

# DDI1

**PDB:**3S8I

**Entry Clone Accession:**ddi1.BC022017.MGC.AT25-B1.pBluescriptR

**Entry Clone Source:**Mammalian Gene Collection

**SGC Clone Accession:**ddi1.0239-0367 (SDC229E10)

**Tag:**NtMGSSHHHHHHSSGLVPRGS

**Host:**Competent BL21 (DE3) cells (Invitrogen, C6000-03)

**Vector:**pET28a-LIC vector (pET28a-LIC: GenBank, EF442785)

## Sequence:

MGSSHHHHHHSSGLVPRGSGQVTMLYINCKVNGHPLKAFVDSGAQMTIMSQACAERCNIMRLVDRRWAGVAKGVGTQRIIGRVHLAQ  
IQIEGDFLQCSFSILEDQPMDDLGLDMLRRHQCSIDLKKNVLVIGTTGTQTYFLPEGELP

## Growth

**Medium:** TB (Sigma, T0918) supplemented with 150 mM glycerol, 100  $\mu$ M kanamycin and 600  $\mu$ l antifoam 204 (Sigma A-8311)

**Procedure:** Competent BL21 (DE3) cells (Invitrogen, C6000-03) were transformed and grown using the LEX system (HarbingerBiotech) at 37 °C in 1L bottles (VWR, 89000-242) containing 900 ml of growth medium. When the OD<sub>600</sub> reached a value of about 6.0, the temperature was reduced to 15 °C, and one hour later the culture was induced with 100  $\mu$ M IPTG (BioShop, IPT001) and incubated overnight (16 hours) at 15 °C.

## Purification

**Procedure:** Protein was purified using the Streamline purification system (1). Briefly, bacterial cultures were directly microfluidized with the outflow connected to a novel column assembly, containing IMAC beads; a few hundred milliliters of regular buffer was also fluidized to wash the beads, and columns were eluted with imidazole by gravity. To each 1L culture bottle the following was sequentially added: NaOH (final pH ~7.5), imidazole (final 8mM), and BME (final 1 mM). Each bottle was fluidized through a Microfluidizer (model M110-EH with H10Z ceramic chamber performed at about 15,000 psi) with the outflow directly connected to a special nozzle-column containing 3 ml of settled HisLink (Promega, V8821). Approximately 200 ml fluidizer buffer was also fluidized to wash the beads. Columns were gravity-washed with 20 mL wash buffer, and protein was gravity-eluted with 8.0 ml elution buffer. Eluate was dialyzed overnight at 4 oC against about 100 volumes of dialyses buffer. Samples were concentrated using a 10 KDa MW cut-off concentrator (Millipore, UFC900524) at 3500 xg to a final value of 15 mg/ml. Protein yield was 7 mg per liter of bacterial culture.

Alenkin D, Yermekbayeva L, Mujib S, Vesterberg A, Newman E, Yamazaki K, Cossar D, Dhe-Paganon S. A centrifugation-free high-throughput protein purification system using in-line

microfluidization. *Protein Expr Purif.* 2011.

## Structure Determination

**MassSpec:** MW = 16392 g/mol

**Crystallization:** Crystals were grown at 18 °C in hanging drop plates (Hampton, HR3-170) by mixing equal volumes of protein (10 mg/ml) and Crystallization Buffer (21% PEG3350, 0.2 M NH<sub>4</sub>(H<sub>2</sub>PO<sub>4</sub>), 0.1 M Bis-Tris pH 6.0). Suitable crystals were cryoprotected by immersion in well solution supplemented with 20 % (v/v) ethylene glycol prior to dunking and storage in liquid nitrogen.