

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
H3BKP8	H3BKP8_MOUSE	Mus musculus	NaN	10.025162	NaN	NaN	34678516	MSVPEPPGEMERAAEEERPPPPTGEGNEEEVVVAAAARASGLVRRRSASSVDEEEEAASETTVVAGGGCKEQELTYELQQGYRILGEFLQEKHRGLTAPFLQPLGGVAAGEEEVAEGRRSGGRGSRVPPQQPSQGMCLLKMEEFSSGQYRGITEFVADFRLMLETQYRLHGVDHWISKQGQKLEMLLEQKLALLSRLREQERKEAEEACQKEVEEWERKLLAQAAPACMENMWEIPAIGHFLCLAQQILNLPEIVFYELERCLLMPQCNAFLSKIMTSLSPHRRPTLHRRPTLPYRTWEAVLRQKVQQWYTAVGQTENPDNCAEKLGLCPQFFKVLGEVNPLEERPFHELPHYQKVWLLKGLCDFVYETQKEVQDAVLGQPIHECREVILGYDYLENAYVHFPPQFCGADVRIYKQRPFPQAP EFPVPPIKVKRVPRIKLEKFKCDYAN TSNGEHRCTKEGLPLAFKKEQEIFDPACCP TKMNFNDHDTLEMEVKS NCDIKVHRPCEIEKTDCCKENLQKPRSPGEVTGFGEPLSPGEIRFIENQEK YGEASKIKTESNPLKENALKSCQIHI NGSHIDHPDINCHKVVRDILLEHSL QSHKKLKLTKMRAKKKKKKKKLKDVLNENLQRKREGLHSLPFKSYKPEIQNKLLIIKKKGKHKHKSGKKSISK KAITKKRKTVTKSPAVPEFQLICTNL DELRELITKIENELKDLENSRKKSGK WYHRRQAVKELHSTLIRLLNELLPW EPKLMKAFQRNRSRLKKDYDDFRR QPDHGQFTRELWATDECEGNPERE APKAEVSKSVDAEPLDTLEKEQEG SDDMKLSEIGFPMARSKLLKKEKLP KDIVKTLPKTLKRQSKQSSYLDDSTK ELSPRKKAKLSTNETSVENLEVNMQ IECLKESKPELPTPEFASKASVPVS TLQKGTKPIQALLAKNIGNKVTLTNQ LPPSTGRSVPAVEKPAISPEDTSPLK PALTCLTSTKGPLQMVYKMPGCGQW LPVDLHSSSVKIQMOPMVDSTKGEK IMQQVLILPKNFVIQHKEGKAVEKEI AAPQQKGPEHCSPGPQTSASC SLVS VPVTSVSTQLPNTVLSKTSTPSSNVS ARSQPLSPVASVSNALTSPVKTSQSE AGKVKSTASSTTLQPHTSPTISSTV QPLL PATTLESTDPGSSIPCFSQQT VDSSEAKQELKTV CIRDSQSILVRTR GGNTGVVKVQTNPEQNSPNSLSSSS VFTFTPQFQAFLVPKSTSCSASSQVA

GVTTSLLPSFSQASTSVSISCGFHP  
PMGKNLKSTQGQTLSSGYVGPMEK  
TSYMPSSPLKPSVSSSLLPSTTNSS  
VSVISISTGNFGQTNTNVIHTSTKPQ  
QVDCITKSYVTRSEATTAVNGDVLG  
ETPGQKLMLVSAPSGLPSGSVPSVN  
TAPEPTSAGVSTQKVFINAPVPGGA  
SSSAIVAESLRQSLPSPLNKTYIKNPE  
QPQIVLIPSTVGAPIKINSSPTVSQIKD  
VKIGLNIGQAIINPPGNLPAMSSINIL  
QSVMPKGEDKSSKSCISPISPNSNSA  
PASSNIVNHSLPAASESARTANTFSG  
IGASVPLSSLVASSPASAGTRPPVLV  
SGNDTSSRIMPVLSNRLCPSNLGNT  
VAISTVKTGHLASSVLISTTQPTVSPK  
CLTPALQIPVTVALPTPVTTSPKIINT  
VPQSATIPGATRPVLSKRQSQTSLQ  
FQSPGTTTIVPTNANTNKPPAELSPS  
ASPGKIVNISNVTSMPNQHMSPLV  
KSTLGNNNSIAGGSAIHTCSPPSNITS  
LAGAPFSEPCIQQKIVINTSTPLAPGT  
QIMMNGARFIVPPQGLGAGSHVLLI  
STNPKYGPPLVLNSGQSILATPVDNP  
VQKIIQTSNSSLGQPLKPSVRNSPK  
TVNSLGSPSSL SAVHTPPHIINTPAK  
VCVPPAPPAALTSVIKSSPATLLAKTS  
LVSAISSNPPLPSSTSVLHLDTSVK  
KLLVSPEGAILNTINTAAKVSSLPSP  
LPPVVSASQNPASVFPAPFPSSALEKP  
DTAAS