

# OTUB1

**PDB:2ZFY**

**Entry Clone Accession:**NP\_060140

**Entry Clone Source:**otub1.BC007519.MGC.AU30F4.pOTB7

**SGC Clone Accession:**otub1.040.271.72C04 (SDC072)

**Tag:**N-terminal: MGSSHHHHHSSGLVPR\*GS (removed)

**Host:**E.coli BL21(DE3)

**Vector:**pET28a-LIC

## Sequence:

mgsshhhhhhssglvpr\*gsEIAVQNPLVSRLELSVLYKEYAEDDNIYQQKIKDLHKKYSYIRKTRPDGNCFYRAFGFSHLEALLD  
DSKELQRFKAVSAKSKEDLVSQGFTEFTIEDFHNTFMDLIEQVEKQTSVADLLASFNDQSTSDFLVVYLRLLTSGYLQRESKFFEHF  
IEGGRTVKEFCQQEVEPMCKESDHIIHIALAQALSVSIQVEYMDRGEGGTTNPHIFPEGSEPKVYLLYRPGHYDILYK

## Growth

**Medium:**TB

**Procedure:** The protein was expressed in E. coli BL21 (DE3) grown in Terrific Broth (TB) in the presence of 50 µg/mL of kanamycin at 37°C to an OD600 of 7.5. Cells were then induced by isopropyl-1-thio-D-galactopyranoside (IPTG), final concentration 0.05 mM, and incubated overnight at 15°C. The culture was centrifuged and the cell pellets were collected and stored at -80°C.

## Purification

**Procedure:** Imac: The cleared lysate from a 2 L culture was loaded onto 3 ml TALON metal-affinity resin column (BD Biosciences) at 4°C. The column was washed with 40 ml Wash buffer, and the protein was eluted with 10 ml Elution buffer.

Tag removal: 1 Unit of thrombin (Sigma T9681) per milligram of protein was added to the 10 mL sample, stored overnight without shaking at 4°C.

Gel-filtration: It was further purified by gel filtration on a HighLoad 16/60 Superdex 200 column (GE Healthcare, Amersham) equilibrated with Gel Filtration buffer and concentrated to 68 mg/ml by ultrafiltration using Amicon Ultra centrifugal filter with 10 kD cutoff and stored at -80 °C.

Protein yield was 17 mg per liter of bacterial culture.

## Extraction

**Procedure:** The cell pellet was defrosted and cells were lysed by sonication, 10 s on, 10 s off at 40% amplitude for 10 min. The lysate was cleared by centrifugation for 45 minutes at 15,500

RPM, 4°C.

**Concentration:**68 mg/mL

## Structure Determination

**MassSpec:**Mass-spectroscopy by LCMS shows that the product was pure and of correct molecular weight

**Crystallization:**Purified protein was crystallized using the hanging drop vapor diffusion method. Crystals grew when the protein (68 mg/mL) was mixed with the reservoir solution in a 1:1 volume ratio, and the drop was equilibrated against a reservoir solution containing 30% PEG 8000, 0.2 M Sodium acetate, and 0.1 M Sodium cacodylate at pH 6.5 in 293 Kelvin.