

UGDH

PDB:2Q3E

Revision

Revision Type:created

Revised by:created

Revision Date:created

Entry Clone Accession:gi|4507813

Entry Clone Source:MGC

SGC Clone Accession:

Tag:N terminal SET1-tag, SBP-tag, TEV protease site (indicated by):

mdpeeasvtsteetltpaqeartraankarkeaelaaataeqtsdekttgwrgghvvveglageleqlrarlehhpqgqrepsggcklglt
tenlyfq*sm

Host:BL21(DE3)-R3-pRARE2

Construct

Prelude:

Sequence:

mdpeeasvtsteetltpaqeartraankarkeaelaaataeqtsdekttgwrgghvvveglageleqlrarlehhpqgqrepsggcklglt
tenlyfq*smFEIKKICCIGAGYVG GPTCSVIAHMCPEIRVTVDNESRINAWSPTLPIYEPGLKEVVESCRGKNLFFS
TNIDDAIKEADL VFISVNTPKTYGMGKGRAADLKYIEACARRIVQNSNGYKIVTEKSTV PVRAAESIRRIFDANTKPNLNQVLS
NPE FLAEGTAIKDLKNPDRV LIGGDETPEGQR AVQALCAVYEHWVPREKILTTNTWSSELS KLAANAFLAQRISSINSISALCE
ATGADV EEVATAIGMDQRIGNKFLKASVGFGGSCF QKDVLNLVYLCEALNLPEVARYWQQVIDM NDYQRRRFASRIIDS LFNTV
TDKKIAILG FAFKKDTGDTRESSSIYISKYLDEGAHL HIYDPKVPREQIVVDSLHSPGVSEDDQVSR LVTISKDPYEACDGAHA
VVICTEWDMFKE LDYERIHKKMLKPAFIFDGRRLVDGLHNE LQTIGFQIETIGKKV

Vector:pBEN1-SGC.

Growth

Medium:

Antibiotics:

Procedure:10ml of an overnight culture was added into 1L of TB supplemented with 50 µg/ml of Ampicillin (total 8L). The cells were cultured at 37°C until the OD600 reached 1.8; the temperature was then decreased to 18°C. IPTG was added to 0.5mM final concentration, and the culture was continued at 18°C overnight.

Purification

Procedure

Column 1: Streptavidin agarose

Column 2: Superdex 200 Hiload 16 60

The Streptavidin agarose (10 ml) was thoroughly washed with 1L of binding buffer. The supernatant was incubated with the washed resin at 4°C for 1 hour by gentle rotation and packed into the column. The column was washed with 2x 50 ml of washing buffer. The protein was eluted with 10 ml of elution buffer and collected in 1.5 ml fractions.

An AKTA Purifier system (GE Healthcare) was used and run at 4°C. Fractions were analysed by SDS - PAGE and the most purified fractions were collected, concentrated and utilized for crystallization.

Enzymatic treatment: For tag removal, 100 m l of TEV protease (6mg/ml) was added to the sample, incubated at 4°C overnight, before a final round of gel filtration chromatography. Purity was analyzed by SDS/PAGE and ESI mass spectrometry.

Extraction

Procedure

The cells were harvested by centrifugation at 4,000 g for 10 min. The pellet from 1 L culture was resuspended in 25 ml of extraction buffer. The sample was sonicated and then centrifuged at 37,500 g. The supernatant was kept for further purification.

Concentration: 20 mg/ml.

Ligand

MassSpec: Experimental mass (ESI TOF mass spectrometry) 52018 Da; (52020 Da theoretical mass)

Crystallization: Crystals were grown by vapor diffusion in sitting drops at 4°C. Before setting up the experiment 5 mM NADH and 10 mM UDP-glucose were added to the protein. A sitting drop consisting of 150 nl protein (20 mg/ml) and 150 nl well solution was equilibrated against well solution containing 16% PEG 3350, 8% ethylene glycol, 160 mM NaBr, 80 mM BTP 6.5. The crystal was transferred to a cryo-protectant consisting of well solution supplemented with 15% ethylene glycol and 3 mM UDP-glucose, mounted in a loop, and flash-cooled in liquid nitrogen.

NMR Spectroscopy:

Data Collection: Data Collection, Resolution: 2.0 Å, X-ray source: Synchrotron SLS-X10SA, single wavelength. The structure was solved by molecular replacement using the structure 2o3j as a search model.

Data Processing: