

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96N67	DOCK7_HUMAN	Homo sapiens	Dedicator of cytokinesis protein 7	11.340476	S190	S30;S180;S182;T450;S452;S862;S864;S882;S888;S896;S900;S905;T907;T909;S910;S929;S964;S1383;S1430;S1432;S1434;S1438;S2129;S1421;S1425;S1429;S1390;S1394;S1398;S1390;S1394;S1398;S1421;S1425;S1429	30379171;28657654;32119511	MAERRAFAQKISRTVAAEVRKQISG QYSGSPQLLKNLNLVGNISHHTTVPL TEAVDPVDLEDYLITHPLAVDSGPLR DLIEFPPDDIEVVYSPRDCRTLVS AVP EESEMDPHVRDCIRSYTEDWAIVIRK YHKLGTGFNPNTLDKQKERQKGLPK QVFESDEAPDGN SYQDDQDDLKRR SMSIDDTPRGSWACSFIDLKNSLPD ALLPNLLDRTPNEEIDRQNDQORKS NRHKELFALHPSPEDEEPIERLSVPD IPKEHFGQRLLVKCLSLKFEIEIEPIF ASLALYDVKEKKKISENFYFDLNSEQ MKGLLRPHVPPAAITTLARSAIFSITY PSQDVFLVIKLEKVLQQGDIGECAEP YMIFKEADATKNKEKLEKLSQADQ FCQRLGKYRMPFAWTAIHLMNIVSS AGSLERDSTEVEISTGERKGSWSER RNSSIVGRRSLERTTSGDDACNLT SF RPATLTVTNFFKQEGDRLSDEDLYK FLADMRRPSSVLRRLRPITAQLKIDIS PAPENPHYCLTPELLQVKLYPDSRVR PTREILEFPARDVYVPNTTYRNLLYIY PQSLNFANRQGSARNITVKVQFMYG EDPSNAMPVIFGKSSCSEFSKEAYTA VVYHNRS PDFHEEIKVKLPATLTDH HHLLFTFYHVSCQKQNTPLETPVG YTWIPMLQNGRLKTGQFCLPVSLEK PPQAYSVLSPEVPLPGMKWVDN HK GVFNVEVVAVSSIHTQDPYLDKFFAL VNALDEHLFPVRIGDMRIMENNLE NELKSSISALNSSQLEPVVRFHLHLL DKLILLVIRPPVIAGQIVNLGQASFEA MASIINRLHKNLEGNHDQHGRNSL LASYIHYVFRLPNTYPN SSSPGPGGL GGSVHYATMARSAVRPASLNLNRSR SLSNSNPDISGTPTSPDDEVRSIIGSK GLDRSNSWVNTGGPKAAPWGSNPS PSAESTQAMDRSCNRMS SHTTETSSF LQTLTGRLPTKKLFHEELALQWVVC SGSVRESALQQA WFFFELMVKSMV HHLYFNDKLEAPRKS RFPERFMDDI AALVSTIASDIVSRFQK DTEMVERLN TSLAFFLNDLLSVM DRGFVFSLIKSC YKQVSSKLYSLPNPSVLVSLRLDFLR IICSHEHYVTLNLPC SLLTPPASPSPS VSSATSQSSGFSTNVQDQKIANMFE LSVPFRQQHYLAGLV LTELAVILDPD AEGFLFGLHKKVINMVHNLSSHDS DPRYSDPQIKARVAMLYLPLIGIIMET VPQLYDFTETHNQ RGRPIC IATDDYE SESGSMISQTVAMA IAGTSVPQLTRP GSFLLTSTSGRQH TTFSAESSRLLI CLLWVLKNADET V LQKWFTDLSVLQ

LNRLDLLLYLCVSCFEYKGGKVFER  
MNSLTFKKSMDRAKLEEAIGSIG  
ARQEMVRRSRGQLGTYTIASPPERS  
PSGSAFGSQENLRWRKDMTHWRQ  
NTEKLDKSRAEIEHEALIDGNLATEA  
NLIILDITLEIVVQTVSVTESKESILGG  
VLKVLHSMACNQSAYVLQHC FAT  
QRALVSKFPELLFEEETEQCADLCL  
RLLRHCS SIGTIRSHASASLYLLMR  
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GTSQNFNEEFLRRSLKTILTYAEDL  
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VKMKEHQEDPEMLIDL MYRIAKGYQ  
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EAAQCLVHSAALVAEYLSMLEDRKY  
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SPDEEGICSGKYFTESGLVGLLEQAA  
ASF SMAGMYEAVNEVYKVLPIHEA  
NRDAKKLSTIHGKLQEA FSKIVHQST  
GWERMFGTYFRVGFYGTKFGDLDE  
QEFVYKEPAITKLAEISHRLEGFYGE  
RFGEDVVEVIKDSNPVDKCKLDPNK  
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EQFKRKTILTTS HAFYIKTRVNVTH  
KEEILTPIEVAIEDMQKKTQELAFAT  
HQDPADPKMLQMV LQGSVGT TVNQ  
GPLEVAQVFLSEIPSDPKLFRHHNKL  
RLCFKDFTKRCEDALRKNKSLIGPD  
QKEYQRELERNYHRLKEALQPLINR  
KIPQLYKAVLPVTCHRDSFSRMSLR  
KMDL