

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5SS00	ZDBF2_MOUSE	Mus musculus	DBF4-type zinc finger-containing protein 2 homolog	15.818779	T459	NaN	36852467	<p> MEAQSRYQEVMMKNNNGQHLFSSQHR SLTRQSRRTATNNTLMERFLQDVL RHHPYNYQDNRSAPNEPEAAAAAA AAADPGSPEVVVLDSDSDEKEDDTA DSGAERNSEDSGSVEEIDYRPGTSQ EHAEVAVRPSVIQKLERGQQSLEL AHKVESGVKKVNSVGVVQATTSGKK LVRPPVICNAPASSLPSGSFERPVAA NSVPRLVLAVASDSFPACDTENLETY FDSPDQGPSNPSSQPKTKDPKKKLSI NLDKLLAQRNLRKAGASFSPVVRVR ELTGSELCSVRAESSELEAGTAGNP RETDTLPEQAREGAIPKRREARSNT VRPQEETRLVLNKPTLLKQKRSVSS EERFSCGSRQAVPGPSQA AVRDL LSLLEEEVEEEKQEEEEEEQEEEEHE EEEEGVDQEDASYESRGSDFDC GSSCQSLSALSELTAREINVSEETHA YSQRNETPTVSGATSDNGSSSRQVI TQNIQLISLVDESYESSGSEVNFDDG DSLPSTSHRPPQVVEVVPLRLVDKS YGSGSSEPCEDSGSSADETPAASRQ QNPVRNTHANLVDENYGSSSFSSDS DAALDHPQVPVQEGSPRGRAVGQG NEEQPSSAEAHPERDGSLETVAHEL QRESQEINLPNQNTSLGDMNCES HGPEVGFHADAQLEADQSPVNPEE VLDLENQSVHSGISNLSFDSNASY QSANDQPQGAWEVNLDELNVDM EVKSNGCSSSELTFFSDSPLLSVTER SLLDFEGLNEDDFNLEDENCVSSSS DITFSDIPDDSVADQPQVAVYEEEP VGLENKSNESCIVSGITFSDIPLHSG NDHPEVAVKEVIKEDENVHLEGKN DNPSGSEICLDSNVPLHSVTNSDVA VKKINPPKEDQVQIEQIEQKENEPTD SELNLD CNSVNSKPGCEDPILRVS ETRLDSHVFPQSVIRKCEVVVKNVC LQKEKHAELTSKSTESHSEVSSASTA SHPVTEPYVGKKAKRKTKHLEEVNS DDEYGGSQPTFKFDVFPRTMTEKPQ PAALKEGHADPKDKITELRGMVNV NTADCLDSVLSQPQLASNENCVELK DTDGKPSDSKASADSTGHFHS LPKQ EFDVVSKMNEWKKEAKVLEQKISDL IYSKIIHDSNVSFRSAMDQLELALKQ ISLGNNDQVSLEDNSQDTCSETNLD SGFSVQAVVEPPEPEVTVLEPEHVE QEGRNNVPCDSEVSV DANGSVQLE </p>

AGQHSESGENRSQKDTDDTEGKRD
DAQFGGITCDSNVPQPLAGHIEVVQ
DIDHWKDHVDLEDKLGESKNSKVN
FHSDEPLQAVTNATQEPVEEINLPR
GRASPNNGCEPYGSTIVPVTNVIFC
SVIQKPQRLQKKCTSLKENSSNPCP
EVNVDSCDGPEVSVDSNDPCQSV
GHLQKPKEMNLKEDHIYLEDKSYK
LVDFEPTYDSDDPVQFVTVPSVEAVS
IKEVNLQKENPDDLENENFQPCCE
VACNSAVHLQSEADPPQVACKEADL
DKKALDIEDKGSVTCVPEVVYDSDVS
FQIVVNQVQTSGETDSPQVVFVDV
VSSDSDCDREVISDSNIPLQLEPPQM
TVKETS DINTDSLGSAANEKYYCKFC
GCDYEASQSVTNQSKESFKIINRKN
DYIILGDSTCPCSGHELNFNVDP
QPTTCQLQQPDRNFIDPEDKNFGSK
CPKRKLNWEDTAHPVTHQLQKTGE
ATSLRKDQKNRFRDRGWESSFA
GDHAAYAVSVIRQTALNSCGSELNF
QDDTFYHSDIDPPQPKKKGQKKT
FDLRVTKYEYPPNPMYGE GEEVAED
DLKEVVVHEASPOGQAPPSIVGKTCP
QGREDDVKTNTQACQGYFYSSYDGG
SETKKILLGEEKTIRSDLNQNTTSI
QHVKGKVGDTGDFSVDLKGQSCSLA
EGLHQHGVQVTSQNVQVEVRCGIQA
SSGKKRKITEQEEDSPKRKCFHHDS
QKKKKAQAGITELPEPQTKVLEPVQP
DSLVYIFSSLSMKEDQSLNPPKTKPG
SDSDLPHIYSCREHTSSGPRRRTVI
NPPQNLMPVEVGIDLNRHDPKPN
GDDSAKRQNLVSTSFAMPKKSVL
KFQRTNQSSFLKKS EDVGATQVPKD
NFQOTLVNRDGAKKFPKSVKEDLE
SQNQRKFWKKMVAANKLCLIKNA
YKTMVLRKSKLASEKLAIWIQLKAT
DIIRKYVSRCHGLMPRRHLSKTVLIR
MQLRKKKIVARKIKEAKRAAEALALK
LSRPSCPPRAPCPPRALCPPRAPCPP
RALCPPRAPCPPRALCTPRAPCPPRA
LCTPRAPCPPRALCTPRAPCPPRALR
APCPPRALCSPRAPCPPRALCPPRAP
CPPRAPRAPCPPRAPCPPRAPCLPRA
PCPPRAPCPPRAPCLPRAPCLPQAPC
PPRALCLPRVSCPPLPLCAPCPPCAP
VLPAGAEELSTTAGPAELPAHLSN
AGGIKRYRKTYFRRRKRLLPVREYDL
RSLSTTNTDRMVTRLASKSKSNEA
K