	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus
0	Q7Z3B3	KANL1_HUMAN	Homo sapiens	KAT8 regulatory NSL complex subunit 1	14.877748	S838	S249;S268;S991;S994;T1003;S1045	34019948;30379171;39302247;40307207;35132862;40596516;35254053	MAAMAPALTDAAAEAHHIRFKLAPP SSTLSPGSAENNGNANILIAANGTKR KAIAAEDPSLDFRNNPTKEDLGKLQ PLVASYLCSDVTSVPSKESLKLQGVF SKQTVLKSHPLLSQSYELRAELLGR QPVLEFSLENLRTMNTSGQTALPQA PVNGLAKKLTKSSTHSDHDNSTSLN GGKRALTSSALHGGEMGGSESGDL KGGMTNCTLPHRSLDVEHTTLYSN NSTANKSSVNSMEQPALQGSSRLSP GTDSSSNLGGVKLEGKKSPLSSILFS ALDSDTRITALLRRQADIESRARRLQ KRLQVVQAKQVERHIQHQLGGFLEK TLSKLPNLESLRPRSQLMLTRKAEA ALRKAASETTTSEGLSNFLKSNSISE ELERFTASGIANLRCSEQAFDSDVTD SSSGGESDIEEEELTRADPEQRHVPL RRRSEWKWAADRAAIVSRWNWLQA HVSDLEYRIRQQTDIYKQIRANKGLI VLGEVPPPEHTTDLFLPLSSEVKTDH GTDKLIESVSQPLENHGAPIIGHISES LSTKSCGALRPVNGVINTLQPVLAD HIPGDSSDAEEQLHKKQRLNLVSSS SDGTCVAARTRPVLSCKKRRLVRPN SIVPLSKKVHRNSTIRPGCDVNPSCA LCGSGSINTMPPEIHYEAPLLERLSQ LDSCVHPVLAFPDDVPTSLHFQSML KSQWQNKPFDKIKPPKKLSLKHRAP MPGSLPDSARKDRHKLVSSFLTTAK LSHHQTRPDRTHRQHLDDVGAVPM VERVTAPKAERLLNPPPPVHDPNHS KMRLRDHSSERSEVLKHHTDMSSS SYLAATHHPPHSPLVRQLSTSSDSPA PASSSSQVTASTSQQPVRRRRGESSF DINNIVIPMSVAATTRVEKLQYKEILT PSWREVDLQSLKGSPDEENEEIEDL SDAAFAALHAKCEEMERARWLWTT SVPPQRRGSRSYRSSDGRTTPQLGS ANPSTPQPASPDVSSSHSLSEYSHG QSPRSPISPELHSAPLTPVARDTPRH LASEDTRCSTPELGLDEQSVQPWER RTFPLAHSPQAECEDQLDAQERAAR CTRRTSGSKTGRETEAAPTSPPIVPL KSRHLVAAATAQRPTHR	True	False	2.642	5.0	5.0	False	False