

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9UIF8	BAZ2B_HUMAN	Homo sapiens	Bromodomain adjacent to zinc finger domain protein 2B	13.682544	S9;S10;S19;S35	S1465;S1467;S1680;T2014;S2019	37217939;35254053;29485866;31492838;30379171	MESGERLPSSAASSTTPTSSSTPSVA SVVSKGGLSTGVASLSSTINPCGHLF RTAGDQPFNLSTVSSAFPMVSHPVF GLHSASSGHSEFGGLGTLGTPTALA AHPQLASFPGAEWWRRTDAHTRTG ATFFPPLLGIPLFAPPAQNHDSSSF HSRTSGKSNRNGPEKGVNGSINGS NTSSVIGINTSVLSTTASSSMGQTKS TSSGGNRKCNQEQSKNOPLDARV DKIKDKKPRKKAMESSNSDSDSGT SSDTSSEGISSSDSDLEEDDEEDQ SIEESEDSDSESEAQHKSNNQVL LHGISDPKADGQKATEKAQEKRIHQ PLPLASESQTHSFQSQKQPQVLSQ QLPFIFQSSQAKEESVNHKHTSVIQST GLVSNVKPLSLVNQAKKETYMKLIV PSPDVLKAGNKNTSESSLTSELRS KREYKQAFPSQLKKQESSKSLKKVI AALSNPKATSSSPAHPKQTLNHP NPFLTNALLGNHQPNGVIVQVIEA PLALTTKTKMQSKINENIAAASSTPF SSPVNLSTSGRRTPGNQTPVMPSAS PILHSQGEKAVSNVNPVKTQHHS HPAKSLVEQFRGTDSDIPSSKDSSEDS NEDEEEDDEEEDDEDEDESDS QSESDSNSESDTEGSEEDDDDKD QDESDDTEGEKTSMKLNKTTSSVK SPMSLTGHSTPRNLHIAKAPGSAP AALCSESQSPALGTSSSTLTSSPHS GTSKRRRVTDERELRIPLEYGWQRE TRIRNFGGRLQGEVAYYAPCGKCLR QYPEVIKYL SRNGIMDISRDNFSFSA KIRVGDFYEARDGPQGMQWCLLKE EDVIPRIRAMEGRRGRPPNPDRQRA REESRMRRRKGRPPNVGNAEFLDN ADAKLLRKLQAQEIARQAAQIKLLRK LQKQEQARVAKEAKKQQAIAAEEK RKQKEQIKIMKQEQEKIKRIQQIRMEK ELRAQQILEAKKKKKEEAANAKLLE AEKRIKEKEMRRQQAVLLKHQERER RRQHMLMKAMEARKKAEEKERL KQEKRDEKRLNKERKLEQRRLELE MAKELKKPNEDMCLADQKPLPELP RIPGLVLSGSTFSDCLMVVQFLRNF GKVLGFDVNIDVPNLSVLQEGLLNI GDSMGEVQDLLVRLLSAAVCDPGLI TGYKAKTALGEHLLNVGVNRDNVSE ILQIFMEAHCGQTELTESLKTAFQA HTPAQKASVLAFLINELACSKSVVSE IDKNIDYMSNLRDRKVVVEGKLRKL RIIHAKKTGKRDTSGGIDLGEQHP GTPTPGRKRRRGGDSYDDDDDD DSDDQGEDDEDEEDKEDKKGKKT DICEDEDEGDQAASVEELEKQIEKLS KQSQYRRKLFDASHSLRSVMFGQ DRYRRRYWILPQCGGIFVEGMESGE GLEEIAKEREKLKKAESVQIKEEMFE

										TSGDSLNCNNTDHCEQKEDLKEKD NTNLFLOKPGSFSKLSKLEVAKMP PESEVMTPKPNAGANGCTLSYQNSG KHSLGSVQSTATQSNVEKADSNNLF NTGSSGPGKFYSPLPNDQLLKLTE KNRQWFSLPRTPCDDTSLTHADM STASLVTPQSQPPSKSPSPTAPLGS SAQNPVGLNPFALSPLOVKGGVSM MGLQFCGWPTGVVTSNIPFTSSVPS LGSGGLGLENGNSFLTNSVASSKS ESPVPQNEKATSAQPAAVEVAKPVD FPSPKPIPEEMQFGWRIIDPEDLKA LLKVLHLRGIREKALQKQIQKHLDYI TQACLKNKDVAIHELNENEENQVTR DIVENWSVEEQAMEMDLSVLQQVE DLERRVASASLQVKGWMCPEPASE REDLVYFEHKSFTKLCKEHDGFTG EDESSAHALERKSDNPLDIAVTRLA DLERNIERRIEDIAPGLRVRRALS EARSAAQVALCQQLOKSIWEKSIM KVYQCICRKGDNELLNLCDCGCDKG CHTYCHRPKITTPDGDWFCPACIAK ASGQTLKIKKLHVKGKKTNESKKGK KVTLTGDEDEDSASTSSSLKRGNK DLKRRMEENTSINLSKQESFTSVK KPKRDDSKDLALCSMILTEMETHED AWPFLLPVNLKLVPGYKKVKKPMD FSTIREKLSSGQYPNLETFALDVRLV FDNCETFNEDDSDIGRAGHNMRKY FEKKWTDTFKVS
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