

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q01477	UBP3_YEAST	Saccharomyces cerevisiae	Ubiquitin carboxyl-terminal hydrolase 3	9.365641	NaN	NaN	33229814	MNMQDANKEESYSMYPKTSSPPPP TPTNMQIPIYQAPLQMYGYTQAPYLY PTQIPAYSFNMVNQNQPIYHQSGSP HHLPPQNNINGGSTTNNNNINKKK WHSNGITNNNGSSGNQGANSSGSG MSYNKSHTYHHNYSNNHIPMMASP NSGSNAGMKKQTNSSNGNGSSATS PSYSSYNSSSQYDLYKFDVTKLKNLK ENSSNLIQLPLFINTTEAEFAAASVQ RYELNMKALNLNSELENSSVEKSS AHHHTKSHSIPKHNEEVKTETHGEE EDAHDKPHASKDAHELKKKTEVK KEDAKQDRNEKVIQEPQATVLPVVD KKEPEESVEENTSKTSSPSPSPAAK SWSAIASDAIKSRQASNKTVSGSMV TKTPISGTTAGVSSTNMAAATIGKSS SPLLSKQPQKKDKKYVPPSTKGIEPL GSIALRMCDFPDFISYVLRNKDVEN KIPVHSIIPRGIINRANICFMSSVLQV LLYCKPFIDVINVLSTRNTNSRVGTS SCKLLDACLTMYKQFDKETYEKKFL ENADDAEKTTESDAKKSSKSKSFQH CATADAVKPDEFYKTLSTIPKFKDLQ WGHQEDAEEFLTHLLDQLHEELISA IDGLTDNEIQNMLQSINDEQLKVFFI RNLSRYGKAEFIKNASPRLKELIEKY GVINDDSTEENGWHEVSGSSKRGK KTKTAAKRTVEIVPSPISKLFGGQFR SVLDIPNNKESQSITLDPFQTIQLDIS DAGVNDLETAFFKKFSEYELLPFKSSS GNDVEAKKQTFIDKLPQVLLIQFKRF SFINNVNKDNAMTNYNAYNGRIEKI RKKIKYGHELIPEESMSSITLKNNTS GIDDRRYKLTGVYHHGVSSDGGHY TADVYHSEHNKWYRIDDVNITELED DDVLKGGEEASDRTAYILMYQKRN