| | 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** | Q01477 | UBP3_YEAST | Saccharomyces cerevisiae | Ubiquitin carboxyl-terminal hydrolase 3 | 24.475439 | NaN | NaN | 33229814 | MNMQDANKEESYSMYPKTSSPPPP<br>TPTNMQIPIYQAPLQMYGYTQAPYLY<br>PTQIPAYSFNMVNQNQPIYHQSGSP<br>HHLPPQNNINGGSTTNNNNINKKK<br>WHSNGITNNNGSSGNQGANSSGSG<br>MSYNKSHTYHHNYSNNHIPMMASP<br>NSGSNAGMKKQTNSSNGNGSSATS<br>PSYSSYNSSSQYDLYKFDVTKLKNLK<br>ENSSNLIQLPLFINTTEAEFAAASVQ<br>RYELNMKALNLNSESLENSSVEKSS<br>AHHHTKSHSIPKHNEEVKTETHGEE<br>EDAHDKKPHASKDAHELKKKTEVK<br>KEDAKQDRNEKVIQEPQATVLPVVD<br>KKEPEESVEENTSKTSSPSPSPPAAK<br>SWSAIASDAIKSRQASNKTVSGSMV<br>TKTPISGTTAGVSSTNMAAATIGKSS<br>SPLLSKQPQKKDKKYVPPSTKGIEPL<br>GSIALRMCFDPDFISYVLRNKDVEN<br>KIPVHSIIPRGIINRANICFMSSVLQV<br>LLYCKPFIDVINVLSTRNTNSRVGTS<br>SCKLLDACLTMYKQFDKETYEKKFL<br>ENADDAEKTTESDAKKSSKSKSFQH<br>CATADAVKPDEFYKTLSTIPKFKDLQ<br>WGHQEDAEEFLTHLLDQLHEELISA<br>IDGLTDNEIQNMLQSINDEQLKVFFI<br>RNLSRYGKAEFIKNASPRLKELIEKY<br>GVINDDSTEENGWHEVSGSSKRGK<br>KTKTAAKRTVEIVPSPISKLFGGQFR<br>SVLDIPNNKESQSITLDPFQTIQLDIS<br>DAGVNDLETAFKKFSEYELLPFKSSS<br>GNDVEAKKQTFIDKLPQVLLIQFKRF<br>SFINNVNKDNAMTNYNAYNGRIEKI<br>RKKIKYGHELIIPEESMSSITLKNNTS<br>GIDDRRYKLTGVIYHHGVSSDGGHY<br>TADVYHSEHNKWYRIDDVNITELED<br>DDVLKGGEEASDSRTAYILMYQKRN | None | None | None | None | None | None | None | None | None |