

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A2RSY1	KANL3_MOUSE	Mus musculus	KAT8 regulatory NSL complex subunit 3	28.939361	T858	S523;S536;S540;S772	26192747;21606357;29187734	MAHRGGERDFQTSARRMGTSLLFQ LSVHERELDLVFLDHSYAKPWSAHP DASSARPTRMLFVTPRRQENTIES DVPIDVETVTATPVPLYDNQKARSV MNECERHVIFARTDADAPPPEDWE EHVNRTGWTVAQNKLFLNKILKALQS DRLARLANEGACNEPVLRRVAVDKC ARRVRQALASVSWDTKLTQWLHTTL VETLSLPMLAAYLDALQTLKGKIPTLI DRMLVSSNTKTGAAGAEALSLLLKR PWDPAVGVLSHNKPSKLPGSPLILIV SSGPSSSVFPASRRHRFWQSQLSCL GKVIPVATHLLNNGSGVGLQCLEH MIGAVRSKVLEIHSFPHKPIILIGW NTGALVACHVSVMEYVTAVVCLGFP LLTVDGPRGDVDDPLLDMKTPVLFV IGQNSLQCHPEAMEDFREKIRAENS LVVVGADDNLRISKAKKKSEGLTQ SMVDRCIQDEIVDFLTGVLTRAEGH VGSEPRDQDAEKKKKPRDLTRRDLA FEIPERGSRPASPAARLPTSPGSED LSSVSSSPTSSPKTKVTTVTSTQKSS QIGTSQLLKRHVQRTEAVLTHRQAQ AQFAAFLKQNMLVRKAFPPGTSSCL FVPISSESVEDIEKEELRVQLKRHHS SSPLPGAKPSKRPKIKVSLISQGDTV GGPCTLSQGGTPEAAGGKPITMTLG ASAGAKELTGLLTTAKSSSSEGGTA STTPSVASSATPNAIHTLQSRIVAT SPGSSLPGTASASSLLQGLSFSLQDI SSKTSGLPGSPSPGPAPQATSVKLPT PMQSLGAIITGTSTIVRTIPVATTLSS LGATPGGKPTAIHQLLTNGSLAKLAS SLPGLAQISNQASGLKVPTTITLTLR GQPSRITTLSPMGSGATPSEEPNSQ MLPSSSQRLPPAP

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