

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A3KFM7	CHD6_MOUSE	Mus musculus	Chromodomain-helicase-DNA-binding protein 6	15.818779	T1715	S1865	36852467	MKMKIQKKEKQLSKLRALNHSPMS DASVNFYDKSPSPFDCSPDQGENIE EAANHCLPHKNLYTTEEEADTLFSR KLTSHNGMEDSGGRGTGVKKRKK KEPGEQEGTKGSKDREP KPRKREP KEPKEPRRAKEPKRAKEPKETKQKD GVKKPRKHREASGTKEGKEKRSDT YGSRTKSKKASREQGTPVERKKKG KRKNETTVESELDHSLPNPSLQSP EEPSESADSQRRSGRQVKRRKYNE DLDFKVDDGETIAVLGAGRTSALS ASTLAWQAEPPEDDANIIEKILASK TVQEVHPGEPFDLELFYVKYRNFS YLHCKWATMEELEKDPRIAQKIKRF RNKQAQMKHIFTEPDEDLFPDYIE IDRILEVAHTKDAETGEEVTHYLVKW CSLPYEESTWELEEDVDPKVKEFE SLQILPEVKHVERPASDAWQKLETS REYRNSNRLREYQLEGMNWLLFN WYNRKNCILADEMGLGKTIQSIAFL SEIFVRGIHGPFLLIAPLSTITNWERE FRTWTEMNAIVYHGSQISRQMIQQY EMVYRDAQGNPLSGVFKFHVITTF EMILADCPKLIHWSCVIIDEAHLR KNRNCKLLEGLKMALEHKVLLTGT PLQNSVEELFSLNLFLEPSQFPSETA FLEEFGLKTEEQVKLQKILKPMML LRLKDDVEKNLAPKQETIIEVELTN IQKKYYRAILEKNFSFLTGANQHN MPNLINTMMELRKCCNHPYLINGA EEKILED FRKAHSSEASDFQLQAMI QAAGKLVLIDKLLPKLIAGGHKVLIFS QMVRCLELDYLIQRRYTYERIDGR VRGNLRQAADRFCCKPDSDFVFL CTRAGGLGINLTAADTCIIFDSDWNP QNDLQAQARCHRIGQSKAVKVYRLI TRNSYEREMFDKASLKLGLDKAILQ DINRKGSTNGVQQLSKMEVEDLLRK GAYGALMDEEEDGSKFCEEDIDQIL QRRTHITIQSEGKGFSAKASFVAS GNRTDISLDDPNFWQKWAKIAELDT EANNEKESLVIDRPRVRKQTKHYNS FEDELMEFSELDSDSERPTRSR LSDRARRYLRAECFRVEKNLLIFGW GRWKDILTHGRFKWPLNEKDMEMI CRALLVYCVKHYKGDEKIKSFIWELI TPSKDGQVQTLQNHSGLSAPVPRGR KGKKTKNQLLLPELKNADWLATCNP EVLHDDGYKHLKQHCNKVLLRV

RMLYYLKAIEILGEAADKAFEGTPARE
LDVLLPDIDYVEIPVDWWDAAEADKS
LLIGVFKHGYERYNAMRADPALCFL
EKVGMPEKSLSAEQGVTDGTS DIP
ERGNIDKEDSAEDKLDGLQKQTASP
SDGSDGIFGEKKDDSQAAQDGSDDP
KSPWPVSSALTARLRLVTTYQR CNR
KELCRPEILGPGNQGYWVQEEVFRR
TSEMDLINKEAQKRWTRREQADFYR
TVSSFGVVYDQEKKAFDWTQFRIISR
LDKKSDESLEHYFYSFVAMCRNVCR
LPAWKDDGPPDASIYVEPITEERA AK
TLYRIELLRKVREQVLMCPQLHERL
QLCRPSLYLPVWWECKHDRDLLIG
TAKHGLNRTDYYIMNDPQLSFLDAY
RNYAQHKRTDTQAPGSLCCLYQSNS
KLYESLTYTPVSRTSESLESEPENLM
RMESRDDHLCLPEGGLPDITCENFV
SKVQEVISLDHDENLLPESLENMIY
GKTGLSQEPHSFQEAPTTNTQSRKN
TITISASRNESCQPPGIEAETSASSL
MSSLEAGVAKMNIKNGKHLVLSISK
EGEPCCSETGRRPETIGHREAKCLV
SPTLDTGHESGFVDLCSLSVYDPKR
NFSSDQQLIDLLENKSLESKLILNQS
DEEEENEDETLAIVASATEKPEVLD
FPKPTVNIPRGKNLSFHQDEAKKGR
LEVVSKTAGPQRVFPFPPANQCHCKH
IERWAHGLGSEDSEVEKPKAYQPD
YRSKANNSTVEGETAVIPTEPFKLKH
ELLKEPWKESSEGGKSFMYAPEGS
EPKPEDMDFENKDDYEKDGTCLSQ
DYPGKYSEEEKSSASGIAGDLGEEA
QEVRAPTIAQLLQEKTLYSFSEWPKD
RVIINRLDNICHVVLKGWPCSHQY
EPGALPTPVLSSSAGSRSSLSEPEA
TEHGFSNGAALAAQIQKESFLAPVFT
KDEQKHRRPYEFEVERDAKARSLEE
YSATHGRPPIVLNGWHGESIDLSC
SSEGPSGATSPFPVSASTPKIGAISL
QGALGMDLSGILQAGLIHPVTGQIV
NGSLRRDDAAMRRRRRGRRKHIEGG
MDLIFLKEQTLQAGILEVHEDAGQT
TLSTHPEVPGATSSAPEPTAAASSQ
AEKAVPSKSLLDWLRQQADYSLDVP
GFGTSFSDKPKQRRPRCKEPGKLDIS
SLGGEERVPAVPKEPGLRGFLPESKF
NHTLAEPVLRDAGPRRRRGRPRNEL
LKAPAIVADSPSGMGPLFMNGLIAG
MDLVGLQNVRNIPGIPLTGLVGFPA
GFATMPTGEEVKNTLSMLPMLPG
MAAVPQMFGVGGLLNTPMATTCTT

