| | 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** | A1Z7F1 | A1Z7F1_DROME | Drosophila melanogaster | alpha-glucosidase | 25.496522 | NaN | NaN | 33925313 | MACFKVLIAAILVLGIHCALGSAAAV<br>DLDLERATTAADTTRDWWQVAQFY<br>QIYPRSYKDSDGDGIGDLQGIISKLDY<br>LKEIGVTATWLSPIYSSPMADFGYDI<br>SDFFDIQPEYGTLADFDELIAEAKKR<br>NIKIILDFVPNHSSDENVWFQKSVK<br>REKGYEDYYMWHDGYVNATTGKRE<br>PPSNWLQAFRGSAWEWNDERQQYY<br>LHQFAVKQPDLNYRNPAVVAQMKR<br>VLTYWLDRGVAGFRMDAVPWCFEV<br>LPDADGRYPDEPLSGYTDDPDDSSY<br>LKHIYTQDLRETVEMVFQWRTLLDD<br>YQRIHGGDTRVIMVETYSGLDYVMQ<br>FYGNRTTKGAQMPFNFQFIIGGNGD<br>KNNTQLNATGFVKIISSWLSQMPAG<br>QTANWVMGNHDQRRVGSRYGENRI<br>DLMNMLQMFLPGVSITYQGEELGM<br>TDLDISWEDSRDPAACNSNSDIYEQ<br>FTRDPARTPFQWSDEANAGFSTNAT<br>TWLPINPNYVTVNAKAENSTSPSHL<br>SLYKQLVDLRKSKTLQFGATRYANV<br>GDNVVAIRRYLSGEPSYVLVANVLDT<br>SVSGIDVASAIYATGSYKIKLLNPQAK<br>ATVGDSVTLSNLTLEPYAALILESV | None | None | None | None | None | None | None | None | None |