

# CBS

PDB:4UUU

## Revision

**Revision Type:**created

**Revised by:**created

**Revision Date:**created

**Entry Clone Accession:**IMAGE:3028099

**Entry Clone Source:**MGC

**SGC Clone Accession:**

**Tag:**N-terminal, TEV protease cleavable hexahistidine tag

**Host:**

## Construct

**Prelude:**

**Sequence:**

"MHHHHHHSSGVDLG TENLYFQSMKPWWHLRVQELGLSAPLTVLPTITCGHTIEILREKGF DQAPVVDEAGVILGMVTLGNMLSSL  
LAGKVQPSDQVGKVIYKQFKQIRLDTLGR LSHILEMDHFALVVHEQQRQMVFGVVT AIDLLNFVAAQE" MHHHHHHSSGVDLG TEN  
LYFQ\*SM is the purification tag plus TEV protease recognition site \*.

**Vector:**pNIC28-Bsa4

## Growth

**Medium:**

**Antibiotics:**

**Procedure:**

## Purification

### Procedure

Cell Lysis Cell pellets were dissolved in approximately 60 ml lysis buffer containing EDTA-free protease inhibitors (diluted 1:1000). The resuspended cell pellets were then treated for 1 hour with 2 mg/ml lysozyme at 4 °C and then broken by homogenization. The insoluble cell debris was separated from the soluble fraction by centrifugation. Talon IMAC The clarified soluble fraction was incubated with 3.0 ml pre-equilibrated Talon resin slurry in 50% ratio (v/v) in binding buffer. for 1 hour at 4°C with rotation. After it was centrifuged with a low brake speed to pellet the resin. The resulting supernatant was removed and the resin pellet was resuspended in the remaining supernatant and was applied to a gravity column. Next the resin was washed with 1 x 10 ml of binding buffer, 5 x 5 ml wash buffer and the protein was eluted with 6 x 3 ml and 2 x 5 ml elution buffer. Size Exclusion Gel Filtration Superdex s75 16/60 Gel Filtration. Pooled elution fractions were then applied to the GF column (pre-equilibrated in GF buffer) at 1.0 ml/min. 1.0

ml fractions were collected. Anionic Exchange Chromatography Combined fractions were concentrated to 5.0 ml and then diluted in low salt buffer. This low salt solution was then applied to a pre-equilibrated Resource Q column and then the protein was eluted with a linear gradient of higher salt concentrations using the high salt buffer. Clean fractions were isolated and combined.

## **Extraction**

### **Procedure**

BL21(DE3)-R3-pRARE2A single colony each was used to inoculate 2 x 50 ml TB media with 50 ug/ml kanamycin and 50 ug/ml chloroamphenicol. These were placed in an incubator at 37 °C, 180 rpm and left to grow overnight. The next day these starter cultures were mixed and used to inoculate 3 L of TB media (10 ml starter culture used per 1 L ) grown at 37 °C, 180 rpm and supplemented with 50 ug/ml kanamycin and 50 ug/ml chloroamphenicol.. When the OD600 reached 1.8 the cells were induced with 0.1 mM IPTG AT 18 °C and left overnight. Cell harvestCells were harvested by centrifugation at 4,500 g for 15 minutes after which the supernatant was discarded and the cell pellets were stored at -80 °C.

**Concentration:**To set up plates the sample was concentrated to approximatley 5 mg/ml using a 10 kDa MWCO concentrator.

### **Ligand**

**MassSpec:**Expected mass: 17425.2 Da Measured mass: 17426.17 Da

**Crystallization:**Crystals were grown by vapour diffusion in sitting drop at 20°C. A sitting drop consisting of 100 nl protein with 1 mM S-adenosyl-L-methionine and 50 nl well solution was equilibrated against well solution containing 36% PEG550MME -- 0.1M tris pH 7.5 -- 0.2M calcium chloride.

### **NMR Spectroscopy:**

**Data Collection:**Resolution: 1.9 Å X-ray source: Diamond Light Source beamline IO2

### **Data Processing:**