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# OPEN ACCESS



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Protocol status: Working We use this protocol and it's working

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## GFP-Clu-tails purification from Escherichia coli cells

Andreas Bracher<sup>1</sup>, F Ulrich Hartl<sup>1</sup>

<sup>1</sup>Department of Cellular Biochemistry, Max Planck Institute of Biochemistry



Patricia Yuste-Checa

#### **ABSTRACT**

This protocol details how to purify the fusion protein GFP-Clu-tails from Escherichia coli.

#### **ATTACHMENTS**

GFP-Clu-tails purification protocol\_protocols.io.docx

#### **MATERIALS**

#### **Buffers**

Binding buffer: PBS + [M] 20 millimolar (mM) imidazole

Elution buffer: PBS + [M] 250 millimolar (mM) imidazole

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# His<sub>6</sub>-Ubiquitin-GFP-Clu-tails expression and cell lysis 1 Express His<sub>6</sub>-Ubiquitin-GFP-Clu-tails in E. coli Bl21 (DE3) codon+RIL cells cultured in A 4 L LB Medium Overnight at \$\mathbb{S}\$ 18 °C with [M] 0.25 millimolar (mM) IPTG. 2 Centrifuge culture and keep pellet. 3 Re-suspend pellet in A 70 mL volume of ice-chilled Binding buffer, add Complete protease inhibitor cocktail (Roche) and [M] 1 millimolar (mM) phenylmethylsulfonyl fluoride (PMSF). 4 2m Lyse cells by ultrasonication in ice bath (15 cycles of 00:00:25 ultrasonication with 00:01:35 intermittent cooling). 5 Clear lysate by centrifugation at 22000 rpm in a JA25.50 rotor at 4 °C

# Ni2+-chelating Sepharose metal affinity chromatography

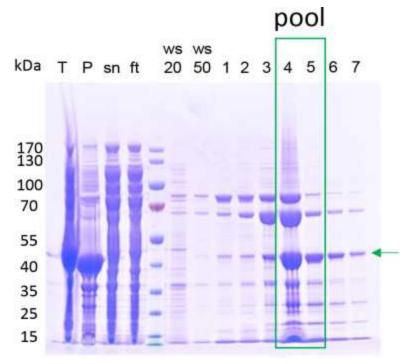
- Load supernatant onto a R B mL Ni-chelating Sepharose (Cytiva 17-0575-01) column previously equilibrated with binding buffer by gravity flow at R 4 °C.
- Wash the column with 5 CV of ice-chilled Binding buffer.



8 Wash the column with 2 CV of ice-chilled PBS + [M] 50 Molarity (m) imdiazole.



- Elute His6-Ubiquitin-GFP-Clu-tails protein with 6x 5 mL of ice-chilled Elution buffer. Collect fractions of 5 mL volume. Bright green color indicates presence of His6-Ubiquitin-GFP-Clu-tails. Store fractions on ice.
- 10 Analyze eluted fractions by SDS-PAGE and Coomassie blue staining.



## Protease cleavage and removal of protease

Pool fractions containing His6-Ubiquitin-GFP-Clu-tails peak (here fractions 4 and 5). Cleave off His6-Ubiquitin with His-tagged Usp2 during 04:00:00 On ice

4h

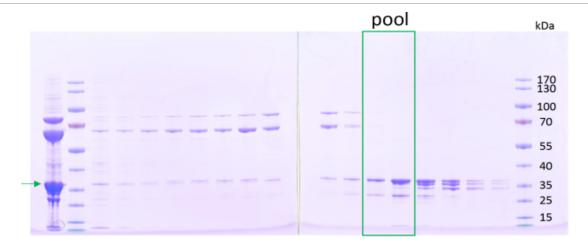
- Exchange protein buffer to Binding buffer with a HiPrep<sup>™</sup> 26/10 Desalting column (Cytiva 17-5087-01) equilibrated with Binding buffer at a containing fractions.
- Remove uncleaved material and His-tagged Usp2 by metal affinity chromatography as above. Collect flow-through and wash fractions with bright green fluorescence.
- Concentrate to L 1.5 mL volume by ultrafiltration using 10 kDa cut-off spin concentrator at L 4 °C

# Size exclusion chromatography on HiLoad 16/600 Superdex-200 (Cytiva 28...

- Apply concentrate on a HiLoad 16/600 Superdex-200 (Cytiva 28-9893-35) column equilibrated with PBS.

  Develop the column at 4 °C and collect 4 3 mL fractions.
- 16 Analyze eluted fractions by SDS-PAGE and Coomassie blue staining.

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Merge fractions 16 and 17. Concentrate to 0.7 mL volume by ultrafiltration using 10 kDa cut-off spin concentrator at 4 °C, aliquot and flash-freeze purified GFP-Clu-tails in liquid nitrogen for storage at -70 °C.

#### Note

Concentrations were determined by absorbance at 488 nm (eGFP!) using absorbance coefficients of  $61,000 \, \text{M}^{-1} \, \text{cm}^{-1}$  or  $1.45 \, \text{Lg}^{-1} \, \text{cm}^{-1}$ .

**Approximate yield**: From 4 L of culture around 2.5 mg of GFP-Clu-tails were obtained.

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