

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appara
Q63ZY3	KANK2_HUMAN	Homo sapiens	KN motif and ankyrin repeat domain-containing protein 2	24.592877	S344;T345	S19;S83;S86;S89;S92;T168;S323;T329;S356;S375;T472;S540;T552	28657654;39531497;38665916;35254053	MAQVLHVPAPFPPTPGPASPPAFPA KDPDPPYSVETPYGYRLDLDFLKYVD DIEKGHTLRRVAVQRRPRLSSLRPGP GSWWTSTESLCSNASGDSRHSAYS CGRGFYPQYGALETRGGFNPRVVERT LLDARRRLEDQAATPTGLGSLTPSAA GSTASLVGVLPPPTPRSSGLSTPVP PSAGHLAHVREQMAGALRKLRLQLE EQVKLIPVLQVKLSVLQEEKRLTVQ LKSQKFLGHPTAGRGRSELCLDLPD PPEDPVALETRSVGTWVRERDLGMP DGEAALAAKVAVLETQLKKALQELQ AAQARQADFPQAWPPDPSFVRVDT VRVVEGPREVEVVAAGAPAQRA QSLEPYGTGLRALAMPGRPESPPVF RSQEVVETMCPVPAATSNVHMVK KISITERSCDGAAGLPEVPAESSSSPP GSEVASLTQPEKSTGRVPTQEPTHR EPTRQAASQSEEEAGGTGGPPAGVR SIMKRKEEVADPTAHRRLQFVGVN GGYESSSEDSSTAENISDNDSTENE APEPRERVPSVAEAPQLRPACTAAAK TSRQECQLSRESQHIPTAEGASGSN TEEEIRMELSPDLISACLALKEYLDN PNALTERELKVAYTTVLQEWLRLAC RSDAHPELVRRHLVTFRAMSARLLD YVNIADSNNGNTALHYSVSHANFPV VQQLLDSGVCKVDKQNRAGYSPIML TALATLKTODDIETVLQFRLGNINA KASQAGQTALMLAVSHGRVDVVKAL LACEADVNVQDDDGSTALMCACEH GHKEIAGLLAVPSCDISLTD RDGST ALMVALDAGQSEIASMLYSRMNIKC SFAPMSDDDESPTSSSAEE	True	False	2.459	2.984	5.0	0.81	0.569