| | 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 | extracellular region |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | V5DCV8 | V5DCV8_TRYCR | Trypanosoma cruzi Dm28c | Coatomer subunit gamma | 19.946178 | NaN | NaN | 30984116 | MQAMPSDRDRYRYEEEDEEETLPFE<br>GLEKTSALQQCRVFNDVRLDISSCL<br>RSMTQCLYLLCTGTTLTETEATDLFF<br>MSTKLLQSTHPKLRRLHYVLMKELS<br>PLVEQRFIASNSLMIDIKSNNDASKC<br>NGIRTLFKVMNSSLYASMDRTIVEA<br>LTSQSSNVVCAALVTGLHIAQVNPE<br>MARKWGTQLTEVIRSCGKAQYAAIA<br>LLHKMRKNDRLSVTRLIDQAKNGVI<br>RSPMALCLVIKMCTELMREDFEGSL<br>DIYKFVTSMMHNNNDLVVFESVKSI<br>CSLRNITAKEVSPAVMVVQLYLNTQ<br>SAVLRFSAIRVLNEVATLHPAAVSPI<br>NSEIENLVTDPNRIIATLAITTLLKTG<br>TEYTIERLITQLSTAGYLRELGDEFK<br>MVIIDAMRVLSAKFPAKYNVFLGFLS<br>KLLAEEGSSQLKENVVDVTMEIAKS<br>NPDSKESVLKHLAEFIDDCNYSQIVR<br>RVLMYLGEEVPLTENPKMFVRYVYN<br>HATLEGPEIRAVAVSTLAKLAAYVPS<br>LRRSIVVLLKRTCNDADDEVRDRAV<br>LYTRIFLNNDENTIRSMVCDVANTV<br>AMSRQARIAARSAVLNTVAEDIGRQ<br>KAIAKEAESGTEETSAVSHAVLQGRE<br>KMRKIKQLLELGEPIVSSEPVPLTDP<br>DSEYVVTVMKHTYISNIVLQFKVKNT<br>MEKVTFKNIAIELDTDEVGVEPQYFI<br>PIESVAPASTSYGYAVLRYEEEQYPS<br>GTLECRFKFAMQEEGGDVEEEEEYP<br>LEGFDINVSDFIAPVNLGGTFETQW<br>EKMQLEETTGTYSLSSMRNLTVAAR<br>ELVEFFGMYVEGGSPGKITTASHTL<br>QMSGTMVNRARSLVLITARVFIAKD<br>KSVALQLMLRGATAELREYLSAALLA | None | None | None | None | None | None | None | None | None |