

UEV3

PDB:3DL2

Revision

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Entry Clone Source:Open Biosystems

SGC Clone Accession:ubc81.171.471; plate SDC076H08

Tag:MGSSHHHHHHSSGLVPR*GS

Host:*E. coli* BL21 (DE3)

Construct

Prelude:

Sequence:

MGSSHHHHHHSSGLVPR*GSSKSWANHENKTVNKITVGGGELGIACTLAISAKGIADRLVLLDLSEGTKGATMDLEIFNLPNVEISKDLSASAHSKVVIFTVNSLGSSQSYLDVVQSNVDMFRALVPALGHYSQHSVLLVASQPVEIMTYVTWKLSTFPANRVIGIGCNLDSQRLQYIITNVLKAQTSKGKEVWVIGEQQEDKVLTWSGQEEVVSHTSQVQLSNRAMELLRVKGQRSWSVGLSVADMVDSIVNNKKKVHSV
SALAKGYDINSEVFLSLPCILGTNGVSEVIKTTLKEDTVTEKLQSSASSIHS LQQQLKL

Vector:PET28a-LIC

Growth

Medium:Terrific Broth (TB) in the presence of 50 µg/ml of kanamycin

Antibiotics:

Procedure:The protein was expressed in *E. coli* BL21 (DE3) grown in growth medium at 37°C to an OD₆₀₀ of 5.5. Protein expression was induced 0.05 mM isopropyl-1-thio-D-galactopyranoside (IPTG) overnight at 15°C. The culture was centrifuged (12,000 x g, 15 minutes) and cell pellets collected and stored at -80°C.

Purification

Procedure

The cleared lysate was loaded onto a 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 10 mL Wash buffer A. The protein was eluted with 6 mL Elution buffer. His-tag was removed by incubation of the protein with thrombin (1 U per mg protein, overnight at 4 °C). The protein was further purified by gel filtration on a HighLoad 16/60 Superdex 200 column (GE Healthcare, Amersham) equilibrated with Gel Filtration buffer. Fractions containing protein (analyzed by SDS-PAGE) were pooled and concentrated by ultrafiltration using an Amicon Ultra centrifugal filter with 10 kD cutoff.

Extraction

Procedure

The cell pellet from a 2 L culture was resuspended in 50 ml Lysis buffer, lysed using a Microfluidizer at 18,000 PSI, and cleared by centrifugation at 40,000 x g for 30 min.

Concentration:

Ligand

MassSpec:

Crystallization: Crystals were grown in hanging drops by mixing 2 ul UBE2V3 solution (6 mg/ml) with 2 ul well solution (1.6 M Na/KPO₄, pH 6.5, 0.2 M NaCl, 2 mM DTT) at 18°C. The crystals were cryoprotected by immersion in the well solution mixed in 1:1 ratio with a water solution containing 20% (w/v) sucrose, 4% (w/v) glucose, 18% (v/v) glycerol and 18% (v/v) ethylene glycol.

NMR Spectroscopy:

Data Collection:

Data Processing: