

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence
Q14839	CHD4_HUMAN	Homo sapiens	Chromodomain-helicase-DNA-binding protein 4	22.983049	NaN	S44;S303;S308;S309;S310;S319;T367;S428;S515;T517;T529;S531;T703;S1209;S1308;S1349;S1370;S1531;S1535;S1537;T1542;T1549;T1553;S1570;S1576;S1602;T1653;T1679	3626323;33214551;34229054;27231347;32522876;34019948;23301498;28510447	MASGLGSPSPCSAGSEEDMDALL NNSLPPHPENEEEDPEEDLSETETP KLKKKKKPKKPRDPKIPKSKRQKKE RMLLCRQLGDSSGEGPEFVEEEEEV ALRSDSEGSdyTPGKKKKKGLGPKK EKKSkskrkeEEEEEDDDDDSKPEK SSAQLLEDWGMEDIDHVFSEEDYRT LTNYKAFSQFVRPLIAAKNPKIAVSK MMMVLGAKWREFSTNNPFKGSSG ASVAAAAAAVAVVESMVTATEVAP PPPPVEVPIRAKTKEGKGNARRKP KGSPrVPDAKKPKKkVAPLkIKLGG FGSKRKRSSSEDDDLVESDFDDAS INSYSVSDGSTSRSSRSRKLRTTK KKKGEEEVTAVDGYETDHQDYCEVC QQGGEIILCDTCPRAYHMVCLDPDM EKAPEGKWSCPHEKEGIQWEAKE DNSEGEEIIEVGGDLEEDDHHM EFCRVCKDGGELLCDCPSSYHIH CLNPPLPEIPNGEWLCPRCTCPALK GKVQKILIWKGWQPPSPTVPRPPD ADPNTPSPKPLEGRPERQFFVKWQG MSYWHCSWVSELQLELHCQVMFR NYQRKNDMDEPPSGDFGGDEEKSR KRKNKDPKFAEMEERFYRYGIKPEW MMIHRILNHSVDKKGHVHYLIKWR DLPYDQASWESEDVEIQDYDLFKQS YWNHRELMRGEEGRPGKkLkVkl RKLERPPETPTVDPTVKYERQPEYLD ATGGTLHPYQMEGLNWLRFswAQG TDTILADEMGLGKTVQTAVFLYSLYK EGHskGPFLVSAPLSTIINWEREFE MwAPDMYVVTYVGDkDSRAIrene FSFEDNAIRGGKkASRMkKEASVKF HVLLTSYELITIDMAILGSIDWACLIV DEAHLKNNQSKFFRVLNGYSLQH KLLLTGTPLQNNLEELFHLLNFLT ERFHNLEGFLEEFADIAKEDQIKKL HDMLGPHMLRRLKADVFKNMPSKT ELIVRVELSPMQKKYKYILTRNFEA LNARGGGNQVSLNVMMDLkKCCN HPYLPVAAMEAPKMPNGMYDGSA LIRASGKLLLQKMLKnlKEGGHRV LIFSQMTKMLDLLEDfLEHEGYKYE RIDGGITGNMRQEAIDRFNAPGAQQ FCFLSTRAGGLGINLATADTVIIYDS DWNPHNDIQAFSRAHRIGQNKKVM IYRFVTRASVEERITQVAKKMMMLT HLVVRPGLGSKTGSMSKQELDDILK FGTEELFKDEATDGGGDNKEGEDSS VIHYDDKAIERLLDRNQDETEDTEL QGMNEYLSSFKVAQYVVREEMGE EEEVEREIIKQEEsvDPDYWEKLLRH HYEQQEDLARNLGKkGRIRKQVNY NDGSQEDRDWQDDQSDNQSDYSV

								ASEEGDEDFDERSEAPRRPSRKGLR NDKDKPLPPLLARVGGNIEVLGFNA RQRKAFNLNMRGMPQDAFTTQ WLVRDLRGKSEKEFKAYVSLFMRHL CEPGADGAETFADGVPREGLSRQHV LTRIGVMSLIRKKVQEFEHVNGRWS MPELAEVEENKKMSQPGSPSPKTPT PSTPGDTQPNTAPVPPAEDGIKIEE NSLKEEESIEGEKEVKSTAPETAIEC TQAPAPASEDEKVVVEPPEGEKVE KAEVKERTEEPMETEPKGAADVEKV EEKSAIDLTPIVVEDKEEKKEEKK EVMLQNGETPKDLNDEKQKKNIKQ RFMFNIADGGFTELHSLWQNEERA ATVTKKTYEIWHRRHDYWLLAGIIN HGYARWQDIQNDPRYAILNEPFKGE MNRGNFLEIKNKFLARRFKLLEQAL VIEEQLRRAAYLNMSQDPSPHSMAL NTRFAEVECLAESHQHSKESMAG NKPANAVLHKVLKQLELLSDMKA DVTRLPATIARIPPVAVRLQMSERNI LSRLANRAPEPTPQQAQQ
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