

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96N67	DOCK7_HUMAN	Homo sapiens	Dedicator of cytokinesis protein 7	10.933704	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;35254053;28657654;34725712;32119511	MAERRAFAQKISRTVAAEVRKQISG QYSGSPQLLKNLNVGNISHHTTVPL TEAVDPVDLEDYLITHPLAVDSGPLR DLIEFFPDDIEVVYSPRDCRTLVS AVP EESEMDPHVRDCIRSYTEDWAI VIRK YHKLGTGFNPNTLDKQKERQKGLPK QVFESDEAPDGN SYQDDQDDLKRR SMSIDDTPRGSWACSFIDLKNSLPD ALLPNLLDRTPNEEIDRQNDQQRKS NRHKELFALHPS PDEEEPIERLSVPD IPKEHFGQRLLVKCLSLKFEIEIEPIF ASLALYDVKEKKKISENFYFDLNSEQ MKGLLRPHVPPAAITTLARSAIFSITY PSQDVFLVIKLEKVLQQGDIGECAEP YMIFKEADATKNKEKLEKLSQADQ FCQRLGKYRMPFAWTAIHL MNIVSS AGSLERDSTEVEISTGERKGSWSER RNSSIVGRRSLERTTSGDDACNLT SF RPATLTVTNFFKQEGDRLSDEDLYK FLADMRRPSSVLRRLRPITAQLKIDIS PAPENPHYCLTPELLQVKLYPDSRVR PTREILEFPARDVYVPNTTYRNLLYIY PQSLNFANRQGSARNITVKVQFMYG EDPSNAMPVIFGKSSCSEFSKEAYTA VVYHNRS PDFHEEIKVKLPATLTDH HHLLFTFYHVSCQKQNTPLETPVG YTWIPMLQNGRLKTGFCLPVSLEK PPQAYSVLSPEVPLPGMKWVDNHK GVFNVEVVAVSSIHTQDPYLDKFFAL VNALDEHLFPVRIGDMRIMENNLE NELKSSISALNSSQLEPVVRFHLHLL DKLILLVIRPPVIAGQVNLGQASFEA MASIINRLHKNLEGNHDQHGRNSL LASYIHYVFRLPNTYPNSSSPGPGGL GGSVHYATMARS AVRPASLNLNRSR SLSNSNPDISGTPTSPDDEVRSIIGSK GLDRNSWVNTGGPKAAPWGSNPS PSAESTQAMDRSCNRMS SHTTETSSF LQTLTGRLPTKKLFHEELALQWVVC SGSVRESALQAWFFFELMVKSMV HHLYFNDKLEAPRKS RFFPERFMDDI AALVSTIASDIVSRFQKDTMVERLN TSLAFFLNDLLSVMDRGFVFSLIKSC YKQVSSKLYSLPNPSVLVSLRLDFLR IICSHEHYVTLNLP CSLTTPASPSPS VSSATSQSSGFSTNVQDQKIANMFE LSVPFRQQHYLAGLV LTELAVILDPD AEGFLGFLHKKVINMVHNLSSHDS DPRYSDPQIKARVAMLYLPLIGIIMET VPQLYDFTETHNQRGRPIC IATDDYE SESGSMISQTVAMA IAGTSVPQLTRP GSFLLTSTSGRQHTT FSAESSRSLLI CLLWVLKNADETVLQKWFTDLSVLQ LNRLDLLLYLCVSCFEYK GKKVFER

MNSLTFKKSMDRAKLEEAIGSIG
ARQEMVRRSRGQLGTYTIASPPERS
PSGSAFGSQENLRWRKDMTHWRQ
NTEKLDKSRAEIEHEALIDGNLATEA
NLILDTLEIVVQTVSVTESKESILGG
VLKVLLHSMACNQSAVYLQHC FAT
QRALVSKFPELLFEEETEQCADLCL
RLLRHCS SIGTIRSHASASLYLLMR
QNFEIGNNFARVKMQVTMSLSSLV
GTSQNFNEEFLLRRSLKILTYAEDL
ELRETTFPDQVQDLVFNLMILSDT
VKMKEHQEDPEMLIDL MYRIAKGYQ
TSPDLRLTWLQNMAGKHSERSNHA
EAAQCLVHSAALVAEYLSMLED RKY
LPVGC VTFQNISSNVLEESAVSDDVV
SPDEEGICSGKYFTESGLVGLLEQAA
ASF SMAGMYEAVNEVYKVLPIHEA
NRDAKKLSTIHGKLQEA FSKIVHQST
GWERMFGTYFRVGFYGT KFGDLDE
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HQDPADPKMLQMV LQGSVGT TVNQ
GPLEVAQVFLSEIPSDPKLFRHHNKL
RLCFKDFTKRCEDALRKNKSLIGPD
QKEYQRELERNYHRLKEALQPLINR
KIPQLYKAVLPVTCHRDSFSRMSLR
KMDL