

UniprotKB ID	Entry name	organism	full name	oglcnaacscore	oglcnaac sites	phosphorylation sites	PMIDS	sequence
Q9P2N6	KANL3_HUMAN	Homo sapiens	KAT8 regulatory NSL complex subunit 3	46.586465	S523;S532;S534;T796;T797;T805;T810;T811;T824;T859;T860;T864	S523;S536;S540;S773	3626323;30379171;20068230;33214551;30059200;31492838;31881804;20305658;28657654;32574038;29351928;29485866;21740066;28450392;34019948;29237092;30620550;23301498	MAHRGGERDFQTSARRMGTSLLFQ LSVHERELDLVFLDHSYAKPWSAHP DASSARPTRMLFVTPRRQHESTIES DVPIDVETVTSTPMPLYDNQKARSV MNECERHVIFARTDADAPPPEDWE EHVNRTGWTMAQNKLFNKILKALQ SDRLARLANEGACNEPVLRRVAVDK CARRVRQALASVSWDTKLIQWLHTT LVETLSLPMLAAYLDALQTLKGGKIPT LIDRMLVSSNTKTGAAGAEALSLLK RPWDPVAVGVLSHNKPSKLPGSPLILI ASSGPSSSVFPTSRHRFWQSQLSC LGKVIPVATHLLNNGSGVGVLCLE HMIGAVRSKVLEIHSHFPHKPIILIG WNTGALVACHVSVMEYVTAVVCLG FPLLTVDGGPRGDVDDPLLDMKTPVL FVIGQNSLQCHPEAMEDFREKIRAE NSLVVVGADDNLRISKAKKSEGL TQSMVDRCIQDEIVDFLTGVLTRAE GHMGSEPRDQDAEKKKPRDVARR DLAFEVPERGSRPASPAKLPASPSG SEDLSSVSSPTSSPKTKVTTVTSAQ KSSQIGSSQLLKRHVQRTEAVLTHK QAQAQFAAFLKQNMLVRKALPPGTS SCLFVPISSEPEEGEKEDLRVQLKR HHPSSPLPGSKTSKRPKIKVSLISQG DTAGGPCAPSQGSAPAEAAGGKPITM TLGQASAGAKELTGLLTTAKSSSEGE GVSASPVPSVSSSTAPSALHTLQSR LVATSPGSSSLPGATSASSLLQGLSFS LQDISSKTSGLPANPSPGPAPQATS KLPTPMQSLGAIITGTSTIVRTIPVAT TLSSLGATPGGKPTAIHQLLTNGGLA KLASSLPGLAQISNQASGLKVPTTITL TLRGQPSRITLSPMGSGAAPSEESS SQVLPSSSQRLPPAP

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