

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9C0D2	CE295_HUMAN	Homo sapiens	Centrosomal protein of 295 kDa	11.234068	S1099;S1593;S1616;S1829;T2128	S14;S654;S938;S1637;T2473	35254053;29351928;37217939;30379171	MKRKVVNTHKLR LSPNEEFILKED YERRRKLRL LQVREQERDIALQIRED IKQRNQQFTR LAEELRAEWEEESQT KIQNLEKLYL ASLRSMGEGHRQAK ENEPDLDALA QRAAERKRKADLRHK EALKVQKNQ KEILLKQKTWHIKARK EALLVEKERS AKITSLPPPPPTLFENI EVKRISAVK TNSSTYHHLHTFVNRE TDTKRPDARL AEEEEAKRLEELQKQ AAQERMERFE KAHVRFQAMKKIH LAQNQEKLMK ELKQLQ QEDLARRR QTVAQMPPQ LVELPYKRSEM KEDW QRELEFAFED MYNADRKVKGNLILH LEPEPLPTV TNQIQDEELDLSMEQE NLGAAEDLPV TEAEICSSSETDVPLVM KTQQIPSKV LFKKLLNKIRSQKSLWT IKSMSSEDE SEMITTVSEIESKAPTVE SGTIASKERT LSSGQEQVVESDTLTI ESGPLASEDK PLSCGTNSGKEQEI ETLPITTVAQ SSVLLHPQEAARIRM SARQKQIMEI EEQKQKQLELLEQIEQ QKLRLETDCF RAQLEEEKRKTQPT GVGIAPASCP VISDEDESHRQMIRNYQ HQLLQONRL HRQSVETARKQLE QTMMLKGRCP SVSAPSLITDSVISVPS WKSERPTAISE HWDQGQRLKLSPN KYQPIQPIQT SKLEQDHFQVARQNH FPQRQVETTE TLRASDILTNOALESQ EHLRQFSQTET QQRDYKLVPKDSET LSRALSHDRQL ISQDARKISETFGAT TFQSLESQQL FSENSENISYHLTEPS SFVPLVPQH SFSSLPVKVESGKIQEP FSAMSKSTV STSHSIISQMHDRPLLP SENITAAQGN MKALQEQLDLQKKVL QATQEAQEQL LCKQKEVEQQTGLS VFLPLVTPD SSALLPSAKADLGRIQE SSPTKNNIAV SSDHHVISQLQDKRLS LSQPILSQON NFKFLQEQLNIQKDS LQARREAQEV LYVHKQSELDRRVC EQAEPSFPFQ VAQHTFTSLPSADTKS GKIQEQHSSK SEKGLVSCQSDIPISQ DGSLSFLQQ FLPLHDSLKLLQEQLT KQRDTLQAR HEAQVELLHRQRDL GDSKSGLV SSSSPVVVQHSVASQA SAKAEPRIQE LYLSEKENVGP SCHL IIPTFQDKS LFPQHSLAQQENLTI QEQSQIQRV ILGAKEGTQEFVHTESE LEKRISSEQT GTSSSLSQVDESERFQ ECISIKSDSTI PLSHPKIPRCQERLLR VSQHMLPLQ DNLEEHQAWLDTEKE AFHFSQKTQ ENTSSSEQTGSSSFIPQL VQLSFTSLA SAESGTILEPLFTESESK IFSSHLQIP QLDRLLRISQLIQPQD NLKALQEQL ATQREAILARQEAREE LLLHQSEWE GRISPEQVDTSSLPLVP QHSFASLPL NESERNQEP CSINSDN IVSSGHSEI PTLPDGLLGLSHLVPQ QDNLIALEE HLAQTDFLPSIEKTQK ELVLSKPC KFEKVSSEHFIQSHHG DLQALQQQLD TQKKAIRSIQEVQEEL

LLQRLSELEKRVSSQVCSFVSQV
PVADSSERTQKSFPKSNLTPSSHR
EIPRLQDRLLSLSKPLPQQDNMTAQ
LDAQREVMYSYKPEEELSLNKQRK
LNKSESAEHTIPSLFLPKETEHSFIPL
PFAEAKPKSTCELYSSQNEHAAPPS
NPVIFGFQDRLLSFSQSVLTQQDNL
GLQKQLDLQREVLHYSQKAQEKLLV
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ISKPTVENDLKTQKMGQLRDWFPNT
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ASEDTSAKQSGEHLEKDLGRRSSKP
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QSFFGSPLAHDPFSCQLVQGENVC
GDDYDEAVKLKESVVENHAVLSYAV
EEEHAYLGPTVKPDDKAKTLYEPLS
SATVSTGSLSYENTDLSLTDPEFS
EHMDDSKQESTTSKEEETNISSIVP
STQDIYQRQNSDVHKSLLPAVDET
TCGHTHFQMQMIDKYINEANLIPEKT
DLQELEHIFPNLHHQLFKPLEPHPD
FDLSSSSGISPDNRDFYQRSDSSSE
SHCATGLSKSTVYFTALRRSMHSS
LNTSPNQPDNLAVHGAHSFATE
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SECSTKHQLESRKESMGFEELSKRG
VVTMLQSQGLIEDNKNETCRVLDIN
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SISSSVSRKGVNKVRASFPEDRKTT
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KTKQEAQAQRARAKEFHKKTLEKL
RAKNTC