| | 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** | E2QCZ3 | E2QCZ3_DROME | Drosophila melanogaster | NaN | 28.06426 | T193 | NaN | 40245499 | MQELKQRFERRQLADDLKQKQVIQ DHLLQKILYSKRLLVDHVEALEHCM QELQTASSASSGRQLLTTRNACLILG GTLLEHLITPRGIRYTMVPSILISGTF AALFAVKSVFVCRNKIVERKLEQLVK TIDEFGNCIRRNMTYFQEIIIMKQAE LIESRQIERAWDCITAAKEVTEILYDA TRKLETDYPLPTKYSAYYAPMEELRE CEYFKNNVTDYSPKHIKDFHNIFAYV QSQYLLRLALTITTRPSISQLSEDLVK IDSLVRQLVQEEEEQHFSNLALALQN KKQLELAELNATRAEQQQSGPIAVL QHSSLKLSACMVAVAAECQALDVTL QKLTVTEAAKKNNSKELVAVANNM HGIENALAVCCDDFQRLMLVYNKFL HSQLDLEGELPKPKQDLDDEFPESIL RVEFSQNVDAPQQRDDFYAYMYDE NEEQHFEAEQNAPFPSPEKELLNFE KRITKGRFKPVLKQLKDRIDPIRQVM LEKEREVLASKGINVDELFGKMERV EQQQEDNRLADAAPLPVYESSSNSD SDSADEEAFRRRSKQHKERDNFAE MRQFLAQKQAINLFKLPPPPVAAGG EELLESEC | True | False | 5.0 | 5.0 | 3.069 | 2.206 | 2.139 | 2.59 | 2.498 |