

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extr |
|--------------|------------|--------------|---------------------------------------|--------------|---------------|---|-------------------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|------|
| Q9Z1X4 | ILF3_MOUSE | Mus musculus | Interleukin enhancer-binding factor 3 | 27.878201 | NaN | T67;T188;S190;T315;S382;S384;S476;S477;S482;S486;T592;S794;S812;S814;S818 | 37453647;28528544 | MRPMPRIFVNDRRHVMMAKHSSVYPT QEELAVQNMVSHTERALKAVSDWI DEQEKGNSELSEAENMDTPPDDDES KEGAGEQKAEHMTRTLGRVVMRVGL VAKGLLLKGDLDLLELVLLCKEKPTTA LLDKVADNLAIQLTTVTEDKYEILQS VDDAAIVIKNTKEPPLSLTIHLTSPVV REEMEKVLAGEITLSVNDPPDVLDRO KCLAAASLRHAKWFQARANGLKSC VIVIRVLRDLCTRVPTWGPLRGWPLE LLCEKSIGTANRPMGAGEALRRVLE CLASGIVMPDGSIGYDPCEKEATDAI GHLDROQREDITQSAQHALLRAAFG QLHKVLGMDPLPSKMPKKPKNEENP VDYTVQIPPTTYAITPMKRPMEEDG EEKSPSKKKKIQKKEEKADPPQAM NALMRLNLKPLGQYKLISQTGPVH APIFTMSVEVDGNSNFEASGPSKKA KLHVAVKVLQDMGLPTGAEGRDSSK GEDSAEESDGKPAIVAPPVVEAVSN PSSVFPDATTEQGPILTKHGKPNV MELNEKRRGLKYEISETGGSHDKR FVMEVEVDGQKFGAGSNKKVAKA YAALAALEKLPDTPLALEANKKKRT PVPVRRGGPKFAAKPHNPGFGMGGP MHNEVPPPNIRGRGRGGNIRGRG RGRGFGGANHGGGYMNAGAGYGSY GYSSNSATAGYSQFYSNGGHSNAG GGSGGGGSSSYSSYQDYSNSP VPPKHAGKKPLHGGQKASYSSGYQ SHQGQQQPYNQSQYSSYGTQGGKQ KGYGHGQGSYSSYSNSYNSPGGGG GSDYSYDSKFNYSGGGRSGGNSY SSGSSSYNTGSHGGYGTSGGSSSY QKGQGGYSSQSNYSPPGSSQSYSGP ASSYQSSQGGYSRNTTEHSMNYQYR | True | False | 1.656 | 5.0 | 2.966 | 1.444 | 0.771 | 1.462 | 1.67 |