	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
0	Q7Z6L1	TCPR1_HUMAN	Homo sapiens	Tectonin beta-propeller repeat-containing protein 1	20.111456	T202	S386;S388;S391;S412;S417;S938;S949	30379171;29351928	MPNSVLWAVDLFGRVYTLSTAGQY WEMCKDSQLEFKRVSATTQCCWGI ACDNQVYVYVCASDVPIRRREEAYE NQRWNPMGGFCEKLLLSDRWGWS DVSGLQHRPLDRVALPSPHWEWES DWYVDENFGGEPTEKGGWTYAIDFP ATYTKDKKWNSCVRRRKWIRYRRYK SRDIWAKIPSKDDPKELPDPFNDLSV GGWEITEEPVGRLSVWAVSLQGKV WYREDVSHSNPEGSSWSLLDTPGE VVQISCGPHDLLWATLWEGQALVRE GINRSNPKGSSWSIVEPPGSENGVM HISVGVSVVWAVTKDWKVWFRRGV NSHNPCGTSWIEMVGEMTMVNVG MNDQVWGIGCEDRAVYFRQGVTPS ELSGKTWKAIIAARECDRSHSGSSSS LLSAGCFFGDEVRGSGESAPSDTDA SSEVERPGPGQILPAEPLDDSKNATG NSASGLGAGRTAEDTVEDACPAEGS REARPNTHPGPAPTPAELPWTNIDL KEAKKVPSHSAAGFPETTSLSSLGLL PLGLEEPYGVDDHPLWAWVSGGGC VVEACAMPRWFTVQAGLSSSVHML SLSITPAQTAAWRKQIFQQLTERTKR ELENFRHYEQAVEQSVWVKTGALQ WWCDWKPHKWVDVRLALEQFTGH DGVRDSILFIYYVVHEEKKYIHIFLNE VVALVPVLNETKHSFALYTPERTRQR WPVRLAAATEQDMNDWLALLSLSC CESRKVQGRPSPQAIWSITCKGDIFV SEPSPDLEAHEHPLPCDQMFWRQM GGHLRMVEANSRGVVWGIGYDHTA WVYTGGYGGGCFQGLASSTSNIYTQ SDVKCVHIYENQRWNPVTGYTSRGL PTDRYMWSDASGLQECTKAGTKPPS LQWAWVSDWFVDFSVPGGTDQEG WQYASDFPASYHGSKTMKDFVRRR CWARKCKLVTSGPWLEVPPIALRDV SIIPESPGAEGSGHSIALWAVSDKGD VLCRLGVSELNPAGSSWLHVGTDQP FASISIGACYQVWAVARDGSAFYRGS VYPSQPAGDCWYHIPSPPRQRLKQV SAGQTSVYALDENGNLWYRQGITPS YPQGSSWEHVSNNVCRVSVGPLDQ VWVIANKVQGSHSLSRGTVCHRTGV QPHEPKGHGWDYGIGGGWDHISVR ANATRAPRSSSQEQEPSAPPEAHGP VCC	True	False	2.527	4.529	1.402	1.282	1.168	2.229	1.