

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
Q6PFX9	TNKS1_MOUSE	Mus musculus	Poly [ADP-ribose] polymerase tankyrase-1	29.227022	NaN	NaN	29187734;24788674	MAASRRSQHHHHHQQQLQPAPG ASAPPPPPPLSPGLAPGTPASPTA GGLAPFASPRHGLALPEGDGSRDPP DRPRSPDPVDGAVCTVAAPAAVPAA SAAVGVAPTPAGGGGGGNNNSASSA SSPTSSSSSPSPGSSLAESPEAAG VGSTATLGAGAAGLPGVPAVSGAL RELLEACRNGDVSrvKRLVDAANVN AKDMAGRKSSPLHFAAGFGRKDVV EHLQMGANVHARDGGGLIPLHNA CSFGHAEVVSLCCOGADPNARDN WNYTPLHEAAIKGKIDVCIVLLQHGGA DPNIRNTDGSALDLADPSAKAVLT GEYKDELLEAARSGNEEKLMAALLT PLNVNCHASDGRKSTPLHLAAGYNR VRIVQLLQHGADVHAKDKGGLVPL HNACSYGHYEVELELLKHGACVNA MDLWQFTPLHEAASKNRVEVCSLL LSHGADPTLVNCHGKSAVDMAPTPE LRERLTYEFKGSLLQAAREADLAK VKKTLALEIINFKQPQSHETALHCAV ASLHPKRKQVAELLLRKGANVNEKN KDFMTPLHVAAERAHNDVMEVLHK HGAKMNALDSLQGTALHRAALAGH LQTCRLLSYGSDPSIISLQGFATAQ MGNEAVQQLSESTPMRTSDVDYRL LEASKAGDLETVKQLCSPQNVNCRD LEGRHSTPLHFAAGYNRVSVVEYLL HHGADVHAKDKGGLVPLHNACSYG HYEVAELLVRHGASVNVADLWKFTP LHEAAAKGKYEICKLLKHGADPTK KNRDGNTPLDLVKEGDTIQDLLRG DAALLDAAKKGCLARVQKLCPTPENI NCRDTQGRNSTPLHLAAGYNNLEV AEYLLEHGADVNAQDKGGLIPLHNA ASYGHVDIAALLIKYNTCVNATDKW AFTPLHEAAQKGRTOQLCALLAHGA DPTMKNQEGQTPDLATADDIRALLI DAMPPEALPTCFKQATVVSASLISP ASTPCLSAASSIDNLTGPLTDLAVG GASNAGGAAGAERKEGEVAGLDM NISQFLKSLGLEHLRDFEFTEQITLD VLADMGHEELKEIGINAYGHRHCLI KGVRELLGGQQGTNPYLFHCVNQ GTILLDLAPEDKEYQSVEEEMQSTIR EHRDGGNAGGIFNRYNVIRIQVNV KKLRERFCHRQKEVSEENHHHNE RMLFHGSPFINAIHKGFDERHAYIG GMFGAGIYFAENSSKSNQYVYIGGG GTGCPHDKDRSCYICHRQMLFCRV LGKSFQFSTMKMAHAPPGHHSVIG RPSVNGLAYAEYVYRGEQAYPEYLIT YQIMKPEAPSQTATAAEQKT	True	False	5.0	4.385	2.19	1.864	4.267	1.96	1.924