

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|---|--------------|---------------|---|----------------------------|---|
| Q0P678 | ZCH18_MOUSE | Mus musculus | Zinc finger CCCH domain-containing protein 18 | 17.391219 | T389;S838 | S6;S33;S45;S58;S64;S71;S75;S80;S92;T105;S106;S114;S169;S483;S528;S530;S532;S838;S847;S863;S888;S891 | 34887587;36852467;34418053 | MDVAESPELDPHSPEDDEEQPALSDD DILRESGSEQDLDGAGERASDLEEE ENATRVQSQEETRSDEEDRASEPKS QDQDSEAHELSRGPAGSPCEEEDD VEEDGTSDLRDEASSVTRELDEHEL DYDEEVPEEPAPAAQEEEEAEKAGAE EEEEKGEAPGEEGKPDVQSVGEQE PTEAAKEKKKEDDDGEIDDDGEIDDD DLEEGEVKDPSDRKVRPRPTCRFFM KGNCTWGMSCRFIHPGVNDKGNYS LITKAEPFPNGAPPLGPHPLMPANP WGGPVVDEILPPPEPPTESAWER GLRHAKVLLKATIRKEQEPDFEEK RFTVTIGEDDREFDKENEVFRDWNS RVPRDVRDITLEPYADPYDYEIERY WRGGQYENFRVQYEAEPYHNYRE RERERERENRQERERERERDRER ERRQERERERERERERDKERQRKE EWERERAKRDEKDRQHRDRDRDKD REKDKEPKPRSPQPPSRQAEPKK ESTSVGPOVKRADEWKDPWRRSRS PKKKLGVSVSPSRARRRRKTSASSAS ASNSSRSSRSRSGSGSSRSRSRS SSYSSYSSSRHSSFSGSRSRRSRFS SSSPSPPTPSPHRPPVRTKGEAPP PGKAGEKSIKKPAPPAPPQATKTTA PGPEPAKPGDLREARRKERQTRTPP RRRTLSGSGSGSSYSGSSRSRSL SVSSVSSVSSATSSSSSVHSVDSDD MYADLSPVSSASSRSPTPAQTKKE RGKSKKEDGVREEKRRRDPSAQPPK SSKAPAGGKASQQAAPQPAVPGQP QQGSFVAHKEIKLTLNKAADKGSR KRYEPSDKDRQSPPAKKANLSPDRG SRDRKSGGRMGSPKPERQRGQNAK APAAPADRKRPLSPQSKGSSKVTSVP GKATDTATAGTKSGKASTLSRRELL KQLKAVEDAIARKRAKIPGKV |