

# ITSN2

**PDB:**3JZY

## Revision

**Revision Type:**created

**Revised by:**created

**Revision Date:**created

**Entry Clone Accession:**BC146779

**Entry Clone Source:**OpenBiosystems EHS1001-99608273 (SGC 30-C2)

**SGC Clone Accession:**HPC098-C04

ITSN2:D1174-E1665

**Tag:**N-terminal tag: mhhhhhhsgrenlyfq\*g

**Host:**BL21-CodonPlus(DE3)-V2R pRARE2

## Construct

**Prelude:**Tag not removed.cDNA sequence verified by sequencing and it matches expected one.The cDNA template used contains a mutation K1525Q compared with reference sequence NP\_062541.2.The construct contains 3 domains, only the C-terminal C2 domain was crystallized.

**Sequence:**

mhhhhhhsgrenlyfqgDTMQPIERKRQGYIHELIQTEERYMADLQLVVEVFQKRMAESGFLTEGEMALIFVNWKELIMSNTKLLK  
ALRVRKKTGGEKMPVQMGDILAAELSHMQAYIRFCSCQLNGAALLQQKTDEDTDFKEFLKKLASDPRCKGMPLOSSFLKPMQRITR  
YPLLIIRSILENTPESHADHSSLKLALERAEELCSQVNEGVREKENSDRLEWIQAHVQCEGLAEQLIFNSLTNCLGPRKLLHSGKLYK  
TKSNKELHGFLFNDFLLLTVKQFAVSSGSEKLFFSKSNAQFKMYKTPIFLNEVLVVKLPTDPSSDEPVFHISHIDRVYTLRTDNIN  
ERTAWVQKIKAASEQYIDTEKKQREKAYQARSQKTSGIGRLMVHVIEATELKACKPNGKSNPYCEISMGSQSYTTRTIQDTLNPKWN  
FNCQFFIKDLYQDVLCLTLFDRQFSPDDFLGRTEIPVAKIRTEQESKGPMTRLLLHEVPTGEVWVRFDLQLFE

**Vector:**pET28-mhl (GI:134105571)

## Growth

**Medium:**Terrific Broth

**Antibiotics:**Kanamycin 50  $\mu$ g/mL + Chloramphenicol 25  $\mu$ g/mL

**Procedure:**LEX Bubbling. The target protein was expressed in E. coli by inoculating 100 mL of overnight culture grown in Luria-Bertani medium into a 1.8 L of Terrific Broth medium in the presence of 50  $\mu$ g/mL kanamycin and 25  $\mu$ g/mL chloramphenicol at 37 degC. When OD600 reached ~3.0, the temperature of the medium was lowered to 15 degC and the culture was induced with 0.5 mM IPTG. The cells were allowed to grow overnight before they were harvested and flash frozen in liquid nitrogen and stored at -80 degC.

## Purification

### Procedure

The lysate was centrifuged at 16,000 rpm for 60 minutes and the supernatant was mixed with 4

mL 50% Ni-NTA beads, and incubated at 4 degC on roller drum for 1 hours. The supernatant was then passed through a gravity column (Poly-Prep, Bio-Rad, Catalog #731-1550) and the beads were washed using 50 mL binding buffer followed by 50mL washing buffer. The protein bound to beads were then eluted using 15 mL elution buffer. The flow-through was collected and loaded onto Supderdex-75 26/60 gel filtration column. Eluted fractions were pooled and concentrated using amicon centrifugal filter (m.w. cut-off 10,000 ). The purity of the proteins was higher than 95% judged by SDS-PAGE.

## **Extraction**

### **Procedure**

Frozen cells from 6L culture were thawed and resuspended in 400 mL extraction buffer with freshly added final concentration of 1mM PMSF/Benzamidine, 0.5% CHAPS and 5U/mL Benzonase (Sigma Catalog # E1014, 250U/ $\mu$ L), and supplemented with 1mL protease inhibitor cocktail (SIGMA Catalog # P8849), and lysed using sonication at 10 seconds 50% duty cycle for 5 minutes at 120W.

**Concentration:** 23.43 mg/mL

### **Ligand**

**MassSpec:** Native: 59018.16, expected 59013.88

**Crystallization:** Stock protein solution was added with a final concentration of 2 mM CaCl<sub>2</sub> and crystallization was setup in sitting drops using Red Wings and SGC-I screens kits. No crystals were observed within one month of setup. Crystal used for data collection was harvested from a drop containing 20% PEG 3350, 0.2M CaCl<sub>2</sub> in the well solution more than 100 days after setup. Paratone was used as cryoprotectant. The original construct includes a RhoGEF, a PH and a C2 domain. Only the C2 domain was crystallized. The protein preparation probably contains residual E.coli protease that cleaved off the RhoGEF and PH domains and left only C2 domain for crystallization. Crystals of the same cell parameters as that of the structure solved one can also be produced in the presence of 1:100 papain (w/w).

### **NMR Spectroscopy:**

### **Data Collection:**

### **Data Processing:**