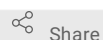


May 24, 2021

# Expression and purification protocol of *Homo sapiens* E2-like enzyme ATG3

Eleonora Turco<sup>1</sup>, Dorotea Fracchiolla<sup>2</sup><sup>1</sup>Sascha Martens Lab – University of Vienna, Max Perutz Labs, Vienna;<sup>2</sup>Sascha Martens Lab – University of Vienna, Max Perutz Labs, Vienna

1 Works for me



Share

[dx.doi.org/10.17504/protocols.io.btgknjuw](https://dx.doi.org/10.17504/protocols.io.btgknjuw)Dorotea Fracchiolla  
Team Hurley

## ABSTRACT

This protocol describes expression and purification procedures for obtaining human recombinant autophagy E2-like enzyme ATG3 (ATG, AuTophagy-related protein) of the ATG8 ubiquitin-like conjugation system.

## THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

A PI3K-WIP1 positive feedback loop allosterically activates LC3 lipidation in autophagy - Published in 10.1083/jcb.201912098

## DOI

[dx.doi.org/10.17504/protocols.io.btgknjuw](https://dx.doi.org/10.17504/protocols.io.btgknjuw)

## PROTOCOL CITATION

Eleonora Turco, Dorotea Fracchiolla 2021. Expression and purification protocol of *Homo sapiens* E2-like enzyme ATG3. **protocols.io**  
<https://dx.doi.org/10.17504/protocols.io.btgknjuw>

## MANUSCRIPT CITATION please remember to cite the following publication along with this protocol

A PI3K-WIP1 positive feedback loop allosterically activates LC3 lipidation in autophagy - Published in 10.1083/jcb.201912098

## KEYWORDS

ATG3, expression, purification, recombinant protein, *Homo sapiens*

## LICENSE

This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

## IMAGE ATTRIBUTION

© Dorotea Fracchiolla

## CREATED

Mar 18, 2021

## LAST MODIFIED

May 24, 2021

## OWNERSHIP HISTORY

Mar 18, 2021  Liz Brydon Protocols.io

May 06, 2021  Dorotea Fracchiolla Team Hurley

## PROTOCOL INTEGER ID

48364

## GUIDELINES

### General information

Insert: *Homo sapiens* ATG3, NP\_071933.2; Expression system: *E. Coli* Rosetta pLyss; plasmid origin: Sascha Martens Lab, Addgene 169079, lab internal construct database number SMC861; backbone: pET-Duet1; plasmid resistance: Ampicillin; tags & cleavage sites: N-term 6xHis, followed by Tobacco Etch Virus (TEV) cleavage site, ATG3 ORF. Ext coeff: 45840 M-1 cm-1, MW 35.8 kDa.

## MATERIALS TEXT

### Materials and Reagents

- *Escherichia coli* Rosetta pLyss cells
- Luria Bertani (LB) medium with antibiotics (final conc. 50µg/ml Ampicillin, 34µg/ml Chloramphenicol)
- IPTG (isopropyl-β-D-thiogalactopyranoside)
- 37°C shaker incubator
- sterile flasks/sterile pipettes
- tip sonicator
- **Lysis Buffer:** 50mM Hepes pH=7.5; 300mM NaCl, 10mM Imidazole, 2mM MgCl<sub>2</sub>, 2mM β-mercaptoethanol, 1mM Pefablock, Complete Protease Inhibitors (EDTA-free CIP tablet, Roche), DNase (Sigma).
- **Buffer A:** 50mM Hepes pH=7.5, 300mM NaCl, 10mM Imidazole (filtered and degassed) + 1mM β-mercaptoethanol
- **Buffer B:** 50mM Hepes pH=7.5, 300mM NaCl, 300mM Imidazole (filtered and degassed) + 1mM β-mercaptoethanol
- **Size Exclusion Chromatography (SEC) Buffer:** 25mM Hepes pH=7.5, 150mM NaCl (filtered and degassed) + 1mM DTT (Dithiothreitol)
- Note: all purification buffers are filtered and degassed. Reducing agents (β-mercaptoethanol and Dithiothreitol) are added after degassing step.

**Columns:** - HT 5ml column (GE Healthcare)  
- S75\_16/60 (GE Healthcare)



**Gels:** 10% SDS-PAGE

## SAFETY WARNINGS




For hazard information and safety warnings, please refer to the SDS (Safety Data Sheet).












## Protein Expression

1 

Transform plasmid DNA (Addgene 169079, SMC861) into *E. Coli* Rosetta pLyss cells and plate on Ampicillin/Chloramphenicol LB agar plate for  **Overnight** growth at  **37 °C**.










2 

The following day, inoculate a  **100 mL LB + Amp/Cam pre-culture** with 1-2 colonies and grow  **Overnight** at  **37 °C** shaking.

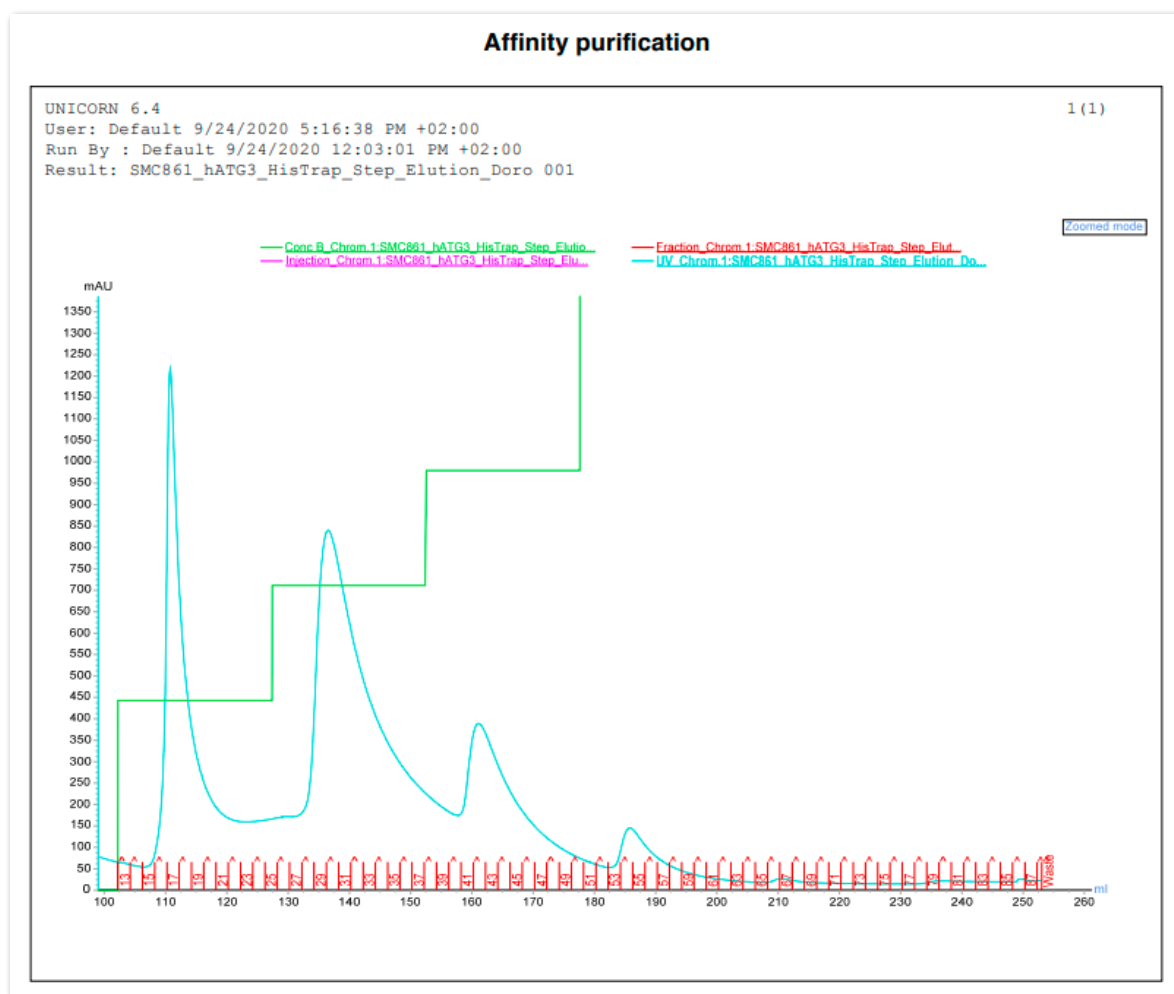
- 3 The following day, use  **20 mL pre-culture** to inoculate  **1 L LB medium + Amp/Cam** at  **37 °C** until an OD<sub>600</sub> (Optical Density at 600nm) of 0.4 is reached.
- 4 Cool down the culture to  **18 °C** and grow until OD<sub>600</sub> = 0.8.
- 5 Induce protein expression with  **100 Micromolar (μM) IPTG** and keep shaking for a further  **16:00:00** at <sup>16h</sup>  
 **18 °C**.
- 6  15m  
Pellet cells at  **4000 rpm, 4°C, 00:15:00** in a Sorvall RC6+ centrifuge (Thermo Scientific), discard supernatant and resuspend pellets in ice cold lysis buffer (25 ml/1 lt culture).
- 7   
Flash freeze resuspended pellets in liquid nitrogen and store at  **-80 °C** until purification.

#### Protein Purification

30m 30s

- 8 Perform His-Trap affinity purification followed by Size Exclusion Chromatography.
- 9 Cells are lysed via freeze/thaw cycles and sonication: thaw pellet corresponding to  **1 L** culture by freeze/thawing in  **Room temperature** water bath. All following steps are to be executed at  **4 °C** or on ice.
- 10 Lyse cells by sonicating them using an immersion tip Sonicator (**2x**  **00:00:30**). Note: adjust times and intensity <sup>30s</sup> according to the available instrument.
- 11  30m  
Clear lysate by spinning it down in a Beckman centrifuge at  **40000 x g, 4°C, 00:30:00**, **Ti45 Rotor**.
- 12 Filter supernatant through a 0.45 μm filter and inject onto a 5ml HT column operating at  **4 °C** pre-equilibrated in Buffer A at 1ml/min flow rate.
- 13   
Wash column with  **5 column volumes (CV) of Buffer A** at 2 ml/min flow rate to remove unspecific bound proteins.

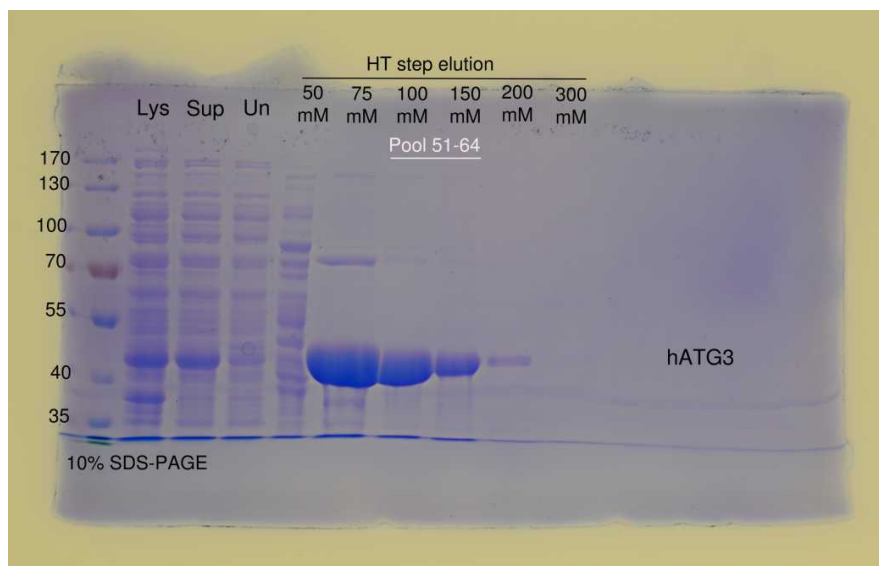
- 14 Elute protein of interest through a step elution gradient in 50mM, 75mM, 100mM, 150mM, 200mM and 300mM Imidazole concentration. Perform elution at 1ml/min flow rate. Collect peak fractions at each step.



## 15

Check peak fractions of each step on a SDS-PAGE (see gel below). Pool and concentrate those containing the protein of interest (usually 100mM and 150mM Imidazole step) by spinning at **4 °C** down in a 10kDa cut-off Amicon Filter to **2 mL final volume**.

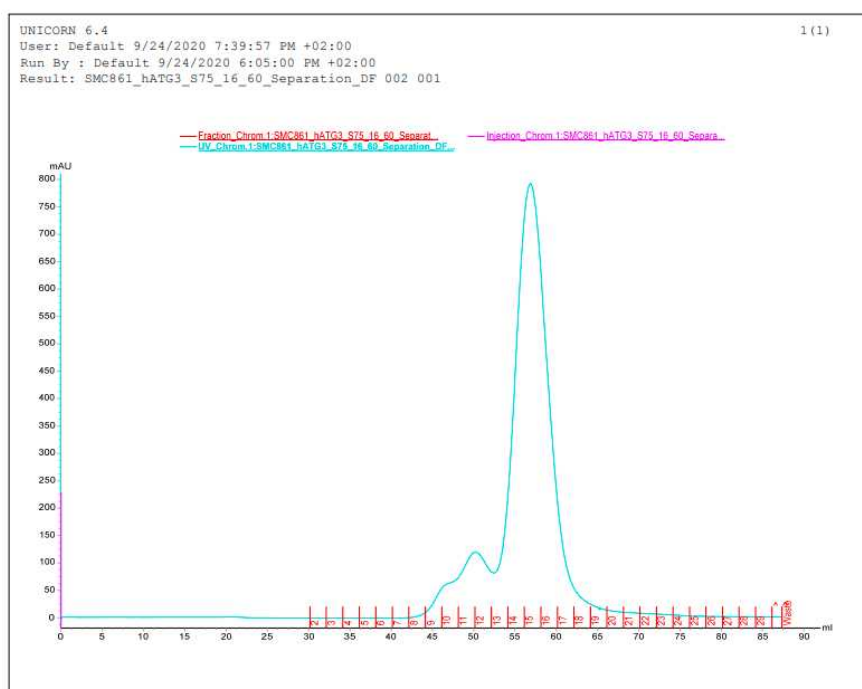
Centrifugation steps are kept short ( **00:05:00** ) to avoid protein local concentration/aggregation on the filter.



Coomassie BB stained gel of His-tag affinity purification for hATG3.

- 16 Inject **2 mL protein** onto a S75\_16/600 column operating at **4 °C** and pre-equilibrated in buffer containing SEC Buffer (see profile below).

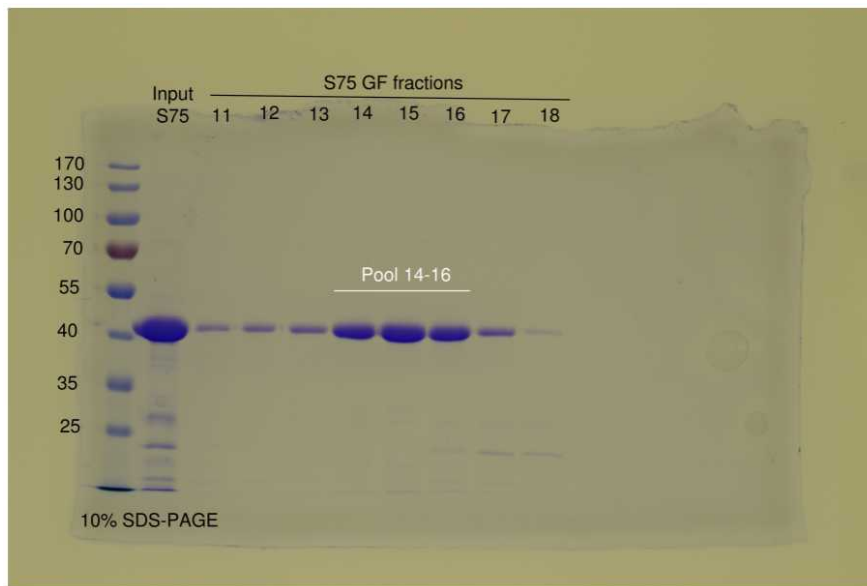
#### Size Exclusion Chromatography



Chromatograph of Size Exclusion purification for hATG3.

- 17 Check fractions on a 10% SDS-PAGE (see gel below). Pool and concentrate those containing the protein of interest down at **4 °C** in a 10kDa cut-off Amicon Filter.

Centrifugation steps are kept short ( **00:05:00** ) to avoid protein local concentration/aqgreagation on the filter.



Coomassie BB stained gel of Size Exclusion Chromatography purification for hATG3.

## 18

Measure protein Absorbance  $A_{280}$  using a Spectrophotometer blanking against SEC buffer (MW = 35,864 kDa ; Extinction coefficient =  $45840\text{M}^{-1}\text{cm}^{-1}$ ).

## 19

Resuspend protein in Glycerol to a final concentration of **[M]30 % (v/v) for glycerol** . Store the protein batch at **-20 °C** . Estimated protein yield: **10 mg per 1 lt culture** . Protein activity is kept for 18 months.