# Purification of His<sup>6</sup>-tagged proteins produced in the BL21(DE3) *E. coli* strain. (also check purification of GST-fusion proteins)

<u>Production of His<sup>6</sup>-tagged proteins</u>

- 1. BL21(DE3) cells transformed with a pET28a derivative, use kanamycin (20 ug/ml).
  - \*No heat shock when transformed.
  - \*Do not leave colonies at 4°C. Use fresh colonies.
- 2. Inoculate a single colony in 25 ml LB+kan, culture @25°C O/N
- 3. Dilute in 750 ml (s ml) LB+kan
- 4. Cultivate at 30°C until an OD660 of approx. 0.8 (you can start induction at lower OD, at least 0.4)
  - \*Take 1ml culture for an SDS-PAGE sample (**pre-induction sample**). (Screw top tube)
  - \*Spin down cells. Remove supernatant.
  - \*Add 1xSDS-PAGE loading buffer (OD x 300 ul) to the pellet.
  - \*Suspend cells by vortex and boil for 2min
  - \*Vortex again briefly and boil another 3min
  - \*Chill on ice, freeze at -20°C
- 5. Add IPTG to the culture (final 0.67 mM IPTG)  $(0.1 \sim 1.0 \text{ mM})$  depending on your protein)
- 6. Cultivate @30°C for 2h (2 h ~5 h depending on you protein)
  - \*Measure OD, and prepare SDS-PAGE sample as described below

(post-induction sample)

- \*Take 1ml culture for SDS-PAGE sample. (Screw top tube)
- \*Spin down cells. Remove supernatant.
- \*Add 1xSDS-PAGE loading buffer (OD x 300 ul = t) to the pellet.
- \*Suspend cells by vortex and boil for 2min
- \*Vortex again briefly and boil another 3min
- \*Chill on ice, freeze at -20°C
- 7. Collect cells in 500 ml centrifuge tube (use two bottle, 375 ml each)

 $(5,000 \text{ rpm}, 15 \text{ min}, 4^{\circ}\text{C})$ 

- 8. Suspend cells in 25ml of TBS, transfer the content to 50 ml tube
- 9. Collect cells (3,200 rpm, 15 min, 4°C)
- 12. Store pellet @-80°C

## Purification of His<sup>6</sup>-tagged protein

## All the procedures must be performed at 4°C or on ice unless otherwise specified

- 1. Add 10 ml (w ml) of ice-cold lysis buffer to cell pellet (4°C)
  - Add 10 mg of lysozyme, suspend cells, incubate on ice for 30 min.
- 2. Sonicate cell (4°C)

With Branson Digital Sonifier: amplitude 20%, processing time 30 sec, ON time 1 sec, OFF time 3 sec. Need to repeat  $3 \sim 5$  times.

Keep the tube on ice while sonicating.

- \*Cell suspension changes its color from white to brown
- \*Avoid foaming. Put down the sonicator tip well below the buffer surface during sonication.
- 3. Transfer cell lysase to a 30ml centrifuge tube (4°C)
- 4. Clarify lysate by centrifugation (JA-17, 15krpm, 30min, 4°C) (use adaptor if necessary)
- 5. During centrifugation, pre-wash Ni-NTA beads
  - a. Take 1 ml (50% slurry, 500 ul bed volume/sample) of beads in 15 ml tube
  - b. Spin down to collect beads, 1000 rpm, 1sec, 4°C (remove supernatant)
  - c. Add 5 ml of lysis buffer, mix gently/thoroughly

d. repeat steps: b, c, b, c and b

- 6. After centrifugation, save (w x 1/s) ml of supernatant for SDS-PAGE sample (sup sample, 4°C)
- 7. Transfer supernatant to the 15ml conical tube containing pre-washed Ni-NTA beads (50% slurry in lysis buffer) (4°C). Do not vortex. Do not pipette up and down. Mix by inversion.
- 8. Rotate for 1h at 4°C
- 9. During the step 8:
  - \*Suspend pellet in 10 ml (w ml) of lysis buffer. Save (w x 1/s = u; 14 ul if the original culture is 750 ml) ml for SDS-PAGE sample (pellet sample, 4°C)
  - \*Prepare sup & pellet samples: sample must contains same concentration of proteins as pre- and post-induction sample
    - \*Add (**u x 0.5**) **ml** of 3XSDS-PAGE loading buffer.
    - \*Add (t-1.5u) ul of 1XSDS-PAGE loading buffer (total t ul)
    - \*Suspend by vortex, and boil for 5min
    - \*on ice, then -20°C.
- 10. Load the lysate-Ni-NTA mixture into a column with the bottom outlet capped (in the cold room). Remove the bottom cap and collect flow-through.
  - \*Save flow-through for SDS-PAGE analysis
  - \*Add (u x 0.5) ml of 3XSDS-PAGE loading buffer.
  - \*Add (t-1.5u) ul of 1XSDS-PAGE loading buffer (total t ul)
  - \*Suspend by vortex, and boil for 5min
  - \*on ice, then -20°C
- 11. Wash Ni-NTA with 5 ml of wash buffer; collect fractions for SDS-PAGE analysis. Do twice, but pool two washes in the same tube.
  - \*Save the wash for SDS-PAGE analysis
  - \*Add (u x 0.5) ml of 3XSDS-PAGE loading buffer.
  - \*Add (t-1.5u) ul of 1XSDS-PAGE loading buffer (total t ul)
  - \*Suspend by vortex, and boil for 5min
  - \*on ice, then -20°C
- 12. Elute the protein 4 times with 500 ul of elution buffer
  - \*Prepare SDS-PAGE sample with each fraction (5 fractions)
  - \*30 ul fraction + 15 ul 1x SDS-PAGE loading buffer
  - \*Suspend by vortex, and boil for 5min
  - \*on ice, then -20°C
  - \*Store the rest of protein solution at 4°C (O/N) or -80°C (longer storage) (make 100 ul aliquots from each fraction)

#### **SDS-PAGE** analysis of purified proteins

Load 15 ul from each step/fraction

## Reagents

Lysis buffer (prepare the amount you need)

	Stock soln		
50 mM NaH <sub>2</sub> PO <sub>4</sub>	1M		
300mM NaCl	5M		
10% Glycerol	60%		
0.25% Tween 20	20%		
10 mM Imidazole	1M		
10 mM beta-Mercaptoethanol**	14M		
1 mM PMSF**	1M		
Total			

\*\*add before use ##Adjust pH to 8.0 using NaOH

Wash buffer (prepare the amount you need)

wash sailer (stepare the amount you need)					
	Stock soln				
50 mM NaH <sub>2</sub> PO <sub>4</sub>	1M				
300mM NaCl	5M				
10% Glycerol	60%				
0.25% Tween 20	20%				
20 mM Imidazole	1M				
10 mM beta-Mercaptoethanol**	14M				
1 mM PMSF**	1M				
Total					

<sup>\*\*</sup>add before use

##Adjust pH to 8.0 using NaOH

Elution buffer (prepare the amount you need)

	Stock soln		
50 mM NaH <sub>2</sub> PO <sub>4</sub>	1M		
300mM NaCl	5M		
10% Glycerol	60%		
0.25% Tween 20	20%		
250 mM Imidazole	1M		
10 mM beta-Mercaptoethanol**	14M		
1 mM PMSF**	1M		
Total			

<sup>\*\*</sup>add before use

##Adjust pH to 8.0 using NaOH

Lysozyme (powder)