

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O14647	CHD2_HUMAN	Homo sapiens	Chromodomain-helicase-DNA-binding protein 2	11.981591	S1075;S1076	S207;S208;T240;S242;S1085;S1087;S1365;S1386;S1807	30620550	MMRNKDKSQEEDSSLHSNASSHSA SEEASGSDSGSQSESEQSDPGSGH GSESNSSSESSESQSESESEAGSKS QPVLPEAKEKPKASKKERIADVKKMW EYYPDVYGVRRSNRSRQEPSRFNIK EEASSGSESGSPKRRGQRQLKKQEK WKQEPSEDEQEQTSAESEPEQKK VKARRPVPRRTVPKPRVKKQPKTQR GKRKKQDSSDEDDDDDEAPKRQTR RRAAKNVSYKEDDDDFETDSDDLIE TGEGVDEQQDNSETIEKVLDSRLGK KGATGASTTVYAIEANGDPSGDFDTE KDEGEIQYLIKWKGWSYIHSTWESE ESLQQQKVGLKKLENFKKKEDEIK QWLGKVSPEDVEYFNCQOELASEL NKQYQIVERVIAVKTSKSTLGQTD AHSRKPAPSNEPEYLCKWMGLPYSE CSWEDEALIGKQNCIDSFHSRNN SKTIPTRECKALKQRPRFVALKKQPA YLGGENLELRDYQLEGLNWLASHW CKNNSVILADEMGLGKTIQTISFLSY LFHQHQLYGPFLIVVPLSTLTSWQR EFEIWAPEINVVVYIGDLMSRNTIRE YEWIHSQTKRLKFNALITTYEILLKD KTVLGSINWAF LGVDEAHRLKNDD LLYKTLIDFKSNHRLITGTPLQNSL KELWSLLHFIMPEKFEFWEDFEED HGKGRENGYQSLHKVLEPFLRRVK KDVEKSLPAKVEQILRVEMSALQKQ YYKWILTRNYKALAKGTRGSTSGFL NIVMELKCCNHCYLIKPPREENERE NGQEILLSLIRSSGKLILLDKLLRLR ERGNRVLIFSQMVRMLDILAEYLTIK HYPFQRLDGSIKGEIRKQALDHFNA DGSEDFCFLLSTRAGGLINLASADT VVIFDSDWNPQNDLQAQARAHRIQ KKQVNIYRLVTKGTVEEIIERAKKK MVL DHLVIQRM DTTGRTILENNSGR SNSNPFNKEELTAILKFGAEDLFKEL EGEESEPQEMDIDEILRLAETRENE VSTSATDELLSQFKVANFATMEDEE ELEERPHKDWDEIIPPEQRKKVEEE ERQKELEEIYMLPRIRSSTKKAQTND SDSDTESKRQAQRSSASESETEDSD DDKKPKRRGRPRSVRKDLVEGFTDA EIRRFIKAYKKFGLPLERLECIARDAE LVDKSVADLKRLGELIHNCSV SAMQ EYEEQLKENASEGKGPGKRRGPTIKI SGVQVNVKSIIQHEEEFEMLHKSIPV

							DPEEKKKYCLTCRVKAAHFDVEWGV EDDSRLLLGIYEHGYGNWELIKTDP ELKLTDKILPVETDKKPKQKQLQTRA DYLLKLLRKGLEKKGAVTGGEEAKL KKRKPRVKKENKVPRLKEEHGIELS SPRHSDNPSEEGEVKDDGLEKSPM KKKQKKKENKENKEKQMSSRKDKE GDKERKSKDKKEKPKSGDAKSSSK SKRSQGPVHITAGSEPVPIGEDEDD LDQETF SICKERMRPVKKALKQLDK PDKGLNVQEQLEHTRNCLLKIGDRI AECLKAYS DQEHKLRWRNLWIFVS KFTEFDARKLHKLYKMAHKKRSQEE EEQKKKDDVTGGKKPFRPEASGSSR DSLISQSHTSHNLHPQKPHLPASHG PQM HGHPRDNYNHPNKRHFSNAD RGDWQRERKFNYGGGNNNPPWGS DRHHQYEQHWYKDHHYGDRRHMD AHRGSYRPNNMSRKRPYDQYSSD RDHRGHRDYDRHHHDSKRRRSDE FRPQNYHQDFRRMSDHRPAMGY HGQGPSDHYSFHTDKLGEYKQPLP PLHPAVSDPRSPPSQKSPHDSKSPL DHRSPLESLQKNNPDYNWNVRK T
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