

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38853	KEL1_YEAST	Saccharomyces cerevisiae	Kelch repeat-containing protein 1	9.365641	NaN	S20;S23;S25;S94;T467;T477;S503;T604;S613;T626;S689;S691;S717;S748;S837;S848;S958;S997;S1003;S1022	33229814	MAGFSFAKKFTHKKHGKTPSDASIS DQSREASLSTPPNEKFFTKQETPQK GRQFSQGYHSNVNKTSSPPMFARK QVSESRIQPSAVPPQQRNVSGPSTTL HKQLSKQREYTVWNRKLNQNSPFPR YRHVASAYVTDKNQIYVIGGLHDQS VYGDTWILTAFDNATRFSTTTIDISEA TPPPRVGHA AVL CGNAFV VFGGDTH KVNKEGLMDDDIYLLNINSYKWTVP APVGPRPLGRYGHKISIIATTQMKT LYVFGGQFDDTYFNDLAVYDLSSFR RPDSHWEF LKPRFTFPPI TNFTMIS YDSKLWVFGGDTLQGLVNDVFM YD PAINDWFIIDTTGEKPPPVQEHATVV YNDLMCVVGGKDEHDAYLNSVYFL NLKSRKWFKLPVFTAGIPQGRSGHS LTLKNDKILIMGGDKFDYARVEEYD LHTSDIDMQRGTIVYTLDLARIKDL PGVMDVPTDTPTPRNGNLDLATPVT PTSHQTKNMNVPISAAPLASAPSPAP KDFSDADRLNREVHNRNVSTEHQN QSHPVNSES HLIAEPNILTPYVPSES SQTPVMKITSNKPFDTPTIQKEPDL ETMDPTVGNQRIPSSIYGDNLTPAN QIKNNSPILETLPSNEIKTPQNGNIE EIKHLPDADEKIDSTTTFDQEINGDK LGTSSMSKVEEDGNVADEDEIGVA QMASSPSKDQFKIKHYNESSELSQN NTEIDKLEPVDITIKKSDTAGHDSA NHVIDASDEKNVSPMGDVPTDTKN EEASVPINRDATTEVVDRALFEKLR ELQSLKELTHEKALEAGAHIKELETE LWQLKSQKNSGTTKEIDELDSVRLQ SKCEILEADNHSLEDKVNELEELVN SKFLDIENLNEVIQFQNEKIKSLELE PNYKEKLEELQIEHENLSRENERLK NESKQHNEIINN VAN YSSQLGSLIS HWKENRANSSFLESSSLISVSDEN GEKTVGEPYGDQSRHHRVVINKLTN RLDDL LERSQELTISKEKLSSEYHAL KMEHSSLSQDVLVKENEIKKI QNDY KESISSMDSASKALMVSQRELEKYK SLNKKLIDELDELKFKNGVCSENFE NGLRSTEESNNVKNSNSIRENQFN IKINDLKAELFITNQERDDLKSEVLE LKKRLLNLENNTKQVNEDADSDLL