

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8BRB7	KAT6B_MOUSE	Mus musculus	Histone acetyltransferase KAT6B	17.689198	S1730;T1737	S356;S866	34678516;36852467	MVKLANPLYTEWILEAVQKIKKQKQ RPSEERICHAVSTSHGLDKKTVSEQL ELSVQDGSVLKVTNKGGLASYKDPDN PGRFSSVKPGTFPKPTKGSKGPPCN DLRNVDWNKLLKRAIEGLEEPPNGSS LKNIEKYLRSQSDLTGTTNHPAFQQ RLRLGAKRAVNNGRLLKEGPQYRVN SGSSDGKGAPQYPSAFPSSLPPVSLL PHEKDQPRADPIPICSFCLGTKESNR EKKPEELLSCADCGSSGHPSCLKFC PELTANVKALRWQCIECKTCSACRV QGKNADNMLFCDSCDRGFHMECC DPPLSRMPKGMWICQVCRPKKKGR KLLHEKAAQIKRRYAKPIGRPKNKLK QRLLSVTSDEGMSAFTGRGSPGRG QKTKVSTTPSSGHAASGKHSSSRLA VTDPTRPGATTKTTTSSTYISASTLKV NKKTKGLIDGLTKFFTPSPDGRRSRG EIIDFSKHYPKRVKVSQKQSCSHVL ATDTDIKISIKQESADVSLVGNKELV TEEDLDVFKQAQELSWEKIECESGV EDCGRYPSVIEFGKYEIQTWYSSPYP QEYARLPKLYLCEFCLKYMKSKNILL RHSKCKGWFHPPANEIYRRKDLSVF EVDGNMSKIYCQNLCLLAKFLDHK TLYYDVEPFLFYVLTKNDEKGCHLV GYFSKEKLCQQKYNVSCIMIMPQHQ RQGFGRFLIDFSYLLSRREGQAGSPE KPLSDLGRLSYLAYWKSIVILEYLYRH HERHISIKAISRATGMCPHDIATTLQ HLHMIDRRDGRFVIIRREKLILGHM EKLNCSRPNELDPESLRWTPMLIS NAVVSEEEERAEKEAERLMEQASC WEKEEQEILSSRVSSRQSSAKVQSK NKYLHSPERRPVAGERGQLELSKE SSEEEEEEEEEDEEEEEEEEEESIQ TSPRLTKPQSVSIKRKRPFVVKKKR GRKRRRINSSVTETISETTEVLNEP FDNSDEERPMPQLEPTCEIPVEEGG RKPVLRKAFPHQPGKKRQTEEEEGE DNHFFKTAALCRKDVEDDDAEHLKE GSKDNPEPLKCRQVWPKGAKRGLS KWKQSKERKTGFKLNLTPPETPME PEDQVTIEEQKELSEDKGSVPGMER EVTETVDALLPQEGSRREETGIPVSP HKSPGGKVDEEDLIRGEEEGEEEGE EEGEREEQEEEEVTTEKDLDGAKS KENPEPEISMEKEDPVHLGDHEEDE DEEEEP SHNEDHDADDEDDGHME

								AANMERGDLPRETFKDALEGQEAF DLSIQPSHSNPEVLMNCGVDLTMSC NSEPKELAGDTGTAPESDAEPPEEQ TQKQDQKNSDGVDAELEEGGPAAV EIDSETAQAVQSLTQENREHDDTFP DCAETQEACRSLQNYHTDQSPQIA TTLDECQQSDHSSPVSSVHSHPGQS VRSVNSPSVPALENSYAQISPDQTAI TVPPLQNMETSPMMDVPSVSDHSQ QVVDSGFSDLGSIESTTENYENPSSY DSTMGGSICGNGSSQNSCSYSSLTS SNLTQNSCAVTQQMSNISGSCSML QQTSSISPPTCSVKSPQGCVVERPPS SSQQLAQCSMAANFTPPMQLADIPE TSNANIGLYERMGQSDFGAGHYPPQ SATFSLAKLQQLTNTLIDHSLPYSHS AAVTSYANSASLSTPLSNTGLVQLSQ SPHSVPGGPQAQATMTPPPNLTPPP MNLPPPLLQRNMAASNIGISHSQRL QTQIASKGHVSMRTKAASLSPAAAT HQSQIYGRSQTVMQGPARTLTMQ RGMNMSVNLMPAPAYNVNSVNMN MNTLNAMNGYSMSQPMMNSGYHS NHGYMNQTPQYPMQMOMGMMGS QPYAQQPMQTPPHANMMYTAPGH HGYMNTGMSKQSLNGSYMRR
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