

L3MBTL

PDB:2PQW

Revision

Revision Type:created

Revised by:created

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Entry Clone Accession:NP_056293

Entry Clone Source:

SGC Clone Accession:L3MBTL1_3:A3-APC053

Tag:N-terminal hexahistidine tag with integrated TEV protease cleavage site:
mhhhhhssgrenlyfq*g

Host:E Coli BL21(DE3) RIL CodonPlus

Construct

Prelude:

Sequence:

GEKKECWSWESYLEEQKAITAPVSLFQDSQAVTHNKNQKLGKMLEGIDPQHPSMYFILTVAEVCGYRLRLHFDGYSECHDFWVNAN
SPDIHPAGWFEKTGHKLQPPKGYKEEFWSQYLRSTRAQAAPKHLFVSQSHSPPLGFQVGMKLEAVDRMNPSLVCVASVTDVVD
RFLVHFDNWDDTYDYWCDPSSPYIHPVGWCQKQKPLTPPDYDPDNFCWEKYLEETGASAVPTWAFKVRPPHSFLVNMKLEAVDR
RNPALIRVASVEDVEDHRIKIHFDDGWSHGDFWIDADHPDIHPAGWCSKTGHPLQPPLGP

Vector:p28a-MHL

Growth

Medium:TB

Antibiotics:

Procedure:A glycerol stock was used to inoculate 20 mL LB media containing 50 µg/mL kanamycin. The culture was grown overnight at 37°C with shaking. The next day this starter culture was used to inoculate 2L of TB medium which contained 50 µg/mL kanamycin. The culture was grown in LEX at 37°C to OD600 ~1.1 and was induced with the addition of 0.5 mM IPTG. The temperature was reduced to 18°C and the culture was incubated for a further 16 hours before harvesting the cells.

Purification

Procedure

Column 1: Affinity purification, open Ni-NTA column
Procedure: The supernatant was incubated with 6mL of 50% slurry Ni-NTA beads by rocking. After 1 hour incubation at 4°C, the beads were washed with 50 mL of lysis buffer. The protein was eluted using ~20mL EB.
Column 2: Gel filtration, HiLoad 26/60 Superdex 200 Prep Grade
Procedure: The eluent from Ni column was

loaded onto the gel filtration column in GF buffer at 2.5 mL/min, fraction size 4mL. The fractions containing protein were identified on a SDS-PAGE gel.

Extraction

Procedure

Cells were harvested by centrifugation and pellets were stored in -80°C. Prior to purification, the cell pellet was resuspended in lysis buffer. Cells were disrupted by sonication (10 minutes twice) and samples were centrifuged for 60 min at 70000 g. The soluble fraction was subjected to further purification by affinity and size exclusion chromatography.

Concentration:20mg/ml

Ligand

MassSpec:

Crystallization:100mM NaAc 4.6, 100mM NaAC, 5% PEG 4K

NMR Spectroscopy:

Data Collection:

Data Processing: