

# Human UBH12

**PDB:**2FAZ

## Revision

**Revision Type:**created

**Revised by:**created

**Revision Date:**created

**Entry Clone Accession:**ubh12.001.076.GSC.#.vec

**Entry Clone Source:**

**SGC Clone Accession:**ubh12.001.076; plate SDC029H12

**Tag:**MGSSHHHHHSSGLVPR\*GS

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

## Construct

**Prelude:**

**Sequence:**

MGSSHHHHHSSGLVPR\*GSMWIQVRTMDGRQTHTVDSL SRLTKVEELRRKIQELFHV EPGLQRLFYRGKQMEDGHTLF DYEVRLND  
TIQLLVRQS

**Vector:**pET28a-LIC

## Growth

**Medium:**

**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 about 4. Cells were induced with 0.05 mM isopropyl-1-thio-D-galactopyranoside (IPTG), and incubated overnight at 15 °C. The culture was centrifuged and cell pellets were collected and stored at -80°C.

## Purification

### Procedure

The clarified supernatant from a 2 L culture was loaded at approximately 1 mL/min by gravity onto 4 mL of Ni-NTA resin (Qiagen 30450). Twelve column volumes of Wash buffer were used to wash the column at approximately 3 mL/min. Samples were eluted from the Ni-NTA resin by exposure to 7.5 mL Elution buffer at 1 mL/min flow rate. EDTA was added to the eluate to final concentration of 1 mM. Approximately 15 minutes later, dithiothreitol was added to the eluate to a final concentration of 2 mM. Protein concentration was determined using molecular coefficients. Thrombin (Sigma T9681; 1 unit per milligram of protein to 50 U maximum) and CaCl<sub>2</sub> (4 mM final concentration) were added to the Ni-NTA eluate in 50 mL conical vials (352096, BD Biosciences) and the conical vial was incubated without shaking, overnight, at 4°C. All gel filtration columns, buffers, and protocols are identical for uncut and thrombin-treated proteins. An XK 16x65 column (part numbers 18-1031-47 and 18-6488-01, GE Healthcare) packed with HighLoad Superdex 200 resin (10-1043-04, GE Healthcare) was pre-equilibrated with Gel filtration buffer using an AKTA Purifier. The full volume of eluate was loaded onto the column at 1.5 mL/min and 2 mL fractions collected using peak fractionation protocols. Peak fractions were analyzed for purity using SDS-PAGE or visual analysis of the chromatogram and pooled. The protein was concentrated by ultrafiltration and analyzed by mass spectrometry.

## Extraction

### Procedure

The cell pellet from a 2 L culture was resuspended in Lysis buffer and lysed using Microfluidizer. Fresh PMSF was added to the lysate and the lysate cleared by centrifugation (24,000 rpm, 30 minutes).

### Concentration:

#### Ligand

#### MassSpec:

**Crystallization:** Purified protein was crystallized using the sitting drop vapor diffusion method. Crystals grew when the protein (13 mg/mL) was mixed with the reservoir solution in a 1:1 volume ratio, and the drop was equilibrated against a reservoir solution containing 20.0% polyethylene glycol 3350, 0.2 M tri-Lithium Citrate in 291 K temperature.

#### NMR Spectroscopy:

#### Data Collection:

#### Data Processing: