| | 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** | Q9VXA3 | Q9VXA3_DROME | Drosophila melanogaster | NaN | 25.496522 | NaN | NaN | 33925313 | MVTMNSEADKTQATNASSTAAPAK<br>TDCTSSNSSDPLAVAIPEPQSATAVV<br>GGGAGGAAPGTATPSPTPLANGSNN<br>GTGSSVSPAATGTGAVATAASAPAAA<br>GTTSASPPASNPTTASTSTASVVGAK<br>TAAATTTSSSSATSSSTAAAGSKQSG<br>SSGGLLTVSGNHHHHHHHAHPQQSA<br>AGQSAAQPPQPPTATALAVPSASHH<br>QRGGKNEDAPNILVYKKMEAIIEKM<br>QAESTGVAVRTVKAFMSKVPSVFTG<br>ADLVAWILKNFDVEDVTEALHFAHL<br>LSSHGYIFPIDDHALTVKNDGTFYRF<br>QTPYFWPSNCWEPENTDYAVYLCKR<br>TMQNKTRLELADYEAENLAKLQKM<br>FSRKWEFIFMQAESQSKVAKKRDKL<br>ERKVLDSQERAFWDVHRPMPGCVN<br>TTEIDIKKAYRRGGSSHGTGSAGASL<br>AKNPVEQLTRIIALRKQKLERRTIKVS<br>KAAEALVAYYDQYNEFDYFITSPELP<br>NPWQTDSTEMWDTEKNSKEVPVRR<br>VKRWAFSLRELLNDAIGREQFTKFL<br>EKEYSGENLKFWESVQEMKALPQS<br>EIKEAIQKIWQEFLAPEAPCPVNVDS<br>KSVELAREAVNSPSGPNRWCFDVAA<br>SHVYHLMKSDSYSRYLRSDMYKDYL<br>NCSRKKIKSIPNLFGVKR | None | None | None | None | None | None | None | None | None |