

# UBE2Q1

**PDB:**2QGX

## Revision

**Revision Type:**created

**Revised by:**created

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**Entry Clone Accession:**NP\_060052

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**SGC Clone Accession:**ubc64.248.414; plate SDC45G03

**Tag:**mgsshhhhhhssglvpr\*gs

**Host:**BL21 (DE3)

## Construct

**Prelude:**

**Sequence:**

mgsshhhhhhssglvpr\*GGSVQVTDRMKELRDIYRSQSFKGGNYAVELVNDSLVDWNVKLLKVDQDSALHNDLQILKEKEGADF  
ILLNFSFKDNFPFDPPFVRVVSPVLSGGYVLGGGAICMELLTKQGWSSAYSIESVIMQISATLVKGKARVQFGANKSQYSLTRAQQSYKSLVQIHEKNGW

**Vector:**p28a-LIC-Thrombin

## Growth

**Medium:**TB

**Antibiotics:**

**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. Protein expression was then induced with 0.05 mM isopropyl-1-thio-D-galactopyranoside (IPTG) overnight at 15°C. The culture was centrifuged (12,000 x g, 15 min) and cell pellets collected and stored at -80°C.

## Purification

### Procedure

**Column 1:** TALON metal-affinity resin column (BD Biosciences)

**Column 2:** HighLoad 16/60 Superdex 200 column (GE Healthcare, Amersham)

The cleared lysate from a 2 liter culture was loaded onto 3 ml TALON metal-affinity resin column (BD Biosciences) at 4°C. The column was washed with 10 ml Wash buffer A, 10 ml Wash buffer B and then 30 ml Wash buffer A, and the protein was eluted with 6 ml Elution buffer.

His-tags was removed by incubation of the protein with 1 U thrombin per mg protein (overnight at 4 °C), and the protein was further purified by gel filtration on a HighLoad 16/60 Superdex 200 column (GE Healthcare, Amersham) equilibrated with Gel Filtration buffer and concentrated by ultrafiltration using Amicon Ultra centrifugal filter with 10kD cutoff.

## Extraction

### Procedure

The cell pellet from a 2 liter culture was resuspended in 50 ml Lysis buffer and lysed using Microfluidizer at 18,000 psi. The lysate was cleared by centrifugation (40,000 x g, 30 minutes).

### Concentration:

### Ligand

### MassSpec:

**Crystallization:** Crystals were grown in hanging drops by mixing 2 microL protein solution (10 mg/ml) with 2 microL well solution (22% PEG4000, 0.1 M Tris-Cl, pH7.5, 0.12 M MgCl<sub>2</sub>, 1 mM DTT) at 21°C. For cryoprotection, the crystals were soaked in well solution supplemented with 20% glycerol.

### NMR Spectroscopy:

### Data Collection:

### Data Processing: