

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
A2RSY1	KANL3_MOUSE	Mus musculus	KAT8 regulatory NSL complex subunit 3	36.507815	S699;S700;T710;T786;S790;T795;T796;T810;T858;T859;T863	S523;S536;S540;S772	29187734;36852467;21606357;34887587;35822049;26192747	MAHRGGERDFQTSARRMGTSLLFQ LSVHERELDLVFLDHSYAKPWSAHP DASSARPTRMLFVTPRRQQENTIES DVPIDVETVTATPVPLYDNQKARSV MNECERHVIFARTDADAPPPEDWE EHVNRGTGWTVAQNKLFNKILKALQS DRLARLANEGACNEPVLRRVAVDKC ARRVRQALASVSWDTKLTQWLHHTL VETLSLPMLAAYLDALQTLKGGKIPTLI DRMLVSSNTKTGAAGAEALSLLKR PWDPAVGVLSHNKPSKLPGSPLILIV SSGPSSSVFPASRRHRFWQSQLSCL GKVIPVATHLLNNGSGVGLQCLEH MIGAVRSKVLEIHSFPHKPIILIGW NTGALVACHVSVMEYVTAVVCLGFP LLTVDGPBGDVEDDPLLDKTPVLFV IGQNSLQCHPEAMEDFREKIRAENS LVVVGADDNLRISKAKKKSEGLTQ SMVDRCIQDEIVDFLTGVLTRAEGH VGSEPRDQDAEKKKKPRDLTRRDLA FEIPERGSRPASPAARLPTSPSGSED LSSVSSSPTSSPKTKVTTVTSTQKSS QIGTSQLLKRHVQRTEAVLTHROAQ AQFAAFLKQNMLVRKAFPPGTSSCL FVPISSESVEDIEKEELRVQLKRHHS SSPLPGAKPSKRPKIKVSLISQGDIV GGPCTLSQGGTPEAAGGKPITMTLG ASAGAKELTGLLTAKSSSEGGGTA STTPSVASSATPNAIHTLQSRVAT SPGSSLPGTASASSLLQGLSFSLQDI SSKTSGLPGSPSPGPAPQATSVKLP PMQSLGAIITGTSTIVRTIPVATLSS LGATPGGKPTAIHQLLTNGSLAKLAS SLPGLAQISNQASGLKVPTTITLTLR GQPSRITLSPMGSGATPSEEPNSQ MLPSSSQRLPPAP