

# Tg-UBC: *Toxoplasma gondii* ubiquitin conjugating enzyme (TgTwinScan\_2218)

PDB:2AYV

## Revision

**Revision Type:**created

**Revised by:**created

**Revision Date:**created

**Entry Clone Accession:**[TgTwinScan\\_2218](#)

**Entry Clone Source:***Toxoplasma gondii* cDNA from the lab of [Prof. David Sibley](#)

**SGC Clone Accession:**TgTwinScan\_2218:A2-A146; plate MAC00T:D2

**Tag:**N-terminal His-tag with integrated TEV protease site: MGSSHHHHHHSSGRENLYFQ\*G

**Host:**E. coli BL21-(DE3)-CodonPlus-RIL from Stratagene

## Construct

**Prelude:**

**Sequence:**

mgsshhhhhssgrenlyfqgALKRINKELNDLSKDPPTNCSAGVPGDDMFHWQATIMGPEDSPYSGGVFFLNHFPSDYPFKPPKV  
NFTTKIYHPNINSQGAICLDILKQWSPAL TISKVLLSISLLTDPNPDDPLVPEIAHLYKSDRMRYDQTAREWSQKYA

**Vector:**p15TV-L

## Growth

**Medium:**Terrific Broth (TB)

**Antibiotics:**50 microG/mL kanamycin and 25 microG/mL chloramphenicol

**Procedure:**A single colony was inoculated into 10 mL of LB with of Antibiotics and incubated with shaking at 250 rpm overnight at 37 degC. The culture was transferred into 50 mL of TB with Antibiotics in a 250 mL shaking flask and incubated at 37 degC for 3 hours. The culture was then transferred into 1.8 L of above-specified growth medium with Antibiotics and 0.3 mL of antifoam (Sigma) in a 2L bottle and cultured using the LEX system to an OD600 of 2.5, cooled to 15 degC and induced with 0.5 mM isopropyl-1-thio-D-galactopyranoside (IPTG) overnight at 15 degC.

## Purification

**Procedure**

The cleared cell lysate was loaded onto a DE52 (Whatman) column packed with 10 g of resin (previously activated with 2.5 M NaCl and equilibrated with Binding Buffer), and subsequently onto a 2.5 mL Ni-NTA column at approximately 1.5 mL/min. When all the lysate was loaded, 20 mL of Binding Buffer was added to the DE52 column. Then the Ni-NTA column was washed with 200 mL of Wash Buffer at 2 □ 2.5 mL/min. After washing, the protein was eluted from the Ni-NTA column with 15 mL of Elution Buffer. EDTA was added immediately to 1 mM. DTT was

then added to 1 mM 15 minutes later. The protein was dialysed in a dialysis cassette (Pierce) overnight dialysis against Crystal Buffer. The following day it was concentrated using a 15 mL Amicon Ultra centrifugal filter device from Millipore (10 kD cutoff). Protein concentration was measured by means of absorbance at OD280. The concentrated protein was flash frozen in N2(l) in 150 microL aliquots and stored at -80 degC.

## **Extraction**

### **Procedure**

Cells were resuspended to approximately 40 mL/L of cell culture in Binding Buffer with protease inhibitor (1 mM benzamidine-HCl and 1 mM phenylmethyl sulfonyl fluoride, PMSF).

Resuspended pellets stored at -80 degC were thawed overnight at 4 degC on the day before purification. Prior to mechanical lysis, each pellet from 1 L of culture was pretreated with 0.5% CHAPS and 500 units of benzonase for 40 minutes at room temperature. Cells were mechanically lysed with a microfluidizer (Microfluidizer Processor, M-110EH) at approximately 18,000 psi. The cell lysate was centrifuged at ~75,000 x g for 20 minutes at 10 degC.

**Concentration:** 26 mg/mL

### **Ligand**

#### **MassSpec:**

**Crystallization:** The protein was crystallized by means of sitting drop vapor diffusion in a 96-well Intelli-Plate. The plate was set with 1 microL cleaved protein and 1 microL buffer in each drop, and 100 microL reservoir volume per well. Crystals grew in two days in 3.5M Na formate, 0.1M NaOAc pH 4.6 at 20 degC.

#### **NMR Spectroscopy:**

#### **Data Collection:**

#### **Data Processing:**