| | 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** | Q03761 | TAF12_YEAST | Saccharomyces cerevisiae | Transcription initiation factor TFIID subunit 12 | 24.475439 | NaN | S129;S286 | 33229814 | MSSNPENSGVNANNNTGTGNADAI<br>TGAQQNMVLQPRQLQEMAAKFRTL<br>LTEARNVGETTPRGKELMFQAAKIK<br>QVYDALTLNRRRQQAAQAYNNTSN<br>SNSSNPASIPTENVPNSSQQQQQQQ<br>QQTRNNSNKFSNMIKQVLTPEENQ<br>EYEKLWQNFQVRHTSIKEKETYLKQ<br>NIDRLEQEINKQTDEGPKQQLQEKK<br>IELLNDWKVLKIEYTKLFNNYQNSK<br>KTFYVECARHNPALHKFLQESTQQQ<br>RVQQQRVQQQQQQQQQQQQQQQQQ<br>QQQQQQQRQGQNQRKISSSNSTEIP<br>SVTGPDALKSQQQQQNTITATNNPR<br>GNVNTSQTEQSKAKVTNVNATASM<br>LNNISSSKSAIFKQTEPAIPISENISTK<br>TPAPVAYRSNRPTITGGSAMNASAL<br>NTPATTKLPPYEMDTQRVMSKRKLR<br>ELVKTVGIDEGDGETVIDGDVEELLL<br>DLADDFVTNVTAFSCRLAKHRKSDN<br>LEARDIQLHLERNWNIRIPGYSADEI<br>RSTRKWNPSQNYNQKLQSITSDKVA<br>AAKNNGNNVASLNTKK | None | None | None | None | None | None | None | None | None |