

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7LBC6	KDM3B_HUMAN	Homo sapiens	Lysine-specific demethylase 3B	19.076303	T366;S381;S386;T409;S420;T425;T426;S428;S429;T430;T433;S437;S465;S469;S470;S480;S647;S650;S651;S654;S657;S659;S660;S661;T667;S668;S679;T696	S492;S546;S556;S560;T614;S766;S773;S778;S779;S798;S1253;S1259	32119511;34725712;35132862;23301498;22661428;28510447;30059200;33214551;35289036;29237092;29351928;30379171;34019948;35254053;37340703;34931806;35138101;30620550;31492838;32574038	MADAAASPVGKRLLLLFDATAASAS ASAPAAAAASGDPGALRTRAWRAG TVRAMSGAVPQDLAIFVEFDGCNWK QHSWVKVHAAEEVIVLLLEGLVWAP REDPVLLQIGRIVSIAQWPALTTFTPLV DKLGLGSSVVPVEYLLDRELRFLSDA NGLHLFQMGTDSQNQILLEHAALR ETVNALISDQKLQEIFSRGPYSVQGH RVKIYQPEGEEGWLYGVVSHQDSIT RLMEVSVTESGEIKSVDPRLIHVML MDNSAPQSEGGTLKAVKSSKGGKKK RESIEGKDGRRRKSASDSGCDPASK KLKGDGRGEVDSNGSDGGEASRGPW KGGNASGEPGLDQRAKQPPSTFVPO INRNIRFATYTKENGRTLVVQDEPVG GDTPASFTPYSTATGQTPLAPEVGGGA ENKEAGKTLQVGGQIVASAAVVT ASSTPNTVRISDTGLAAGTVPEKQKG SRSQASGENSRNSILASSGFGAPLPS SSQPLTFGSGRSQSNGLVATENKPL GFSFGCSSAQEAQKDDLSKNLFFQ CMSQTLPTSNYFTTVSESLADSSSS RDSFKQSLESLSGLCKGRSVLGTDT TKPGSKAGSSVDRKVPAESMPTLTP AFPRSLLNARTPENHENLFLQPPKL SREEPSNPFLAFVKEVHSPFSSFAS QASGSSSSATTVTSTKVPASWPESH SADSASLAKKKPLFITDSSKLVSGV LGSALTSGGPSLSAMGNRSSSPTS SLTQPIEMPTLSSSPTEERPTVGGPQ QDNPLLKTFSNVFGRHSGGFLSSPA DFSQENKAPFEAVKRFLDERSLAC RQDSDSSTNSDLSLDSEEQLQAK TGLKGIPEHLMGKLGPNGERSAELL LGKSKGKQAPKGRPRTAPLKVGQSV LKDVSKVKKLQSGEPFLQDGCIN VAPHLHKCRECLERYRKFKEQEED DSTVACRFFHFRRLIFTRKGVLRVEG FLSPQQSDPDAMNLWIPSSSLAEGI DLETSKYILANVGDQFCQLVMSEKE AMMMVEPHQKVAWKRAVRGVREM CDVCETTLFNIHWVCRKCGFGVCLD CYRLRKSRRPSETEEMGDDEEVFSWL KCAKQGSHEPENLMPTQIIPGTALY NIGDMVHAARGKWGIKANCPCISRO NKSVLRPAVTNGMSQLPSINPSASS GNETTFSGGGGPAPVTTPEPDHVPK ADSTDIRSEEPLKTDSSASNSNSELK AIRPPCPDTAPPSSALHWLADLATQK AKEETKEAGSLRSVLNKESSHSPFGL DSFNSTAKVSPLTPKLFNSLLLGPTA SNNKTEGSSLRDLLHSGPGKLPQTP LDTGIPFPVVFSTSSAGVKSASLPN FLDHIIASVVENKKTSDASKRACNLT DTQKEVKEMVMGLNVLDPHTSHSW LCDGRLCLHDPNSKNNWKFIFREC WKQGQPVLVSGVHKKLKSELWKPE AFSQEFGDQVDLVNCRNCAIISDV KVRDFWDGFEIICKRLRSEDGQPMV LKLKDWPPGEDFRDMMPTRFEDLM ENLPLPEYTKRDGRNLASRLPSYFV RPDLGPKMYNAYGLITAEDRRVGTT NLHLDVDSDAVNMVYVGIPIGEGAH

