

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitoch
Q9HCK8	CHD8_HUMAN	Homo sapiens	Chromodomain-helicase-DNA-binding protein 8	12.224462	S113;S117;S123;T171;T184;S256;S2519;T2522;T2523	S432;S553;S562;S1420;S1424;S1976;S1978;T1993;S1995;S2008;S2046;T2051;S2069;S2071;S2182;S2200;S2202;T2204;S2211;T2215;S2223;S2519	30379171;38665916;35132862;33214551;37340703;35138101;34019948;35254053;32119511;29351928	MADPIMDLFDDPNLFGDLSLTDSSFNQVTDPIEALGLPSSLDLQDMNQDGGGGDVGNGSSASELVPPPEETAPTELSKESTAPAPESITLHDDYTTQPASQEQAQPVLTSTPTSGLLQVSKSQEILSQGNPFMVGVSATAVSSSSAGGQPPQSAPKIVLKAPPSSSVTGAHVAVQIQAGGITSTAQPLVAGTANGKVTFTKVLGTPLRPGVSVSGNTVLAQVPGNQAQVQIRVQPSRPVKQLVLPVKGSAPAGNPGATGPPLKPAVLTSTPTQGSKRITLVLPQPSGGPQGRHVVLGSLPGKIVLQGNQLAALTQAKNAQGQPAKVVTIQLQVQQPQQKIQIVPQPPSSQPOQPQPPSTQPVTLSVQQAQIMGPGQSPGQRLSVPVKVVLQPOAGSSQGASSGLSVKVLASAEVAALSSPASSAPHSGGKTGMEENRRLEHQKQKQKANRIVAEIARARARGEQNIPRVLNEDELPSVRPEEEGKRRRKSAGERLKEEKPKKSTSGASKTKGKSKLNTITPVVCKKRRNTSSDSDVEVMPAQSPREDEESSIQRRSNRQVKRKYTEDLDIKITDDEEEEDVVTGPIKPEPILPEPVQEPDGETLPSMQFFVENPSEEDAAIVDKVLSMRIVKKELPISQYTEAEFFVKYKNYSYLHCEWATISQLEKDKRIHQKLRKFKTKMAQMRHFFHEDEEFPNPDYVEVDRILDSEHSIDKDNQEPVIYLVKWCSLPYEDSTWELKEDVDEGKIREFKRIQSRHPELKRVNRPOASAWKLELSHEYKRNRLREYQLEGVNWLLFNWYNRQNCILADEMGLGKTIQSIQIAFLQEVYVGIHGFVIAPLSTITNWEREFNTWTEMTIVYHGLASRQMIQYEMYCKDSRGRIPGAYKFDALITTFEMILSDCPELREIEWRCVIIDEAHLKRNCKLLDSLKHMDLEHKVLLTGTPLQNTVEELFSLHFLFEPSPQFSESEFLKDFDGLKTEEQVQKLAALKPMMLRRLKEDVEKNLAPKQETIIEVELTNIQKYYRAILEKNFSFLSKGAGHTNMPNLLNTMMELRKCCNHPYLINGAEKILTEFREACHIIPDFHLQAMVRSAGKLVLDKLLPKLKAGGHKVLIFSQMVRCLDILEDYLIQRRYLYERIDGRVGRNLRQAADRFSKPDSDRFVLLCTRAGGLGINLTAADTCIIFDSWNPQNDLQAQARCHRIGQSKAVKVYRLITRNSYEREMFDKASLKLGLDKAVLQSMSGRDGNITGIQFQSKKEIEDLRRKGAYAAIMEEDDEGSKFCEEDIDQILLRRTTTITIESEKGSTFAKASFVASENRTDISLDDPNFWOKWAKKADLMDLLNSKNNLVIDTPRVKQTRHFSTLKDDDLVEFSDL ESEDDERPRSRRDRHHAYGRDTCFRVEKHLVYVWGRWRDILSHGRFKRRMTERDVETICRAILVYCLLHYRGDENIKGFIWDLISPAENGKTKELQNHSGLSIPVPRGRKGGKVKVQSTFDIHKADWIRKYNPDTLFDQESYKHLKHQCNKVLRLVRMLYLRQEVIGDQAEKVLGGAIASEIDIWFVVDQLEVPITWWDSEADKSLIGVFKHGYEKYNTMRADPALCFLEKAGRPDDKAAIAEHRVLDNFSDIVEGVDFDKCEDPEYKPLQGPDKQDDEGDPLMMMDDEISVIDGDEAQTQOPGHLFWPPGSALTARLRLVTAYQRSYKREQMKEAAERGDRRRRCEAAFKLKEIARREKQQRWTRREQTDYRVVSTFGVEYDPTMQFHWDRFRTFARLDKKTDESITKYFHGFVAMCRQVCRLLPPAAGDEPPDPNLFIPIEERASRTLYRIELRLREQLVCHPLLEDRLALCQPPGPELPKWVEPVRHDGELLRGAARHGVSQTD CNIMQDPDFSLAARMNMYQNHQAGAPAPSLSRCSTPLHQQYTSRTASPLPLRPDAPVEKSPETATQVPSLES	None	None	None	None	None

								LTLKLEHEVVARSRPTQDYEMRVS PSDTPPLVSRVPPVKLEDDSDSE LDLSKLSPPSSSSSSSSSSSTDES EDEKEEKLTDOSRSKLYDEESLLSLT MSQDGFNEDGEQMTPELLLQER ORASEWPKDRVLINRIDLVCQAVLS GKWPSRRSQEMVTGGILGPGNHL LDSPSLTPGEYGDSPVPTPRSSAAS MAEEEASAVSTAAQFTKLRGMDE KEFTVQIKDEEGLKLTQKHKLMAN GVMGDGHPLFHKKGNRKKLVELE VECMEEP NHLVDLETRIPVINKVD GTLVGEDAPRAELEMWLGHP FAVDPRFLAYMEDRRKQKWQRCKK NNKAELNCLGMEPVQTANSRNGKK GHHTETVFNRLPGPIAPESKKRA RRMRPDL SKMMALMQGGSTGSLSL HNTFOHSSGLQSVSSLGHSSATSA SLPFMPFVMGGAPSPHVDSSTML HHHHHPHPHHHHHPGLRAPG YPSSPVTASGTTLRLPLQPEEDDD EDEEDDDLSQGYDSSERDFSLIDD PMMPANSDESSEDADD				
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