| | 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** | Q9VYJ1 | Q9VYJ1_DROME | Drosophila melanogaster | NaN | 28.06426 | T564 | NaN | 40245499 | MKPPLSSITRLVVILGVMLLKLAAAQ DCQQQLDNDTDAEVKPCENFHQYA CGNWYSNTSQRILGAHDMRSKWVA NLKLQLIRYLEESSSTDSFSANTSVT TSTAQLGTLYRCCLLSGQSIRTYTDA LSRSGANFPLLANHTGGSFDWVLA NAALRTYGAQGLWRLLVKDNWQSA DQRIFYMLPPKFDLLGRDDELNEFL YQRYLKILLLELGLRVRRAALLAEKL VAFEKSLQKLLPNDAEQTLVLRKPQ SLKDLESQLPRLELKRYFETILEGID WDSKTLLLVADLDYLQGLQRLLAEH TSDPMVLSTWLLLQLPAYFELRLHD DDKLGNQRNHCLEQLRKLMPGQLG RLQMQLVLGESYQDVYDTAIQQIAVL FRDLKQQFELVLNETEVFNEDLVTR NLARDKLRAMRLLTPRLQDPINGGP GTPQLGGNCDANLLQLSLSQAKLEY RQVFGEPVTSVPADPLSVNAYYRLKL NRIELPLGLMATPLIQQQCSKEADTR AKLSAGLGYILAHEMVHAFDLDGLN YDATGQLANGQWSARAIIRFGLRAG CYLGVRYSNATLTVNENIADTEGLRL ALDTYRHRSGATNLRTFFVAFAQNW CGTVASTSGLSQHAGHAERVNNVV GNMPEFGDTFECKATSQMNPVNKC HIW | False | False | 1.676 | 1.754 | 1.829 | 1.794 | 1.629 | 4.318 | 2.914 |