

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q12906	ILF3_HUMAN	Homo sapiens	Interleukin enhancer-binding factor 3	23.186007	S258;S382	S62;T188;S190;T315;S382;S384;S476;S477;S482;T592;S792;S810;S812;S816;S482;S482;S482	32522876;30379171;30397120;20305658;28510447;33465208;30444036;33214551;35083852;21740066;34019948;22121020;34725712;34229054;32119511;23301498	MRPMPRIFVNDHRHVMKHSVYPT QEELEAVQNMVSHTERALKAVSDWI DEQEKGSSEQAESDNMDVPPEDDS KEGAGEQKTEHMTRTLRGVMRVGL VAKGLLLKGDLDLELVLLCCEKPTTA LLDKVADNLAIQLAAVTEDEKYEILQS VDDAAIVIKNTKEPPLSLTIHLTSPVV REEMEKVLAGETLSVNDPPDVLDRQ KCLAALASLRHAKWFQARANGLKSC VIVIRVLRDLCTRVPTWGPLRGWPLE LLCEKSIGTANRPMGAGEALRRVLE CLASGIVMPDGSGIYDPCEKEATDAI GHLDRQQREDITQSAQHALRLAAFG QLHKVLGMDPLPSKMPKKPKNENP VDYTVQIPPSTTYAITPMKRPMEDG EEKSPSKKKKKIQKKEEKAEPQAM NALMRLNQLKPGLYKLVSTGPV HAPIFTMSVEVDGNSFEASGPSKKT AKLHVAVKVLQDMGLPTGAEGRDSS KGEDSAEETEAKPAVVAPAPVVEAVS TPSAAFPSDATAEQGPILTKHGKNPV MELNEKRRGLKYELISETGGSHDKR FVMEVEVDGQKFQAGSNKKVAKA YAALAALEKLPDTPALDANKKKRA PVPVRGGPKFAAKPHNPGFGMGGP MHNEVPPPPNLRGRGRGGSIRGRG RGRGFGGANHGGYMNAGAGYGSYG YGGNSATAGYSQFYSNGHSGNAS GGGGGGGGSSGYGSYYQGDNYNS PVPPKHAGKKOPHGGQKPSYGSY QSHQGOQQSYNQSPYSNYGPPQ GK QKGYNHGQGSYSYSNSYNSPGGGG GSDYNYESKFNYSGSGGRSGGNSYG SGGASYNPGSHGGYGGGSGGSSY QGKQGGYSQSNYNSPGSQNYSGP PSSYQSSQGGYGRNADHSMNYQYR