

| 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 |
|--------------|------------|--------------|-----------------------------------|--------------|---------------|------------------------|--|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|
| P17987 | TCPA_HUMAN | Homo sapiens | T-complex protein 1 subunit alpha | 16.123428 | NaN | S6;T181;S491;S544;S551 | 16408927;38665916;35083852;33214551;23301498;32870666;35138101;34019948;30397120;32119511;34725712;37217939;28510447 | MEGPLSVFGDRSTGETIRSQNVMAA ASIANIVKSSLGVPGLDKMLVDDIGD VTITNDGATILKLEVEHPAAKVLCE LADLQDKEVGDGTTSVVIAAELLKN ADELVKQKIHPSTVISGYRLACKEAV RYINENLIVNTDELGRDCLINAAKTS MSSKIIGINGDFFANMVVDAVLAIKY TDIRGQPRYPVNSVNLKAHGRSQM ESMLISGYALNCVVGSGQMPKRIVN AKIACLD FSLQTKMKLGVQVITDP EKLDQIRQRES DITKERIQKILATGA NVILTTGGIDDMCLKYFVEAGAMAV RRVLKRD LKRIAKASGATILSTLANL EGEETF EAAMLGQAEV VQERICDD ELILIKNTKARTSASIILRGANDFMC DEMERSLHDALCVVKRVLESKSVVP GGGAVEAALSIYLENYATSMGSREQ LAIAEFARSLLVIPNTLAVNAAQDST DLVAKLRAFHN EAQV NPERKNLKW I GLDLSNGKPRDNKQAGVFEPTIVKV KSLKFATEA AITILRIDDLIKLHPESK DDKHGSYEDAVHSGALND | True | False | 5.0 | 2.986 | 2.023 | 1.862 | 3.567 | 2.7 |