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Datasheet for ABIN4368209 INDIGO Ni-Agarose

3 Images



## Overview

Characteristics:

Quantity:	10 mL
Application:	Purification (Purif)
Product Details	
Purpose:	Specific binding and purification of his-tagged proteins
Brand:	HighSpec
Sample Type:	Cell Culture Supernatant, Cell Lysate, Tissue Lysate

Novel His affinity ligand providing stability in 20 mM EDTA and 20 mM DTT

- Binding capacity up to 80 mg/mL
- Suitable for high flow rates in batch and FPLC
- pH tolerant from pH 4-9.
- Delivered as 20 mL of a 50 % suspension
- Average agarose bead size: 100  $\mu m$

Components:	Affinity Agarose
Material not included:	Lysis Buffer
	Wash Buffer
	Elution Buffer
	Ice bath
	Refrigerated centrifuge for 50 mL tube (min 10,000 x g)
	• 50 mL centrifuge tube
	Micropipettor and Micropipetting tips
	Disposable gravity flow columns with capped bottom outlet 2 ml

• pH meter

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### • End-over-end shaker

SDS-PAGE buffers, reagents and equipment

Optional: Western Blot reagents and equipment

Bead Size:	100 um
Bead Matrix:	Agarose beads
Bead Ligand:	Ni-INDIGO

# Application Details

Comment:	Sample Volume for an assay: >200 mL E.coli culture volume or corresponding quantity.
	Protocol can be scaled up easily.
Sample Volume:	200 mL
Assay Time:	4 - 5 h
Protocol:	Purification of his-tagged protein in batch gravity flow or on FPLC columns
Reagent Preparation:	A Purification under native conditions:
	<ul> <li>Native Lysis buffer: NaH2PO4 50 mM, NaCl 300 mM, Imidazole 10 mM, pH 8 Optional: Add 1 tablet protease inhibitor cocktail to the Lysis Buffer. Tip: Lysis Buffer contains 10 mM imidazole to prevent binding of untagged proteins. If His- tagged proteins do not bind under these conditions, reduce the imidazole concentration to 1- 5 mM.</li> <li>Native Wash buffer: NaH2PO4 50 mM, NaCl 300 mM, Imidazole 20 mM, pH 8</li> <li>Native Elution buffer: NaH2PO4 50 mM, NaCl 300 mM, Imidazole 250- 500 mM, pH 8 Additional chemicals required: Lysozyme, Benzonase® nuclease, Optional: Protease inhibitor cocktail</li> </ul>
	B Purification under denaturing conditions:
	<ul> <li>Denaturing Lysis buffer: NaH2PO4 100 mM, Tris base 10 mM, Urea 8 M, pH 8.0, Optional: Benzonase® nuclease (e.g. Merck Milipore, #707464)</li> <li>Denaturing Wash buffer: NaH2PO4 100 mM, Tris base 10 mM, Urea 8 M, pH 6.3NaH2PO4 100 mM</li> <li>Denaturing Elution buffer: NaH2PO4 100 mM, Tris base 10 mM, Urea 8 M, pH 4.5 Tip: If the target protein is acid-labile, elution can be performed with 250-500 mM imidazole. Note: Due to urea dissociation, adjust the pH immediately before use.</li> </ul>
Assay Procedure:	A Protocol for purification under native conditions:

- 1. Thaw the E. coli cell pellets corresponding to 200 mL bacterial culture on ice for 15 min. Optional: Freezing the cell pellet at -20 °C for 30 min prior to incubation at room temperature improves lysis by lysozyme.
- 2. Resuspend the cell pellet in 10 mL Native Lysis Buffer supplemented with 1 mg/mL lysozyme, and pour it into a 50 mL conical centrifuge tube.
- 3. If the solution is very viscous, add 3 units Benzonase® per mL E.coli culture volume to the lysis buffer. Alternatively or additionally, sonicate the lysate to improve cell disruption.
- 4. Incubate on an end-over-end shaker at room temperature for 30 min, or at 4 °C for 1 h, depending on the temperature stability of the protein.
- 5. Centrifuge the lysate for 30 min at 10,000 x g and 2-8 °C. Carefully collect the supernatant without touching the pellet.

Note: The supernatant contains the cleared lysate fraction. We recommend to take aliquots of all fractions for SDS-PAGE analysis.

6. Resuspend the HighSpec Ni-NTA Agarose by inverting the bottle until the suspension is homogeneous. Transfer 1 mL of the 50 % suspension (corresponding to 500 μL bed volume) to a 15 mL conical centrifuge tube. Allow the resin to settle by gravity and remove the supernatant.

Tip: Alternatively, resin equilibration can be performed directly in the disposable gravity flow column.

- 7. Add 2.5 mL Native Lysis Buffer and gently resuspend the slurry to equilibrate the resin. Allow the resin to settle by gravity and remove 2 mL supernatant.
- 8. Add 10 mL cleared lysate to the equilibrated HighSpec Ni-NTA Agarose resin and incubate at 4 °C for 1 h on an end-over-end shaker.

Tip: Alternatively, batch binding can be performed directly in a gravity flow column with closed bottom and top outlets.

- 9. Transfer the binding suspension to a disposable gravity flow column with a capped bottom outlet. Use Lysis Buffer to rinse the centrifuge tube and remove resin adhered to the wall.
- 10. Remove the bottom cap of the column and collect the flow-through.
- 11. Wash the column with 5 mL Native Wash Buffer. Repeat the washing step at least 3 times.
- 12. Elute the His-tagged protein 5 times using 0.5 mL Native Elution Buffer. Collect each eluate in a separate tube and determine the protein concentration of each fraction. Optional: Incubate the resin for 15 min in Elution Buffer before collecting the eluate to increase protein yields.
- Analyze all fractions by SDS-PAGE.
   Note: Do not boil membrane proteins. Instead, incubate samples at 46 °C for 30 min in preparation for SDS-PAGE analysis.
- 14. Optional: Perform Western Blot experiment using PentaHis Antibody.

## B Protocol for purification under denaturing conditions:

- 1. Thaw the E. coli cell pellet on ice.
- Resuspend the cell pellet in 10 mL Denaturing Lysis Buffer. Optional: Benzonase® can be added to the lysate to reduce viscosity caused by nucleic acids (3 U/mL bacterial culture). Nucleic acids can also be sheared by passing the lysate 10 times through a fine-gauge needle.
- 3. Incubate at room temperature for 30 min on an end-over-end shaker.

	<ul> <li>4. Centrifuge the lysate for 30 min at room temperature and 10,000 x g. Collect the supernatant. Note: The supernatant contains the cleared lysate fraction. We recommend to take aliquots of all fractions for SDS-PAGE analysis.</li> <li>5. Resuspend the HighSpec Ni-NTA Agarose by inverting the bottle until the suspension is homogeneous. Transfer 1 mL of the 50 % suspension (corresponding to 0.5 mL bed volume) into a 15 mL conical centrifuge tube. Allow the resin to settle by gravity and remove the supernatant.</li> <li>6. Add the cleared lysate to the resin and incubate the mixture for 1 h at room temperature on an end-over-end shaker. Tip: Alternatively, batch binding can be done directly in a gravity flow column with closed top and bottom outlet.</li> <li>7. Transfer the binding suspension to a disposable gravity flow column with a capped bottom outlet. Use Lysis Buffer to rinse the centrifuge tube and remove resin adhered to the wall.</li> <li>8. Remove the bottom cap of the column and collect the flow-through.</li> </ul>
	9. Wash the column with 5 mL Denaturing Wash Buffer. Repeat the washing step at least 3 times.
	10. Elute the His-tagged protein 5 times using 0.5 mL Denaturing Elution Buffer. Collect each eluate in a separate tube and determine the protein concentration of each fraction. Tip: If the target protein is acid-labile, elution can be performed with 250-500 mM imidazole.
	<ol> <li>Analyze all fractions by SDS-PAGE.</li> <li>Note: Do not boil membrane proteins. Instead, incubate samples at 46 °C for 30 min in preparation for SDS-PAGE analysis.</li> <li>Optional: Perform Western Blot experiment using PentaHis Antibody.</li> </ol>
Calculation of Results	Analyze by SDS-PAGE Bradford Assay or spectrophotometrically
Postrictions:	Ear Desearch Lise only
	TO Research use only
Handling	
Format:	Liquid
Storage:	4 °C



Elution Run 1

Elution Run 4

Elution Run 8

Image 1. Fig.2: PureCube 100 INDIGO Ni-Agarose outperforms competitor products. His-tagged GFP was purified on PureCube 100 INDIGO Ni-Agarose and two leading competitor matrices. Yields obtained with the INDIGO matrix were considerably higher at comparable purity. Buffer conditions: Sodium phosphate buffer pH 7.4, 10 mM DTT, 20 mM EDTA. Imidazole concentrations: Binding step: 10 mM, Wash: 20 mM, Elution: 250 mM. Fig.3: PureCube 100 INDIGO Ni-Agarose can be re-used multiple times without regeneration. GFP was spiked into E.coli lysates and purified in eight aliquots on the same 1 ml column filled with PureCube 100 INDIGO Ni-Agarose. Between each run, the column was briefly washed with loading buffer containing PBS and 10 mM imidazole. No decrease in performance was observed, even after eight consecutive runs. Left: Chromatogram; Right: SDS-PAGE. M: Marker, L: Lysate, S: Lysate spiked with GFP.

### SDS-PAGE

**Image 2.** Fig.3: PureCube 100 INDIGO Ni-Agarose can be reused multiple times without regeneration. GFP was spiked into E.coli lysates and purified in eight aliquots on the same 1 ml column filled with PureCube 100 INDIGO Ni-Agarose. Between each run, the column was briefly washed with loading buffer containing PBS and 10 mM imidazole. No decrease in performance was observed, even after eight consecutive runs. Left: Chromatogram; Right: SDS-PAGE. M: Marker, L: Lysate, S: Lysate spiked with GFP.

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#### SDS-PAGE

**Image 3.** Fig.1: PureCube 100 INDIGO Ni-Agarose is compatible with 20 mM EDTA and 20 mM DTT. SDS-PAGE of GFP expressed in E.coli and purified with PureCube 100 INDIGO Ni-Agarose in the presence of 20 mM EDTA and 20 mM DTT. High yield (>80 mg/ml) and purity were obtained.

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