

| 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 | extracel region |
|--------------|------------|--------------|----------------------------------|--------------|---------------|-----------------------|--|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|-----------------|
| P80314       | TCPB_MOUSE | Mus musculus | T-complex protein 1 subunit beta | 46.458321    | NaN           | S3;S260;T261          | 28135057;37507081;28528544;37453647;29187734;34887587;25153642;24788674;33300544;21540332;39627609;39927985;22645316 | MASLSLAPVNIKAGADEERAETAR<br>LSSFIGAIAIGDLVKSTLGPKGMDKIL<br>LSSGRDAALMVTNDGATILKNIGVD<br>NPAAKVLVDMSRVQDDEVGDGTTT<br>VTVLAELLREAESLIAKKIHPQTII<br>GWREATKAAREALLSSAVDHGSDEA<br>RFWQDLMNIAGTTLSSKLLTHHKD<br>HFTKLAVEAVLRLLKSGNLEAIHVIK<br>KLGGS LADSYLDEGFLDKKIGVNO<br>PKRIENAKILANTGMDTDKIKIFGSR<br>VRVDSTAKVAEIEHAEKEKMEKEVE<br>RILKHGINCFINRQLIYNYPEQLFGA<br>AGVMAIEHADFAGVERLALVTGGEI<br>ASTFDHPELVKLGSCKLIEEVMIGED<br>KLIHFSGVALGEACTIVLRGATQIL<br>DEAERSLHDALCVLAQTVKDPRIVY<br>GGGCSEMLMAHAVTQLANRTPGKE<br>AVAMESFAKALRMLPTHADNAGYD<br>SADLVAQLRAAHSEGHITAGLDMKE<br>GTIGDMAVLGITESFOVKROVLLSAA<br>EAAEVILRVDNIIKAAPRKRVPDHPHC | True          | False         | 5.0     | 0.562   | 0.469         | False                 | False           | 0.188           | False           |