

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q66JQ7	KNL1_MOUSE	Mus musculus	Kinetochorescaffold 1	15.818779	S523	S32;S60;S538;T540;S794;S878;S1243;S1464	36852467	MDGVYSEANEENDNTQRPVRRQHS SILKPPRSPLQDLKCGNQTNQEPNP PRKRKSSRRVSFADTIKVFQTESHM KTERNSEISGMNTLLCAPIQTQMQQ KEFSITDCNHERKHANDQTVIFSDE NQMDLTASHTVMITKGLSDCTKNE NSTKIDTTSFLENLKHHAANSRIKK DLACSTVLSQNFSEKINSDFIKR LKTGKHISSTELDKENAEIPVYSKD SNSASSTYQMHASLGVDENSSNRT RIFREQDDGMNLTQCHTACIKTWIP PSTEAKIGEFKGDKTIYGNCEMELTT NYTIQVLSSENNLSERETQTQNGM NVTTVDGATAPAPEKKTALKDKLNA AFQGSFPNPENKIHKCHPIESETH TVTQISSQSASTLAVTSKICSSPAIE GYKTIFHSSNDAMELTKCLSAMEE EKLLKADDKYSKICTNPDAGPLREK TIYLEEDSMDITKSHTVAIDNKIFKH DQENIKKEIIAIPFEKEMVLRNLMP MSKDEKRDVNYISVPQVSKEQLORS QTNTLSVSLTDKKMEFLADEDMDLT KSHTTKLSQVIPTTFDLASKNVTKSY SHSKSPLNEWESLDKQVVLGQHSK LPLPQRKDRDDPDCSHHKIMYSEEL QTMDLTKSHTIVIGFGPSEVQEHSKI NLEHKNSQLTAESIQTAVNVPAANS RVVTTNDMDMLKDRSTHKPELLKE KQNIKIYGRKSIGRLKIDKILFSEGN EGDMDITKSCTVKINHRSLDKHDS HLVSLAGTSKILHARGQVEMEINR SHTTPLECKIISPSDITPGDLDKTMM SIDDHEELDMTKSHTVFIDYQAEAK GVLPRDLDFQLSKKESLQPKVTSLS AEEIYISKNSSESNHLPKGSQTLILEE GSNSGLGEETNDAQKPGFLNELLSG KTQRRKSLSLKNKSITFPENDKSYRE IPQSSAVEINNETRLEDKGFVPL AGTSKPVLSAYGPEDMEISIGOTTAS EYKTVPEEITIPMDKTMFVDNFG DLDVTRSHTVFIDCQAKEKVLDEYT NLGIQKTKTLGSEGDTHIQEITKNP AAQHKHHMTTVIPSSTVSDQSSMK IKFHKADRDEEVKGKEVEANMLKQT KPESCLLNITDGKNVDFTSSYTADVC RSSDKYSSLPNISSSDNSGGNTMSL CDKNKEKAYNCQVPNEFTYAAILPS TYHMDSKLSVFPPCPSKEVTQTES AIALLKDEDPVVEPLGEMATFNSKH

							VSLNLAKDQTEAFVDVSVASQPHLS AQQSPSTQKGQDVARREDEGILAKAG KKALPFLLNVAASTWENESKIPTN VEHFAVTYEKELSISIQTDKCNTNVQ SPSNSALTTQVIQTHANAEGALDFLV PSTVSCFSSTKPSLSNLNRKTEEVLD FQTVNLLPPAEQLLEEGSQAHSMSI VQATEIYRLGSRNDRDEESKTCNE AETTSVPLKTAVKDKTRRCSLGIFLP KLPSKRSCSITGVDDLEQILADAADL TQLETQPVC SKDPGIGSVAAKLNLSP SQFINEENLPVYPGEILSSDSVSLDIE ESVLIDTSQRESLPSENKTENCRAQ KRTRVEENDVTNEKKIRTHDSAQDQ EVSCLEPKNVLEF
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