

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9C0D5	TANC1_HUMAN	Homo sapiens	Protein TANC1	10.041052	S300;S1662;S1676;S1814	S63;S66;S67;S207;S270;S465;S1439;S1668;S1676;S1677	30379171;30059200;31492838;29237092;34019948;30620550	MLKAVLKKSSREGGKGGKKEAGSDF GPETSPVLHLDHSADSPVSSLPTAED TYRVSLAKGVSMSPSSPLLPRQSHL VQSRVNKKSPGPPVVRKPKYVESPRVP GDAVIMPFREVAKPTPEDEHEAKAD NEPSCSPAAQELLTRLGFLLEGIGPS ATHITIEDKNETMCTALSQGISPCST LTSSTASPSTDSPCSTLNSCVSKTAA NKSPCETISSPSTLESKDSGIIATITS SSENDDRSGSSLEWNKDGNLRLGV QKGVLDHRRADNCSPVAEEETTGSA ESTLPAEASSAGDGPVPYSQSSSLI MPRPNSVAATSSTKLEDLSYLDGQR NAPLRTSIRLPWHNTAGGRAQEVKA RFAPYKQDILLKPLLFEVPSITTDV FVGRDWLFHQIEENLRNTELAENR GAVVVGNVGFGKTAIISKLVALSCHG SRMRQIASNSPGSSPKTSDPTQDLH FTPLLSPSSSTSASSTAKTPLGSISAE NQRPREDAVKYLASKVVAYHYCQAD NTYTCLVPEFVHSIAALLCRSHQLAA YRDLLIKEPQLQSMLSLRSCVQDPV AAFKRGVLEPLTNLRNEQKIPEEEYII LIDGLNEAEFHKPDYGDTLSSFITKII SKFPAWLKLVTVRANFQEIISALPFV KLSLDDFPDNKDIHSDLHAYVQHRV HSSQDILSNISLNGKADATLIGKVSS HLVLRSLGSYLYLKLTLDFQRGHLV IKSASYKVPVSLSELYLLQCNMKF MTQSAFERALPILNVALASLHPMTD EQIFQAINAGHIQGEQGWEDFQQR MDALSCFLIKRRDKTRMFCHPSFRE WLVWRADGENTAFLCEPRNGHALL AFMFSRQEGKLNROQTMELGHHIL KAHIFKGLSKKTGISSSHLQALWIGY STEGLSAALASLRNLYTPNVKVSRLI ILGGANVNYRTEVLNNAPILCVQSH LGHEEVVTLLEFGACLDGTSNGM TALCYAAAAGHMMLVCLLTKKGVRV DHLDDKGGQCALVHSALRGHGDILQY LLTCEWSPGPPQPGTLRKSHALQQA LTAAASMGHSSVQCLLGMEKEHE VEVNGDTLWGETALTAAGRGKLE VCELLLGHGAAVSRTNRRGVPPLFC AARQGHWQIVRLLLERGCDVNLSDK QGRTPLMVAACEGHLSTVEFLLSKG AALSSLDKEGLSALSWACKGHRV VQYLVEEGAAIDQTDKNGRTPDLA AFYGDAETVLYLVEKGAVIEHVDHS GMRPLDRAIGCRNTSVVVALLRGA KLGNAAWAMATSKPDILILLQKLM EEGNVMYKKGKMKEAAQRYQYALR KFPREGFGEDMRPFNELRVSLYLNL SRCRRKTNDFGMAEEFASKALELKP KSYEAFYARARAKRNSRQFVAALAD LQEAVKLCPTNQEVKRLARVEEFC KQLQRSQQKQKQGPLPAPLNDSEN EEDTPTPGLSDHFHSEETEEETSP QEESVSPTPRSQSSSVPSYIRNLQ

EGLQSKGRPVSPQSRAGIGKSLREPV
AQPGLLLOPSKQAQIVKTSOHLGSG
QSAVRNGSMKVQISSQNPPSPMPG
RIAATPAGSRTQHLEGTGFTTRAGC
GHFGDRLGPSQNVRLQCGENGAH
PLPSKTKTTERLLSHSSVAVDAAPPN
QGGLATCSDVRHPASLTSSGSSGSP
SSSIKMSSSTSSLTSSSFSDGFKVQ
GPDTRIKDKVVTHVQSGTAEHRPRN
TPFMGIMDKTARFQQSNPPSRSW
HCPAPEGLLTNTSSAAGLOSANTEK
PSLMQVGGYNNQAKTCSVSTLSASV
HNGAQVKELEESKQIPVHSQENRI
TKTVSHLYQESISKQQPHISNEAHR
HLTAAKPKRSFIESNV