

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	39.566891	S523;S532;S534;S700;S703;S710;S714;S717;S719;T720;T765;T787;S791;T796;T797;T799;S800;T801;T805;T810;T811;T824;T859;T860;T862;T864	S523;S536;S540;S773	35132862;23301498;21740066;30059200;20068230;35289036;33214551;31881804;29351928;29237092;30379171;28450392;34019948;35254053;20305658;37340703;29485866;28657654;35138101;30620550;31492838;32574038	MAHRGGERDFQTSARRMGTSLLFQ LSVHERELDLVFLDHSYAKPWSAHP DASSARPTRMLFVTPRRQHESTIES DVPIDVETVTSTPMPLYDNQKARSV MNECERHVIFARTDADAPPPEDWE EHVNRTGWTMAQNKLFNKILKALQ SDRLARLANEGACNEPVLRRVAVDK CARRVRQALASVSWDTKLIQWLHTT LVETLSLPMLAAYLDALQTLKGGKIPT LIDRMLVSSNTKTGAAGAEALSLLK RPWDPAVGVLSHNKPSKLPGSPLILI ASSGPSSSVFPTSRRHRFWQSQLSC LGKVIPVATHLLNNGSGVGVLCLE HMIGAVRSKVLEIHSHFPHKPIILIG WNTGALVACHVSVMEYVTAVVCLG FPLLTVDGGPRGDVDDPLLDMKTPVL FVIGQNSLQCHPEAMEDFREKIRAE NSLVVVGADDNLRISKAKKSEGL TQSMVDRCIQDEIVDFLTGVLTRAE GHMGSEPRDQDAEKKKPRDVARR DLAFEVPERGSRPASPAAKLPASPSG SEDLSSVSSPTSSPKTKVTTVTSAQ KSSQIGSSQLLKRHVQRTEAVLTHK QAQAQFAAFLKQNMLVRKALPPGTS SCLFVPISSEPEEKEKEDLRVQLKR HHPSSPLPGSKTSKRPKIKVSLISQG DTAGGPCAPSQGSAPAAAGGKPITM TLGQASAGAKELTGLLTTAKSSSEGE GVSASPVPSVSSSTAPSALHTLQSR LVATSPGSSSLPGATSASSLLQGLSFS LQDISSKTSGLPANPSPGPAPQATSV KLPTPMQSLGAIITGTSTIVRTIPVAT TLSSLGATPGGKPTAIHQLLTNGGLA KLASSLPGLAQISNQASGLKVPTTITL TLRGQPSRITTLSPMGSGAAPSEESS SQVLPSSSQRLPPAP

0