

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6ZRS2	SRCAP_HUMAN	Homo sapiens	Helicase SRCAP	5.800368	S196;S1124;S1125;S1126;S1186;T2406;T2412;S2416;T2420	S1172	31492838;34725712;34019948;30620550;34846842;35254053;35132862;35289036;30379171;33214551	MQSSPSPAHPQLPVLQTMVSDGM TGSNPVSPASSSSPASSGAGGISPQH IAQDSSLDGPPGPPDGATVPLEGFSL SQAADLANKGPKWEKSHAEIAEQAK HEAEIETRIAE LRKEGFWSLKRLPKV PEPPRPKGHWDYLCEEMQWLSADF AQERRWKRGVARKVVRMVIRHHEE QRQKEERARREEQAKLRRIASTMAK DVRQFWSNVEKVVFQKQSRLEEK RKKALDLHLD FIVGQTEKYS DLLSQS LNQPLTSSKAGSSPCLGSSSAASSPP PPASRLDDEDGDFQPQDEEEDDEE TIEVEEQQEGNDAEAQRREIELLRRE GELPLEELLRSLPPQLLEGSPSSPQT PSSHSDTRDGPPEGAE EEPQVLEI KPPPSAVTQRNKQPWHPDEDDEEF TANEEEA EDEEDTIAAEEQLEGEVD HAMELSELAREGELSMEELLQOYA GAYAPGSGSSEDEDEDEV DANSSDC EPEGPVEAE EPPQEDSSSQSDSVED RSEDEE DEHSEEEETSGSSASEESE SESEDAQSQSQADEEEDDDDFGVE YLLARDEEQSEADAGSGPPTPGPTTL GPKKEITDIAAAAE SLQPKGYTLATT QVKTP IPI LLLRGQLREYQHIGLDWL TMYEKKLN GILADEMGLGKTIQTISL LAHLACEKGNWGP HLIIVPTSVMLN WEMELKRWCP SFKILTYGAQKERK LKRQGWTKPNAFHCITSYKLVLQD HQA FRRNWRYLILDEA QNIKNFKS QRWQSL LNFN SQRRLLLTGTPLQNS LMELWSLMHFLMPHV FQSHREFKE WFSNPLTGMIEGSQEYNEGLVKRLH KVL RPFLLRRVKVDVEKQMPKKYEH VIRCRLSKRQRCLYDDFMAQTTTKE TLATGHFM SVINILMQLRKVCNHPN LFDPRPVTSPFITPGICFSTASLV LRA TDVHPLQRIDMGRFDLIGLEGRVSR YEADTF LPRHRLSRRV LLEVATAPDP PPRPKPVKMKVNRMLQPVPKQEGR TVVVVNNPRAPLGPVVRPPPGPELS AQPTPGVPVQVLPASLMVSASPAGPP LIPASRPPGPVLLPPLQPN SGLPQV LPSPLGVL SGT SRPPTPLSLKPTPPA PVRLSPAPPPGSSSLLKPLTVPPGYTF PPAAATTT TTTATATTTAVPAPTAP QRLILSPDMQARLP SGEVVSIGQLAS LAQRPVANAGGSKPLTFQIQGNKLT LTGAQVRQLAVGQPRPLQRNVVHLV SAGGQHHLISQPAHVALIQAVAPTPG PTPVSVLPSSTPSTTPAPTGLSLPLAA NQVPPTMVNNTGVVKIVVRQAPRDG LTPVPLAPAPRPPSSGLPAVLNPRP TLTPGRLPTPLGTARAPMPTPLVR PLLKLVHSPSPEVSASAPGAAPLTISS PLHVPSSLPGPASSMPPIPNSPLAS

PV SSTV SVPLSSSLPISVPTTL P APAS
APLTIPISAPLTVSASGPALLTSVTPPL
APVVPAAPGPPSLAPSGASPSASALT
LGLATAPSLSSSQTPGHPLLLAPTSS
HVPGLNSTVAPACSPVLVPASALASP
FPSAPNPAPAQASLLAPASSASQALA
TPLAPMAAPQTAILAPSPAPPLAPLPV
LAPSPGAAPVLASSQTPVPVMAPSST
PGTSLASASPVPAPT PVLAPSSTQTM
LPAPVPSPLPSPASTQTLALAPALAPT
LGGSSPSQTL SLGTGNPQGFPTQT
LSLTPASSLVPTPAQTL SLAPGPPLGP
TQTL SLAPAPPLAPASVGPAPAHTL
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PTLGAAAQTLALAPASTQSPASQAS
SLVVSASGAAPLPVTMVSRLPVSKDE
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GRLVTVVEEKELVRRRRRQQRGAAST
LVPGVSETSASPGSPSVRSMSPGESS
PPIGGPCEAAPSSSLPTPPQPF IARR
HIELGVTGGGSPENGDGALLAITPPA
VKRRRGRPPKKNRSPADAGRGVDEA
PSSTLKGKTNGADVPVGPETLIVADP
VLEPQLIPGPQLGQP VHRPNPLLS
PVEKRRRGRPPKARDLPIPGTISSAG
DGNSESRTPPPHPSLTLPLPLLVC

