

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96HP0	DOCK6_HUMAN	Homo sapiens	Dedicator of cytokinesis protein 6	8.666114	S185	S178;S872;S880;S884;S1308;S2036	35254053	MAASERRAFAHKINRTVAAEVRKQV SRERSGSPHSSRRCSSSLGVPLTEVV EPLDFEDVLLSRPPDAEPGPLRDLVE FPADDLELLLQPRECRTTEPGIPKDE KLDAQVRAAVEMYIEDWVIVHRRYQ YLSAAYSPVTTDTQERERQKGLPRQVF EQDASGDERSGPEDSNDSRRGSGSP EDTPRSSGASSIFDLRNLAADSLPS LLERAAPEDVDRRNETLRRQHRPPA LLTLYPAPDEDEAVERCSRPEPPREH FGQRILVKCLSLKFEIEIEPIFGILALY DVREKKKISENFYFDLNSDSMKGLL RAHGTHPAISTLARS AIFS VTYPSPDI FLVIKLEKVLQOGDISECCEPYMVLK EVD TAKNKEKLEKLR LA AEQFCTRL GRYRMPFAWTAVHLANIVSSAGQLD RDS DSEGERRPAWTD RRRRRGPQDR ASSGD DACFSGFRPATLTVTNFFK QEAERLSDEDLFKFLADMRRPSSLL RRLRPVTAQLKIDISPAPENPHFCLS PELLHIKYPDPGRPTKEILEFPARE VYAPHTSYRNLLYVYPHSLNFSSRQ GSVRNLAVRVQYMTGEDPSQALPVI FGKSSCSEFTREAFTPVYHKNKSPE FYEEFKLHLPACVTENHLLLFTFYH VSCQPRPGTALETPVGFTWIPLLQH GRLRTGPFCLPVSVDQPPPSYSVLT DVALPGMRWVDGHKGVFSVELTAV SSVHPQDPYLDKFFTLVHVLEEGAF PFRLKDTV LSEGNVEQELRASLAAL RLASPEPLVAFSHHVLDKLVRLVIRP PIISGQIVNLGRGA FEAMAHV VSLVH RSLEAAQDARGHCPQLAAYVHYAFR LPGTEPSLPDGAPPVTVQAATLARGS GRPASLYLARSKSISSSNPDLAVAPG SVDDEVSRI LASKLLHEELALQWVV SSSAVREAILQHAWFFFQLMVKSMA LHLLLGQR LDTPRKLRFPGRFLDDIT ALVGSVGLVITRVHKDVELAEHLN ASLAFFLSDLLSLVDRGFVFSLVRAH YKQVATRLQSSPNPAALLTLRMEFT RILCSHEHYVTNLNLPCCPLSPPASPS PSVSSTTSQSSTFSSQAPDPKVTSMF ELSGPFRQQHFLAGLLLTELALALEP EAEGAFLLHKKAISAVHSLLCGHDT DPRYAEATVKARVAELYLPLLSIARD TLPRLHDFAE GPGQRSRLASMLDSD TEGEGDIAGTINPSVAMAIAGGPLAP GSRASISQGPPTASRAGCALSAESSR

TLLACVLWVLKNTPEALLQRWATDL  
TLPQLGRLLDLLYLCLAAFEYKGGKA  
FERINSLTFKKSMDMKARLEEAILGT  
IGARQEMVRRSRERSPFGNPENVR  
WRKSVTHWKQTSRVDKTKDEME  
HEALVEGNLATEASLVVLDTLEIVQ  
TVMLSEARESVLGAVLKVVLYSLGSA  
QSALFLQHGLATQRALVSKFPELLFE  
EDTELCADLCLRLLRHCGSRISTIRT  
HASASLYLLMRQNFEIGHNFARVK  
MQVTMSLSSLVGTTONFSEEHLRRS  
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VNEVYKNLIPILEAHRDYKKLAAVHG  
KLQEAFTKIMHQSSGWERVFGTYFR  
VGFYGAHFGDLDEQEFVYKEPSITKL  
AEISHRLEEFYTERFGDDVVEIHKDS  
NPVDKSKLDSQKAYIQITYVEPYFDT  
YELKDRVITYFDRNYGLRFLCTPFT  
PDGRAHGELPEQHKRKTLSTDHAF  
PYIKTRIRVCHREETVLTPEVAIED  
MQKKTRELAFAEQDPPDAKMLQM  
VLQGSVGPTVNQGPLEVAQVFLAEIP  
EDPKLFRHHNKLRLCFKDFCKKCE  
DALRKNKALIGPDQKEYHRELERNY  
CRLREALQPLLTQRLPQLMAPTPPG  
LRNSLNRASFRKADL