

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9C0D5	TANC1_HUMAN	Homo sapiens	Protein TANC1	16.369165	S213;S214;S291;S300;S1477;S1537;T1590;S1662;S1671;S1676;S1814	S63;S66;S67;S207;S270;S465;S1439;S1668;S1676;S1677	29237092;35254053;30379171;38253038;30620550;31492838;34019948;30059200;35289036	MLKAVLKKSREGGKGGKKEAGSDF GPETSPVLHLDHSDSPVSSSLPTAED TYRVSLAKGVSMSLPSSPLLPRQSHL VQSRVNNKSPGVPVRKPKYVESPRVP GDAVIMPFREVAKPTEPDEHEAKAD NEPSCSPAAQELLTRLGFLLEGIPG ATHITIEDKNETMCTALSQGISPCST LTSSTASPSTDSPCSTLNSCVSKTAA NKSPCETISSPSTLESKDSGIIATITS SSENDDRSGSSLEWNKDGNLRLGV QKGVLDHRRADNCSPVAEEETTGSA ESTLPAESSAGDGPVPYSGSSSLI MPRPNSVAATSSTKLEDLSYLDGQR NAPLRTSIRLPWHNTAGGRAQEVKA RFAPYKQDILLKPLLFEVPSITDSDV FVGRDWLFHQIEENLRNTELAENR GAVVVGNGVFGKTAIISKLVALSCHG SRMRQIASNSPGSSPKTSDPTQDLH FTPLLSPPSSSTSASSTAKTPLGSISAE NQRPREDAVKYLASKVVAYHYCQAD NTYTCLVPEFVHSIAALLCRSHQLAA YRDLLIKEPQLQSMLSLRSCVQDPV AAFKRGVLEPLTNLRNEQKIPEEYII LIDGLNEAEFHKPDYGDTLSSFITKII SKFPAWLKLVTVRANFQEIISALPFV KLSLDDFPDNKDIHSDLHAYVQHRV HSSQDILSNISLNGKADATLIGKVSS HLVLRSLGSYLKLTLDLDFQRGHLV IKSASYKVVVSLSELYLLQCNMKF MTQSAFERALPILNVALASLHPMTD EQIFQAINAGHIQGEQGWEDFQQR MDALSCFLIKRRDKTRMFCHPSFRE WLWVRADGENTAFLCEPRNGHALL AFMFSRQEGKLNROQTMELGHHL KAHIFKGLSKKTGISSSHLQALWIGY STEGLSAALASLRNLYTPNVKVSRLI ILGGANVNYRTEVLNNAIPILCVQSH LGHEEVVTLLEFGACLDGTSENGM TALCYAAAAGHMKLVCLLTKKGVRV DHLDDKKGQCALVHSALRGHDILQY LLTCEWSPGPPQPGTLRKSHALQQA LTAASMGHSSVVQCLLGMKEHE VEVNGTDTLWGETALTAAGRKLE VCELLLGHGAAVSRNRRGVPLFC AARQGHVQIVRLLLERGCDVNLSDK QGRTPLMVAACEGHLSTVEFLLSKG AALSSLDKEGLSALSWACLKGHRAV VQYLVEEGAAIDQTDKNGRTPLDLA AFYGDAETVLYLVEKGAVIEHVDHS GMRPLDRAIGCRNTSVVVALLRKGA KLGNAAWAMATSKPDILHLLQKLM EEGNVYKKGKMKKEAAQRYQYALR KFPREGFGEDMRPFNELRVSLYLN SRCRRKTNDFGMAEEFASKALELKP KSYEAFYARARAKRNSRQFVAALAD LQEAVKLCPTNQEVKRLARVEEEC KQLQRSQQQKQQGPLPAPLNDSEN EEDTPTPGLSDHFHSEETEEETSP QEEVSPTPRSQPSSSVPSYIRNLQ EGLQSKGRPVSPQSRAGIGKSLREPV AQPGLLLQPSKQAQIVKTSQHLGSG QSAVRNGSMKVQISSQNPPSPMPG

								RIAATPAGSRTQHLEGTGTFTRAGC GHFGDRLGPSQNVRLQCGENPAH PLPSKTKTTERLLSHSSVAVDAAPPN QGGLATCSDVRHPASLTSSGSSGSP SSSIKMSSTSSLTSSSSFSDFKQVQ GPDTRIKDKVVTHVQSGTAEHRPRN TPFMGIMDKTARFQQQSNPPRSW HCPAPEGLLTNTSSAAGLQSANTEK PSLMQVGGYNNQAKTCSVSTLSASV HNGAQVKELEESKQIPVHSQENRI TKTVSHLYQESISKQOPHISNEAHR HLTAAKPKRSFIESNV
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