

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9WV30	NFAT5_MOUSE	Mus musculus	Nuclear factor of activated T-cells 5	9.983052	NaN	S120;S134;T135;S155;S560	35822049	MPSDFISLLSADLDLESPKSLYSRES VYDLLPKELQLPPPRETSVASMSQTS GGEAGSPPPAVVAADASSAPSSSSM GGACSSFTTSSSPTIYSTSVTDSKAM QVESCSSAVGVSNRNVSEKQLTGNT VQQHPSTPKRHTVLYISPPEDLLDN SRMSCQDEGCGLESEQSCSMWME DPSNFSNMSTSSYNDNTEVPRKSR KRNPKQRPVKKRRDCEESNMDIFD ADSAKAPHYVLSQLTTDNKGNKAG NGTLDSQKGTGVKKSPLCGQYPV KSEGKELKIVVQPETQHRARYLTEGS RGSVKDRTOQGFPVTKLEGHNEPVV LQVFGNDSGRVKPHGFYQACRVTG RNTTPCKEVDIEGTTVIEVGLDPSNN MTLAVDCVILKLRNADVEARIGIAG SKKKSTRARLVFRVNITRKDGSTLTL QTPSSPILCTQPAGVPEILKSLHSCS VKGEEVFLIGKNFLKGTQVIFQENV SDENSWKSEAEIDMELFHQNHLIVK VPPYHDQHITLPVSVGIYVVTNAGRS HDVQPFYTPDPAAGALNVNVKKEI SSPARPCSFEEAMKAMKTTGCNVD KVTLPNALITPLISSMIKTEDVTPM EVTSEKRSSPIFQTTKSIGSTQQTLET ISNIAGGAPFSSPSSSSHLTPESENQ QQLQPKAYNPETLTTIQTQDISQPGT FPAVSAASQLPSSDALLQATQFQT REASRDTIQSDTVVNLSQLTEASQ QQQSPLQEQAQTLQQQIPSNIFSPS SVSQLQSTIQQLQAGSFTGSAAGGR SGSVDLVQQVLEAQQQLSSVLFSTP DGNENVQEQLNADIFQVSIQNSVS PGMFSSAESAVHTRPDNLLPGRADS VHQQTENTLSNQQQQQQQQQVM ESSAAMVMEMQQSICQAAAQIQSEL FPSAASASGSLQQSPVYQQPSHMMS ALPTNEDMQMQCELFSSPPAASGN ETSTTTTPQVATPGSTMFPQTPSSGD GEETGAQAKQIQNSVFQTMVQMQR SGDSQPQVNLFSSTKNIMSVQNG TQQQGNLFFQQGSEMMSLQSGNFL QQSSHSQAQLFHPQNPIADAQNLSQ ETQGSIFHSPNPIVHSQTSTASSEQL QPSMFHSQNTI AVLQGSVVPQDQQS PNIFLSQSSINNLQTNVAQEEQISF FAAQNSISPLQSTSNTEQQAQFQQQ PPISHIQTPILSQEQAQPSQQGLFQP QVALGSLPPNPMPQNQQGPIFQTR

