| | 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** | A9UNF4 | A9UNF4_DROME | Drosophila melanogaster | NaN | 28.06426 | T562 | NaN | 40245499 | MPRICRCCCKRDLKLLGLFEASQPT<br>LANTTVASASAAASVASNKSSGTLAK<br>TCATEAETETPQTLSANPVNPGTKAT<br>SSPSEAATTLSTSTRPASASFSSTSFS<br>TSTSPMRRRTTNTTAPSASPAPRGRS<br>SDNAVDYTLSGAHSSMDIVLEEMTI<br>WMLNINRDDGLPQEICRQCMAQFL<br>MVAKFRRKCVRMQQRLQDFAQEIS<br>DRQTARQAKRQQKVQKKTHDRWLQ<br>LEPTSHEEQQPIRKRRTVSESDGDST<br>AAAAAAEAATATATTPAKQVRTDFES<br>LPPPKPSPQQHRVRKYRTLSIDCEM<br>ERPAASINASSMPWKTKAARKELSE<br>NWASSCSTAVMSPNRQRSSTSSSE<br>HMSDFSVGPCCGFESDSNPSSTSNS<br>ARRTVILVKQQSPEKTSPMYDLMPR<br>RGRRPRKAPIYMPAKRQRRSLVRKE<br>PLYRKAQPVNEENKKSDQKDIYPKE<br>SRMEGLAPSCDEAIDSTSDAPMEVA<br>SNVVQKKCDEICLNGLIDSRNQLGA<br>MEMEPVNEMIASNGVDKELTLSVQ<br>NNKETTSEEQIFEATSVIKESVQPEQ<br>SAEDIQVRTEIESIANDIRAEADLVQS<br>EELTSSTETSESQPKAVEQLESMGK<br>QESYALPQNMEAHESVQQTQTVDE<br>SDSADQREIIVSIPLEVLDENLQRRL<br>CVDPETEAANRQSADPKEFALRPDD<br>VNNNENDENFCQSATADSFNSAVA<br>LRTNKIEATPPGGTDSDDTDTDEVPI<br>VPSGPTPMDFMAEFAKHCANGIEQL<br>KVTTPVPSPTPSDLVPLNDLEAKQKL<br>EVEMNDVETTLNGILNEMQDQHMY<br>TPNCAHIDEFLTPADYATLTEQESSV<br>SSPPNPFEQSSVAEPKDEKPSATEDP<br>FDNSNFTNELIGFQNDIPCFENIETP<br>EEGQNLHLELTQFLRDLQ | NaN | NaN | NaN | NaN | NaN | NaN | NaN | NaN | None |