

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region	
0	O95271	TNKS1_HUMAN	Homo sapiens	Poly [ADP-ribose] polymerase tankyrase-1	19.994096	NaN	NaN	37217939	MAASRRSQHHHHHHQQQLQPAPG ASAPPPPPPLSPGLAPGTTPASPTA SGLAPFASPRHGLALPEGDGSRDPP DRPRSPDPVDGTSCCSTTSTICTVAA APVVPAVSTSSAAGVAPNPAGSGSN NSPSSSSSPTSSSSSPSPGSSLAE SPEAAGVSSSTAPLGPAAAGPTGVPA VSGALRELLEACRNGDVSrvKRLVD AANVNAKDMAGRKSSPLHFAAGFG RKDVVEHLLQMGANVHARDGGLI PLHNACSFHAEVVSLLCQGGADPN ARDNWNYPPLHEAAIKGKIDVICIVLL QHGDAPNIRNTDGGKALDLADPSAK AVLTGEYKDELLEAARSGNEEKL ALLTPLNVNCHASDGRKSTPLHLAA GYNRVRVQQLLQHGADVHAKDKGG LVPLHNACSYGHYEVELELLKHGAC VNAMDLWQFTPLHEAASKNRVEVC SLLL SHGADPTLVNCHGKSAVDMAP TPELRRLTYEFKGHSLQAAAREADL AKVKKTLALEIINFKQPQSHETALHC AVASLHPKRKQVTELLLRKGANVNE KNKDFMTPLHVAERAHNDVMEVL HKHGAKMNALDTLGQTALHRAALA GHLQTCRLLSYGSDPSIISLQGF AQMGNEAVQQLSESTPIRTSDVDY RLEEASKAGDLETVKQLCSSONVNC RDLEGRHSTPLHFAAGYNRVSVVEY LLHHGADVHAKDKGGLVPLHNAC YGHYEVAELLVRHGASVNVADLWKF TPLHEAAAKGKYEICKLLKHGADPT KKNRDGNTPLDLVKEGDDIQLLR GDAALLDAKKGCLARVOKLCTPEN INCRDTQGRNSTPLHLAAGYNNLEV AEYLLHGGADVNAQDKGGLIPLHNA ASYGHVDIAALLIKYNTCVNATDKW AFTPLHEAAQKGRTOCALLAHGA DPTMKNQEGQTPDLATADDIRALLI DAMPPEALPTCFKQATVVSASLISP ASTPSCLSAASSIDNLTGPLAELAVG GASNAGDGAAGTERKEGEVAGLDM NISQFLKSLGLEHLRDFETEQITLD VLADMGHEELKEIGINAYGHRHCLI KGVRLGGQGTNPYLTFHCVNQ GTILLDLAPEDKEYQSVEEMQSTIR EHRDGGNAGGIFNRYNVIRIQKVVN KKLRERFCHRQKEVSEENHNHNE RMLFHGSPFINAIIHKGFDERHAYIG GMFGAGIYFAENSSKSNQVYVYIGG GTGCPHDKDRSCYICHRQMLFCRVT LGKSFLLQFSTMKMAHAPPCHHSVIG RPSVNGLAYAEYVIYRGEQAYPEYLIT YQIMKPEAPSQTATAAEQKT	True	True	4.59	4.844	2.366	1.755	5.0	3.132	1.946