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Purification of CCPG1-GST

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Elias Adriaenssens¹

¹Sascha Martens lab, University of Vienna, Max Perutz Labs - Vienna



Elias Adriaenssens

Sascha Martens lab, University of Vienna, Max Perutz Labs - ...

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We use this protocol and it's working

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Abstract

This protocol details the purification of CCPG1-GST, its analysis by SDS-PAGE and Coomassie staining.

Materials

Lysis Buffer:

| A | B |
|--|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 300 mM |
| Triton X-100 | 1% |
| Glycerol | 5% |
| MgCl ₂ | 2 mM |
| DTT | 1 mM |
| β-mercaptoethanol | 2 mM |
| cOmplete EDTA-free protease inhibitors (Roche) | |
| CIP protease inhibitor (Sigma) | |
| DNase (Sigma) | |

Wash Buffer:

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 300 mM |
| DTT | 1 mM |


Salt wash Buffer:

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 700 mM |
| DTT | 1 mM |

SEC Buffer:

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 25 mM |
| NaCl | 300 mM |
| DTT | 1 mM |



-  Rosetta™(DE3)pLysS Competent Cells - Novagen **Merck Catalog #70956-4**
- pET-Duet1 vector encoding CCPG1 (1-212aa)-thrombin-GST (available from Addgene)



Purification procedure

1d 2h 45m 30s

- 1 To purify CCPG1-GST, fuse the cytosol-exposed domain of CCPG1 (1-212aa) to a C-terminal GST-tag through gene synthesis and Gibson cloning into a pET-DUET1 vector.
- 2 After the transformation of the pET-DUET1 vector encoding CCPG1-GST in *E. coli* Rosetta pLysS cells (Novagen Cat# 70956-4), grow cells in 2x Tryptone Yeast extract (TY) medium at 37 °C until an OD₆₀₀ of 0.4 and then continue at 18 °C .
- 3 Once the cells reach an OD₆₀₀ of 0.8, induce the protein expression with 100 micromolar (μM) isopropyl β-D-1-thiogalactopyranoside (IPTG) for 16:00:00 at 18 °C .
- 4 Collect the cells by centrifugation and resuspend in lysis buffer.

16h



Lysis Buffer:

| A | B |
|--|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 300 mM |
| Triton X-100 | 1% |
| Glycerol | 5% |
| MgCl ₂ | 2 mM |
| DTT | 1 mM |
| β-mercaptoethanol | 2 mM |
| cOmplete EDTA-free protease inhibitors (Roche) | |
| CIP protease inhibitor (Sigma) | |
| DNase (Sigma) | |

- 5 Sonicate the cell lysates twice for 00:00:30 and clear by centrifugation at 18000 rpm, 4°C, 00:45:00 in a SORVAL RC6+ centrifuge with an F21S-8x50Y rotor (Thermo Scientific).

45m 30s



- 6 Collect the supernatant and incubate with pre-equilibrated Glutathione Sepharose 4B beads (GE Healthcare) for  02:00:00 at  4 °C with gentle shaking to bind CCPG1-GST.

2h




- 7 Centrifuge the samples to pellet the beads and remove the unbound lysate. Then wash the beads twice with wash buffer, once with high salt wash buffer, and two more times with wash buffer.

**Wash Buffer:**

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 300 mM |
| DTT | 1 mM |

Salt wash Buffer:


| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 700 mM |
| DTT | 1 mM |

- 8 Incubate the beads  Overnight with  4 mL of [M] 50 millimolar (mM) reduced glutathione dissolved in wash buffer at  4 °C , to elute CCPG1-GST from the beads.

8h

**Wash Buffer:**

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 50 mM |
| NaCl | 300 mM |
| DTT | 1 mM |

- 9 To collect the supernatant, collect the beads by centrifugation. Wash the beads twice with  4 mL of wash buffer, and collect the supernatant each time.





10 Pool the supernatant fractions, filter through a 0.45 µm syringe filter, concentrate with 30 kDa cut-off Amicon filter (Merck Millipore), and load onto a pre-equilibrated Superdex 200 Increase 10/300 GL column (Cytiva).

11 Elute the proteins with SEC buffer.

SEC Buffer:

| A | B |
|-----------------|--------|
| Tris-HCl pH 7.4 | 25 mM |
| NaCl | 300 mM |
| DTT | 1 mM |

12 Analyze the fractions by SDS-PAGE and Coomassie staining. Pool the fractions containing purified CCPG1-GST.



13 After concentrating the purified protein, aliquot the protein and snap-frozen in liquid nitrogen.