S759;S761;S860	36852467	MDVPEPQDPDPDGGDGPHEPGGSP QDELDFSILFDYDYLNPPIEEPIAHK AISSPSGLAYPDDVLDYGLKPCNPLA SLSGEPPGRFGEPSDSIGFQNFLSPVK PAGASGSPRIEITPSHELMQAGGAL RGRDAGLSPEQPALALAGVAASPRF TLPVPGYEGYREPLCLSPASSGSSAS FISDTFSPYTSPCVSPNNAGPDDLCP QFQNI PAHYSPRTSPIMSPRTSLAED SCLGRHSPVPRPASRSSSPGAKRRH SCAEALVAPLPAASPQRSRSPSPQPS PHVALQDD SIPAGYPPTAGSAVLMD ALNTLATDSPCGIPSKIWKTS PDPTP VSTAPSKAGLARHIYPTVEFLGPCEQ EERRNSAPESILLVPPTWPKQLVPAI PICSIPVTASLPPELWPLSNQSGSYEL RIEVQPKPHHRAHYETEGSRGAVKA PTGGHPVVQLHGYMENKPLGLQIFI GTADERILKPHAFYQVHRITGKTVTT TSYEKIVGNTKVL EIPLEPKNNMRAT IDCAGILKLRNADIELRKGETDIGRK NTRVRLVFRVHVPEPSGRIVSLQAAS NPIECSQRS AHELPMVERQDM DSC LVYGGQQMILTGQNFTAESKVV FME KTTDGGQI WEMEATVDKDKSQPNM LFVEIPEYRNKHIRVPVKVNFYVING KRKRSQPQHFTYHPVPAIKTEPSDEY EPSLICSPAHGGLGSQPYYPQH PML AESPCLVATMAPCQQFRSGLSSPD ARYQQQSPAAALYQRSKSLSPGLLGY QQPSLLAAPLGLADAHRSVLVHAGS QGQGQGSTLPHTSSASQQASPVIHY SPTNQQLRGGGHQEFQHIMYCENF GPSSARPGPPPINQGQRLSPGAYPTV IQQQTAPSQRAAKNGPSDQKEALPT GTVKQEQNL DQTYLDDVNEIIRKE FSGPPSRNQT