#### I. Introduction

This protocol is provided for **Capturem His-Tagged Purification 96-Well Plate** (Cat. No. 635714), a single-use disposable 96-well plate that allows simple, rapid purification of his-tagged proteins from mammalian or bacterial cell samples in up to 1 ml of clarified lysate per well, using either vacuum filtration or centrifugation. Each well requires a minimum elution volume of  $100 \mu l$ . This plate is suitable for use under native or denaturing conditions, in the presence of additives such as DTT (up to 10 mM),  $\beta ME$  (up to 30 mM), TCEP (up to 5 mM), EDTA (up to 10 mM), or glycerol (see reagent compatibility table for more information).

## II. Materials and Reagents

#### A. Components

• 1 Capturem His-Tagged Purification 96-Well Plate

#### B. Additional Materials Required

- Lysis Buffer: We recommend using xTractor™ Buffer (Cat. Nos. 635625, 635626, or 635671), but you may substitute your standard lysis buffer. We strongly recommend that you begin the purification procedure without using imidazole in your lysis and wash buffers when purifying standard proteins. We only suggest adding imidazole to the wash buffer if you notice significant background binding. xTractor Buffer does not contain imidazole.
- Wash Buffer: 20 mM Na<sub>3</sub>PO<sub>4</sub>, 150 mM NaCl, pH 7.6
- Elution Buffer: 20 mM Na<sub>3</sub>PO<sub>4</sub>, 500 mM NaCl, 500 mM imidazole, pH 7.6
- **96-well collection plates:** Each purification will require four standard 96-well collection plates. These plates should be used throughout the protocol to collect flowthrough and wash samples that will be saved for SDS-PAGE analysis and/or colorimetric protein assays, e.g., Bradford assays.
- **96-well plate filtration device:** The Capturem His-Tagged Purification 96-Well Plate is compatible with standard 96-well plate filtration devices, such as the NucleoVac 96 Vacuum Manifold (Cat. No. 740681).
- **96-well plate centrifuge:** The Capturem His-Tagged Purification 96-Well Plate is compatible with standard 96-well plate centrifuges.
- Multi-channel pipette

# III. Sample Preparation

Express your his-tagged protein(s) of interest in 1 ml cultures grown in 96-well plates or 2–5 ml cultures grown in 6-well plates. Before beginning the protein purification protocol in Section IV, it is necessary to prepare crude cell lysates from each culture and clarify the lysates by centrifugation or filtration. We recommend using xTractor Buffer (Section II) for lysis. Lysis protocols using xTractor Buffer are provided in the xTractor Buffer & xTractor Buffer Kit User Manual. Individual protocols are also available for preparing cell lysates from bacterial, mammalian, baculovirus, and yeast cultures.

#### • Bacterial Cell Samples

We recommend starting with a fresh or frozen cell pellet from 2–5 ml of overnight bacterial culture, which should yield 200–1,000 µl of cleared lysate.

**NOTE:** When working with bacterial cells, the volume of lysate (containing the overexpressed his-tagged protein of interest) is determined by the amount of wet cell pellet obtained from a starting culture volume of 2–5 ml. For example, a log-phase *E. coli* culture (O.D. = 0.6–0.8), induced for 2–4 hr, would be expected to provide  $\sim$ 20–80 mg of bacterial pellet from 2–5 ml of culture. We recommend adding  $\sim$ 400 µl of xTractor Buffer to each  $\sim$ 20 mg of wet bacterial cell pellet.

#### • Mammalian Cell Samples

We recommend starting with a fresh or frozen cell pellet from 2 ml of mammalian cell culture, (e.g., from from a single well of a 6-well culture plate), which should be resuspended in  $100-500 \,\mu l$  xTractor Buffer, yielding up to  $600 \,\mu l$  of cleared lysate). Adherent cells may be harvested by treating them with trypsin and spinning them down, or scraping them directly from the well in the presence of xTractor Buffer. Suspension cells may be harvested by spinning down the liquid culture.

NOTE: When lysing mammalian cells, you may substitute your standard lysis buffer for xTractor Buffer.

## IV. His-Tagged Protein Purification

Purification of his-tagged proteins from the cleared cell lysates prepared in Section III may be performed using vacuum filtration (Protocol A) or centrifugation in a standard 96-well plate centrifuge (Protocol B).

## A. Protocol: Protein Purification Using Vacuum Filtration

- 1. Assemble the Capturem His-Tagged Purification 96-Well Plate in a 96-well plate filtration device together with a 96-well collection plate (supplied by the user—see Section II.B).
- 2. Add 400 μl Lysis Buffer or PBS to each well of the Capturem His-Tagged Purification 96-Well Plate using a multi-channel pipette, to equilibrate the wells. Use vacuum (~0.6–0.7 bar) to run the buffer through the Capturem plate wells into the collection plate. Remove the flowthrough and discard it along with the collection plate. Reassemble the Capturem His-Tagged Purification 96-Well Plate in the 96-well plate filtration device together with a new collection plate.

**NOTE:** The required pressure may vary between 0.2–0.8 bar, depending on the sample. When performing vacuum filtration, use the following formula to convert between different vacuum pressure units:

#### **Pressure Conversion Formula:**

1 mbar =  $100 \text{ Pa} = 0.750 \text{ mm Hg} = 14.504 \text{ x } 10^{-3} \text{ psi} = 0.987 \text{ x } 10^{-3} \text{ atm}$ 

- 3. Load 200–1,000 µl cleared lysate (from Section III) onto each well of the equilibrated Capturem His-Tagged Purification 96-Well Plate. Use a vacuum (~0.6–0.7 bar) to run the buffer through the Capturem plate wells into the collection plate. Save the collection plate containing the lysate flowthrough for protein analysis. Reassemble the Capturem His-Tagged Purification 96-Well Plate in the 96-well plate filtration device together with a new collection plate.
- 4. Add 300 μl Wash Buffer to each well of the Capturem His-Tagged Purification 96-Well Plate. Use a vacuum (~0.6–0.7 bar) to run the buffer through the Capturem plate wells into the collection plate. Save the collection plate containing the lysate flowthrough for protein analysis. Reassemble the the Capturem His-Tagged Purification 96-Well Plate in the 96-well plate filtration device together with a new collection plate.
- 5. Add 300 μl Elution Buffer to each well of the Capturem His-Tagged Purification 96-Well Plate. Use a vacuum (~0.6–0.7 bar) to run the buffer through the Capturem plate wells into the collection plate.

The wells of the collection plate should contain your eluted tagged protein, which is now ready for analysis.

**NOTE:** ≥90% of your tagged protein can be eluted with 100 µl of Elution Buffer.

- 6. Measure the amount of protein in your flowthrough samples from Steps 3 and 4, and your eluate from Step 5, using a Bradford assay or other colorimetric protein analysis method.
- 7. Analyze the samples that were quantified in Step 6 using SDS-PAGE.

## B. Protocol: Protein Purification Using Centrifugation

- 1. Place the Capturem His-Tagged Purification 96-Well Plate securely on top of a 96-well collection plate (supplied by the user—see Section II.B).
- 2. Add 400 μl Lysis Buffer or PBS to each well of the Capturem His-Tagged Purification 96-Well Plate using a multi-channel pipette, to equilibrate the wells. Centrifuge at 2,000g for 3 min at room temperature. Remove the flowthrough and discard it along with the collection plate. Place the Capturem His-Tagged Purification 96-Well Plate securely on top of a new 96-well collection plate.

**NOTE:** When performing centrifugation, use the following formula to convert rpm to centrifugal force units (RCF or g) if the centrifuge does not automatically provide this information:

#### **Centrifugal Force Conversion Formula:**

RCF or  $g = (1.12) x (R) x (rpm/1,000)^2$ 

where R is the distance (in millimeters) from the center of the rotor to the end of the spin bucket when held horizontally and away from the center.

If the solution does not fully drain from the column, perform a second centrifugation at 2,000g for 3 min. If the solution is still not draining completely, refer to Appendix A. Troubleshooting Guide and re-examine your lysate for viscosity, particles or cloudiness. For lysate preparation instructions, refer to the xTractor Buffer and xTractor Buffer Kit User Manual.

- 3. Load 200–1,000 µl cleared lysate (from Section III) onto each well of the equilibrated Capturem His-Tagged Purification 96-Well Plate. Centrifuge at 2,000g for 3 min at room temperature. Save the collection plate containing the lysate flowthrough for protein analysis. Place the Capturem His-Tagged Purification 96-Well Plate securely on top of a new 96-well collection plate.
- 4. Add 300 μl Wash Buffer to each well of the Capturem His-Tagged Purification 96-Well Plate. Centrifuge at 2,000g for 3 min at room temperature. Save the collection plate containing the lysate flowthrough for protein analysis. Place the Capturem His-Tagged Purification 96-Well Plate securely on top of a new 96-well collection plate.
- Add 300 μl Elution Buffer to each well of the Capturem His-Tagged Purification 96-Well Plate.
   Centrifuge at 2,000g for 3 min at room temperature. The wells of the collection plate should contain your eluted tagged protein, which is now ready for analysis.

**NOTE:** ≥90% of your tagged protein can be eluted with 100 µl of Elution Buffer.

- 6. Measure the amount of protein in your flowthrough samples from Steps 3 and 4, and your eluate from Step 5, using a Bradford assay or other colorimetric protein analysis method.
- 7. Analyze the samples that were quantified in Step 6 using SDS-PAGE.

# **Appendix A. Troubleshooting Guide**

| Problem                           | Possible Explanation   | Solution  |
|-----------------------------------|--|---|
| Background bands/<br>low purity   | Nonspecific binding of proteins to the membrane  | <ul> <li>Add an additional wash step after binding with Wash Buffer.</li> <li>Before loading the lysate in Section IV.A or IV.B, include a blocking step between Steps 1 and 2 by adding BSA (100 µg) in a phosphate- or acetate-based buffer at pH 5 and spin at 2,000<i>g</i> for 3 min.</li> </ul>   |
| Low percentage recovery           | The sample contains more histagged protein than the Capturem His-Tagged Purification 96-Well Plate has the capacity to bind. | Reduce the amount of sample added to each well. If you need to purify more his-tagged protein, consider using the Capturem His-Tagged Purification 24-Well Plate, the Capturem His-Tagged Purification Maxiprep Kit or Capturem His-Tagged Purification Large Volume, which have higher binding capacities.   |
| Low yield of his-tagged protein   | Lysis Buffer contains imidazole, which interferes with his-tag binding.  | Make sure that Lysis Buffer is free of imidazole. Our xTractor Buffer does not contain imidazole.   |
|                                   | Too much imidazole in Wash Buffer can elute his-tagged protein.  | Make sure the imidazole concentration in Wash Buffer is no higher than 40 mM.   |
| His-tagged protein does not elute | Elution conditions are too mild, or elution buffer does not contain enough imidazole.  | Follow the instructions using the recommended elution buffer containing appropriate amount of imidazole.  |
| Wells do not fully<br>drain       | Clogging due to particles or a very viscous sample   | <ul> <li>Prepare the lysate according to the xTractor Buffer and xTractor Buffer Kit User Manual.</li> <li>If the lysate is not clear, centrifuge it a second time at 10,000–12,000g for 20 min or use a 0.45-micron filter (cellulose acetate) for further clarification.</li> <li>Consider adding more DNase I to your lysate or lysozyme if appropriate (see xTractor Buffer and xTractor Buffer Kit User Manual).</li> <li>Repeat Capturem His-Tagged Purification 96-Well Plate centrifugation at 2,000g for 3 min. If necessary, repeat this centrifugation one more time.</li> </ul> |

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