

UniprotKB ID	Entry name	organism	full name	oglcnaescore	oglcnaesites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regio
Q86TV6	TTC7B_HUMAN	Homo sapiens	Tetratricopeptide repeat protein 7B	18.379595	NaN	S160;S202;S625;S629;S630;S673;S677;S678;S681	30379171	MATKKAGSRLETEIERCSEQWER IPELVKQLSAKLIANDDMAELLGSES KLEQYLKEHPLRQGASPRGPKPQLT EVRKHLTAALDRGNLKFSEFLQESNL IMAKLNVYEGDYKEALNIYARVGLD DLPLTAVPPYRLRVIAEAYATKGLCL EKLPISSSTSNLHVDREQDVITCYEK AGDIALLYLQEIERVILSNIQNRSPKP GPAPHDQELGFFLETGLQRAHVLYF KNGNLTRGVGRFRELRAVETRITQ NLRMTIARQLAEILLRGMCEQSYWN PLEDPPCQSPLEDDPLRKGANTKTYTL TRRARVYSGENIFCPQENTEEALLLL LISESMANRDAVLSRIPEHKSDRLIS LQSASVVYDLLTIALGRRGOYEMLSE CLERAMKFAFEFFHLWYQFALSML AAGKSARAVKVLKECIRLKPDDATIP LLAAKLCMGSLHWLEEAKEFAKTIV DVGEKTSFKAKGYLALGLTYSLQAT DASLRGMQEVLRKALLAFQRAHSL SPTDHQAAFYLAQLAISROIPEALG YVRQALQLQGDDANSLHLLALLLSA QKHYHDALNIIDMALSEYPENFILLF SKVKLQSLCRGPDEALLTCKHMLQI WKSCYNLTNPDSGRGSSLLDRITIA DRRQLNTITLPDFSDPETGSVHATSV AASRVEQALSEVASSLQSSAPKQGPL HPWMTLAQIWLHAAEVYIGIKPAE ATACTQEAANLFPMSHNVLYMRGOI AELRGSMDARRWYEEALAIPTHV KSMQRLALILHQLGRYSLAEKILRDA VQVNSTAHEVWNLGELVLAQAGND AAATECFLTALELEASSPAVPFTIIPR VL	True	True	5.0	3.288	0.741	1.022	1.258	5.0	0.569