

Np95-like ring finger (NIRF) protein isoform b

PDB:1Z6U

Revision

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Entry Clone Accession:

Entry Clone Source:MGC

SGC Clone Accession:ubh14.672.802; plate SDC013:C4

Tag:N-terminal His-tag with integrated thrombin-cleavage site MGSSHHHHHHSSGLVPRGS.

Host:E.coli BL21 (DE3)

Construct

Prelude:

Sequence:

gsPSASKVYKASDSAEAEAFQLTPQQQLIREDCQNQKLWDEVLSHLVEGPNFLKKLEQSFMCVCCQELVYQPVVTECFHNVCKDC
LQRSFKAQVFSCPACRHDLGQNYIMIPNEILQTLDDLFFPGYSKGR

Vector:p28a-LIC

Growth

Medium:TB

Antibiotics:

Procedure:Using the SGC LEX bubbling system, Np95-like ring finger protein was expressed in E. coli BL21 (DE3) grown in Terrific Broth (TB) in the presence of 50 µg/mL of kanamycin at 37°C to an OD₆₀₀ of 7.5. Cells were then induced by isopropyl-1-thio-D-galactopyranoside (IPTG), final concentration 0.05 mM, and incubated overnight at 15°C. The culture was centrifuged and the cell pellets were collected and stored at -80°C.

Purification

Procedure

The cleared lysate was loaded onto a TALON metal-affinity resin column from BD Biosciences at 4°C. The column was washed with wash buffer A (10mM Tris-HCl pH 8.0, 0.5 M NaCl, 5% glycerol, 10 mM imidazole, 1 mM β -mercaptoethanol), wash buffer B (same as wash buffer A but containing 0.05% Tween 20) and again wash buffer A, and the protein was eluted with 10 mM Tris-HCl, pH 8.0, 0.5 M NaCl, 5% glycerol, 200 mM imidazole, 1 mM β -mercaptoethanol. The protein was further purified by gel filtration on a HighLoad 16/60 Superdex 200 column (GE Healthcare, Amersham) equilibrated with 20 mM Tris-HCl, pH 8.0, 0.5 M NaCl, 5% glycerol, 2mM dithiothreitol and concentrated by ultrafiltration.

Extraction

Procedure

The cell pellet was resuspended in lysis buffer (10 mM Tris-HCl, pH 8.0, 0.5 M NaCl, 5% glycerol, 2 mM imidazole, 1 mM β -mercaptoethanol) with protease inhibitor (0.1mM phenylmethyl sulfonyl fluoride, PMSF) and lysed using Microfluidizer. The lysate was cleared by centrifugation.

Concentration: 68 mg/mL

Ligand

MassSpec:

Crystallization: Purified Np95-like ring finger protein 3 was crystallized using the hanging drop vapor diffusion method. Crystals grew when the protein (34 mg/mL) was mixed with the reservoir solution in a 1:1 volume ratio, and the drop was equilibrated against a reservoir solution containing 0.9M sodium citrate, 0.1M Tris-HCl, pH 8.5.

NMR Spectroscopy:

Data Collection:

Data Processing: