| | 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** | Q8T038 | Q8T038_DROME | Drosophila melanogaster | NaN | 28.06426 | S566 | NaN | 40245499 | PSWPQQAPAHSLIVDSTPLFNVEEIY<br>PKHHLNLPRNDLKKESPAHHIAKGN<br>SYRVPRHEKKEHQQLSPVKKRVKES<br>SPPHQQRYQRAAHVSPQYHAHHNC<br>NYGNGSGYSASAGSVVASSAASSSNI<br>VNGSSSSSHHHHVPVAHAQPYSSSS<br>GHHIVSNCNANGGAGGAVVYQQPP<br>AHAHQQHGQQPHPQSSQHPQHVK<br>QPTITIHDTPSPTAVITISDSEDEGGE<br>AGGAQVPVLKQRAHAQSQTSGSLLQ<br>HAPSSSCSRSSAHQQPHPHPQQQQ<br>QQQINYGDHDPEDARRRHHAAAAA<br>AVGSPKHHHHQQQQQPAPPPQQQQ<br>QQYQPQPPPQQVLPQPQPSIKYEPG<br>QSQKKRILAMAQSECGYQPQVPSQP<br>SSSASLPHIPTKQEPAEFYPEYAAAPP<br>QQLDTKRSSWAPTSSGSGVSALPLA<br>HPKREAPSVAPISYVAPSVAPPLAHS<br>KSSSASSSSSISAATAAAAAAAAAAA<br>ASVGPPSWGPPQVYRQPSQPPPGSV<br>GLPGSTQPPPSSVAPHPHHHSHGH<br>HHHQAGTPLGGSPSSTAAAALLQPD<br>IYAQGDIYRRPTVFVSQAAPSYAYAN<br>RAVVAPPPAHNSSSRQVIPSHPLPAH<br>IQIPTQYSQFGPLSPAQVAASKHAAH<br>FAPTNIWYGAE | NaN | NaN | NaN | NaN | NaN | NaN | NaN | NaN | None |