

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A0A0C4DFX4	A0A0C4DFX4_HUMAN	Homo sapiens	Helicase SRCAP	14.559543	T2229;S2239	NaN	35254053;31492838;30620550	VSDGMTGSNPVSPASSSSPASSGAG GISPQHIAQDSSLDGPPGPPDGATVP LEGFSLSQAADLANKGPKWEKSHAE IAEQAKHEAEIETRIAELRKEGFWSL KRLPKVPEPPRPKGHWYDLCEEMQ WLSADFAQERRWKRGVARKVVRMV IRHHEEQRQKEERARREEQAKLRRI ASTMAKDVRQFWSNVEKVVQFKQQ SRLEEKRKALDLHLDFIVGQTEKYS DLLSQLNQPLTSSKAGSSPCLGSSS AASSPPPPASRLDDEGDGDFQPQEDE EEDDEETIEVEEQQEGNDAEAQRRE IELLRREGELPLELLRSLPQLLEG PSSPSQTPSSHSDTRDGPPEGAAE EPPQVLEIKPPPSAVTQRNKQPWHP DEDDEEFTANEEEADEEDTIAAEE QLEGEVDHAMELSELAREGELSME ELLQQYAGAYAPGSGSSEDEDEDEV DANSSDCEPEGVVEAEPPQEDSSS QSDSVEDRSEDEEDEHSEEEETSGS SASEESESESEDAQSQSQADEEEE DDDFGVEYLLARDEEQSEADAGSGP PTPGPTTLGPKKEITDIAAAAESLQPK GYTLATTQVKTIPILLLGRQLREYQH IGLDWLVTMYEKKLNGILADEMGLG KTIQTISLLAHLACEKGNWGPPLIIV PTSVMLNWEMELKRWCPSPFKILTY GAQKERKLRQGWTKPNAFHVCITS YKLVLDHQAFRRKNWRYLILDEAQ NIKNFKSQRWQSLNFNNSQRRLLT GTPLNQSLMELWLSLMHFLMPHVQ SHREFKEWFSNPLTGMIEGSQEYNE GLVKRLHKVLRPFLRRVKVDVEKQ MPKKYEHVIRCRLSKRQRCLYDDFM AQTTTKETLATGHFMSVINILMQLR KVCNHPNLFDRPVTSPFITPGICFS TASLVL RATDVHPLQRIDMGRFDLIG LEGRVSRYEADTF LPRHLSRRV LLE VATAPDPPPRPKPVKMKNRMLQPV PKQEGRTVVVVNNPRAPLGPVVRP PPGPELSAQPTPGVPQVLPASLMVS ASPAGPPLIPASRPPGPVLLPPLQPN SGSLPQAGEVVSIGQLASLAQRPVAN AGGSKPLTFQIQGNKLTLTGAQVRQ LAVGQPRPLQMPPTMVNNTGVVKIV VRQAPRDGLTPVPLAPAPRPPSSGL PAVLNPRPTLTPGRLPTPTLGTARAP MPTPTLVRPLLKLVHSPSPEVSASAP GAAPLTISSPLHVPSSLPGPASSPMPI PNSSPLASPVSSTVSVPLSSSLPISVP

TTLPAPASAPLTIPISAPLTVSASGPAL  
LTSVTPPLAPVVPAAPGPPSLAPSGA  
SPSASALTLGLATAPSLSSSQTPGHP  
LLLAPTSSHVPGLNSTVAPACSPVLV  
PASALASPFPSAPNPAPAQASLLAPA  
SSASQALATPLAPMAAPQTAILAPSP  
APPLAPLPVLAPSPGAAPVLASSQTP  
VPVMAPSSTPGTSLASASPVPAPTPV  
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MVSRLPVSKDEPDTLTLRSGPPSPPS  
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MTRMLDVLEQFLTYHGHLYLRDGDG  
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MKFLEASLEEVSREELKQAEQVEA  
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GDESSCGTGGGTHRRSKKAKAPERP  
GTRVSERLRGARAETQGANHTPVIS  
AHQTRSTTTPRCSPARERVPRPAPR  
PRPTASAPAAIPALVPVPSAPVPIS  
APNPITILPVHILPSPPPPSQIPPCSSP  
ACTPPPACTPPPAHTPPPAQTCLVTP  
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AELCAQALASPESLELASVASETSS  
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EEKELVRRRRQQRGAASTLVPGVSE  
TSASPGSPSVRSMSPGESSPPIGGPC

EAAPSSSLPTPPQQPFIARRHIELGVT  
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RPPKKNRSPADAGRGVDEAPSSTLK  
GKTNGADPVPGPETLIVADPVLEPQL  
IPGPQLGPQVHRPNLLSPVEKRR  
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AVSHRGRKAKT