EconoFit Nuvia IMAC Ni-Charged Columns, 1 and 5 ml

Instruction Manual

Catalog numbers

12009285

12009286

12009287

12009288

Please read the instructions in this manual prior to using EconoFit Nuvia IMAC Ni-Charged Columns. If you have any questions or require any further assistance, please contact your Bio-Rad Laboratories representative.



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Introduction

EconoFit Nuvia IMAC Ni-Charged Columns are convenient, disposable, prepacked low-pressure chromatography columns. EconoFit Columns offer both increased run-to-run uniformity and high purity of protein through the column design and novel resin technology. Compatible with aqueous buffers most commonly used for protein purification, EconoFit Columns offer improved performance for your protein separation needs.

Immobilized metal affinity chromatography (IMAC) is an excellent chromatography technique for purification of histidine-tagged proteins. The principle of IMAC is based on the affinity histidine has for metal ions. Side chains on the nitrilotriacetic acid (NTA) functional ligand selectively bind recombinant histidine-tagged proteins when the resin is charged with Ni²⁺ or other metals. The advantage of this technique is that proteins can often be purified close to homogeneity in a single step.

These columns are packed with Bio-Rad's specially designed Nuvia IMAC Ni-Charged Resin. Such characteristics as the polymeric nature, optimized ligand density, and open pore structure of the Nuvia IMAC Bead result in superb mechanical strength with high stringency, low nonspecific binding, and the ability to perform separations at high flow rates.

Section 2

Product Information

EconoFit Columns are disposable, easy-to-use, prepacked chromatographic columns supplied ready for use in convenient 1 and 5 ml sizes. They are quickly connected to liquid chromatography systems using 10-32 fittings. Columns are available for a variety of chromatographic techniques, including desalting (size exclusion [SEC]), ion exchange (IEX), affinity (AC), mixed-mode, and hydrophobic interaction chromatography (HIC). See Table 1 for specifications. Refer to bio-rad.com/ResinsandColumns for a complete listing of items in the EconoFit Column product line.

Table 1, EconoFit Nuvia IMAC Ni-Charged Column specifications

Property	Description
Size	1 and 5 ml bed volumes
Dimensions	1 ml: 80 mm length x 15 mm inner diameter 5 ml: 80 mm length x 23 mm inner diameter
Recommended flow rate	1 ml: 1–2 ml/min (240–480 cm/hr) 5 ml: 5–10 ml/min (240–480 cm/hr)
Maximum flow rate	1 ml: 6 ml/min (1,440 cm/hr) 5 ml: 20 ml/min (963 cm/hr)
Fittings	10-32 (1/16"), female inlet and male outlet
Column material	Polypropylene
Frit material	High-density polyethylene
Shipping solution	20% ethanol
Storage conditions	20% ethanol
Autoclavability	Not autoclavable

Nuvia IMAC Ni-Charged Resin is also available in bottles. Refer to Ordering Information in section 10 of this manual. See Table 2 for specifications. Go to bio-rad.com/ResinsandColumns for more information.

Table 2. Nuvia IMAC Ni-Charged Resin specifications.

Property	Description
Functional ligand	Nitrilotriacetic acid
Base bead	UNOsphere
Particle size	38–53 μm
Mean particle size	50 μm
Metal ion capacity	≥18 µmol Cu²+/ml
Dynamic binding capacity*	≥40 mg/ml
Recommended flow rate	480 cm/hr
Maximum operating pressure	45 psi
Chemical compatibility	See Table 3
Storage temperature	4°C to ambient temperature
Storage conditions	20% ethanol
Shelf life in 20% ethanol	>3 years at ambient temperature
Operational temperature	4-40°C
Autoclavability	Not autoclavable

^{*} Q10% determination of 1 mg/ml histidine-tagged pure protein (40 kD). Note: Dynamic binding capacity will vary from protein to protein.

Nuvia IMAC Columns are compatible with the aqueous buffers most commonly used with IMAC purification techniques (see Table 3).

Table 3. Chemical compatibilities for Nuvia IMAC Ni-Charged Columns.*

Reagent	Stability
Chelating Agent	
EDTA	1 mM
Sulfhydryl Reagents	
β-mercaptoethanol	20 mM
DTT	10 mM
TCEP	20 mM
Detergents	
Nonionic	2% (Triton, Tween)
Zwitterionic	1% (CHAPS, CHAPSO)
Denaturing Agents	
Guanidine HCI	6 M
Urea	8 M
Other Additives	
NaCl	2 M (include at least 300 mM NaCl in buffers)
MgCl ₂	100 mM (use HEPES to prevent precipitation)
CaCl ₂	5 mM (use HEPES to prevent precipitation)
Glycerol	20% (backpressure may increase significantly, slower flow rates may be required)
Ethanol	20%

^{*} Nuvia IMAC binding capacities are unaffected up to the concentrations given when employing typical reagents used for histidine-tagged protein purification.

Buffers and Methods

IMAC methods can be run using either native or denaturing purification protocols. Under native conditions, proteins are purified using buffers that help retain the natural folded structure of the target protein. Under denaturing conditions, strong chaotropic agents (typically 6-8 M urea or 6 M guanidine) are added to the buffers, allowing target proteins to be purified in their unfolded states. The recommended buffer compositions and formulations are provided in Table 4.

Table 4. Suggested buffer composition.

Step	Buffer*			
	NaCl, mM	Na phosphate, mM	Imidazole, mM	Urea, M
Native lysis/wash buffer 1	300	50	5	NA
Native wash buffer 2	300	50	25	NA
Native elution buffer	300	50	500	NA
Denaturing lysis/wash buffer 1	300	50	5	6
Denaturing wash buffer 2	300	50	25	6
Denaturing elution buffer	300	50	500	6

^{*} For all buffer formulations, adjust pH to 8.0 with KOH or H₉PO₄ and filter through a 0.2 µm filter. Native buffers can be stored up to 1 year at 4-22°C. Denaturing buffers must be made fresh and used within 7 days or frozen in aliquots at -20°C for later use.

Section 4

Quick Solubility Screening Protocols

Before choosing a native or denaturing purification protocol, it is useful to determine both the approximate expression level of a protein and whether the overexpressed target protein partitions into the soluble or insoluble fraction. Soluble proteins are typically purified with the native purification procedure while insoluble proteins must be solubilized in stringent denaturants (urea or guanidine) and are purified with the denaturing procedure.

The following procedure provides a quick screen for solubility and expression level:

- Pellet ~2 ml of Escherichia coli culture by centrifugation at 4,000 x g for 10 min at 4°C.
- 2. Resuspend the pellet in 500 µl of phosphate buffered saline (PBS) and sonicate on ice for 60 sec, in 10 sec pulses. Remove 50 µl of sonicate and label the sample Total. Centrifuge the lysate at 12,000 x g for 10 min at 4°C. Transfer the supernatant to a clean tube. Remove 50 µl of the supernatant and label the tube Soluble.
- 3. Resuspend the insoluble pellet in 500 µl of 6 M urea in 1x PBS and sonicate on ice for 60 sec, in 10 sec pulses. Centrifuge the lysate at 12,000 x g for 10 min at 4°C. Remove 50 µl of the supernatant and label the tube Insoluble.
- 4. To each of the 50 μl samples, add 150 μl of Laemmli buffer and boil for 5 min at 95°C.
- 5. Load 10 µl of each sample on an SDS-PAGE gel.
- 6. Examine the soluble and insoluble fractions for the target protein. Approximate the expression level and determine partitioning of the target protein.

Preparation of E. coli Lysates

For E. coli cultures expressing medium to high levels of histidine-tagged proteins (≥10% of total protein). 200 ml of culture will yield sufficient material for a 1 ml column purification, and 1,000 ml of culture will yield sufficient material for a 5 ml column purification run. For cultures expressing protein at low levels (~10% of total protein), the culture volumes will need to be determined empirically for each protein.

Native Lysates

- 1. Harvest the cell pellet by centrifugation at 8,000 x g for 10 min at 4°C.
- 2. Determine the weight of the pellet and resuspend in 10 culture volumes of native lysis/wash buffer 1 (200 ml of culture typically yields 0.8 g of paste and results in 8 ml of lysate).
- 3. Sonicate the lysate on ice four times at 1 min intervals.
- 4. Centrifuge the lysate at 12,000 x g for 20 min at 4°C.
- 5. Remove the supernatant and filter it through a 0.2 µm filter immediately before applying to the column.

Denatured Lysates

- 1. Harvest the cell pellet by centrifugation at 8,000 x g for 10 min at 4°C.
- 2. Determine the weight of the pellet and resuspend in 10 culture volumes of denaturing lysis/wash buffer 1 (200 ml of culture typically yields 0.8 g of paste and results in 8 ml of lysate).
- 3. Sonicate the lysate four times at 1 min intervals.
- 4. Centrifuge the lysate at 12,000 x g for 20 min at 4°C.
- 5. Remove the supernatant and filter it through a 0.2 µm filter immediately before applying to the column.

Section 6

Preparing a Column and Subsequent Purification

Prepare buffer sets for either the native or denaturing purification protocols using a single buffer set throughout the procedure. To prepare the column for the procedure, remove the top closure and connect the column to the chromatography system. Open the bottom closure and connect the column outlet to the system. Flush the packing solution (20% ethanol) from the column by running 2 column volumes (CV) of water at a flow rate of 2 ml/min (1 ml column) or 10 ml/min (5 ml column). The column is now ready for the purification steps. Flow rates are given in ml/min and are specific to the 1 ml column.

If using a 5 ml column for a procedure, substitute the higher flow rate in the method (see Table 5).

Table 5. Purification method suggestions.

Step	Column Volumes, CV	1 ml Column Flow Rate, ml/min	5 ml Column Flow Rate, ml/min
Equilibrate	5	2	10
Lysate load	Varies based on sample volume	1*	5*
Wash 1	6	1	5*
Wash 2	6	2	10
Elute	5	2	10

^{*} Depending on sample viscosity.

Standard methods that are compatible with any type of chromatography system are listed in the following steps. To maximize binding capacity with large proteins (>100 kD), for purification at 4°C, or for purifications under denaturing conditions, the lysate load flow rate can be decreased (to 0.5 ml/min for the 1 ml column and 2 ml/min for the 5 ml). Whether this decrease maximizes flow rate will have to be determined empirically for individual proteins.

- Equilibrate the column with 5 CV of equilibration/wash buffer 1 at 2 ml/min.
- Load the sample lysate at 1 ml/min.
- Wash the column with 6 CV of wash buffer 1 at 1 ml/min.
- 4. Wash the column with 6 CV of wash buffer 2 at 2 ml/min.

Note: Equivalent to 5% buffer B/elution buffer wash.

- 5. Elute the purified protein with 10 CV of elution buffer at 2 ml/min.
- 6. Prior to quantitation of the protein concentration, the purified protein should be exchanged into a nonimidazole buffer (imidazole can absorb at 280 nm). Purified protein from denaturing purifications should be exchanged into another buffer through dialysis.

The chromatogram in Figure 1 illustrates a representative purification of a high-expressing soluble protein purified using the native buffer set and method described in Table 5.

Note: IMAC buffers made with potassium salts are more stable than sodium salt-based buffers. However, potassium will complex with sodium dodecyl sulfate (SDS) in Laemmli buffer and precipitate out of solution. Prior to analyzing IMAC samples on gels, the samples must be diluted at least 1:7 with Laemmli buffer to prevent precipitation.

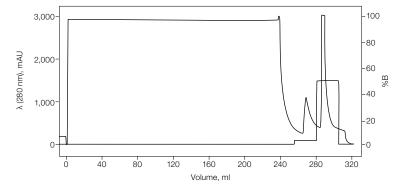


Fig. 1. Typical IMAC purification. A histidine-tagged protein was purified from the soluble fraction using the standard Nuvia IMAC native purification protocol. Clarified E. coli lysate was loaded onto a 5 ml Nuvia IMAC Ni-Charged Column. The column was washed with 6 CV of wash buffer 1 followed by 6 CV of wash buffer 2. Purified protein was eluted with 5 CV of elution buffer.

Scaling Up

EconoFit Columns are available in 1 and 5 ml formats. The Nuvia IMAC Resin is also available in various amounts, from 25 ml bottles to larger bulk quantities, for scaling up methods developed using the columns. For quick scale-up, two or three columns of the same type can be connected in series, so take care to maintain an overall system pressure ≤45 psi.

In addition, Bio-Rad carries an extensive line of empty chromatography columns from laboratory to process scale. Ask your local Bio-Rad representative or go to bio-rad.com/ResinsandColumns for more information.

Section 8

Regenerating, Cleaning, Sanitizing, and Storing Columns

Protein cross-contamination, frit clogging, and increased backpressure can result from running a column beyond the recommended number of uses. After repeated use, a column may run slower or produce high backpressure, a result that should be expected due to the nature of the sample mixture. The following cleaning and regeneration procedures may be used. However, we recommend that you dispose of a column after several uses. To avoid cross-contamination, we recommend that each column be designated for a single protein. To maintain good flow properties, we recommend that the columns be cleaned between uses.

Run the cleaning protocol at 2 ml/min for 1 ml columns. For 5 ml columns, run the cleaning protocol at 5 ml/min.

High Salt/Acid Cleaning

- 1. Rinse the column with 2 CV of water at 2 ml/min.
- 2. Wash the column with 5 CV of 500 mM NaCl, 50 mM Tris, pH 8.0 at 2 ml/min.
- 3. Wash the column with 5 CV of 500 mM NaCl, 100 mM NaOAc, pH 4.5 at 2 ml/min.
- 4. Rinse the column with 2 CV of water at 2 ml/min.
- 5. Store the column in 20% ethanol at 4°C.

Chaotropic Cleaning

- 1. Rinse the column with 2 CV of water at 2 ml/min.
- 2. Wash the column with 5 CV of 6 M guanidine HCl at 2 ml/min.
- 3. Wash the column with 2 CV of water at 2 ml/min.
- 4. Rinse the column with 2 CV of water at 2 ml/min.
- 5. Store the column in 20% ethanol at 4°C.

Other Cleaning Guidelines

In situations where it is desirable to run different proteins over the same column, completely sanitize, strip, and recharge the column between sample runs. Take care when handling and disposing of metal-containing solutions.

- 1. Clean the column with 10 CV of 1 M NaOH.
- 2. Rinse the column with 10 CV of water.
- 3. Strip metal ions with 5 CV of 0.1 M EDTA.
- 4. Rinse the column with 10 CV of water.
- 5. Recharge the column with 5 CV of 0.1 M nickel sulfate, pH 4.5.
- 6. Rinse the column with 10 CV of water.
- 7. Store the column in 20% ethanol.

Troubleshooting Guide

Column Clogging or Slow Flow Rate Particulates in sample Filter all samples and buffers through 0.2 µm filter prior to application Sample too viscous Add nuclease to lysate to degrade DNA. Centrifuge and filter lysate again No Target Protein in Eluate Low level of target Check expression level of protein in starting SDS-PAGE material Target protein not binding • Check levels of target protein in lysate, flowthrough, wash fractions, and eluted fractions • Check for presence of histidine tag with antihistidine antibody Target Protein in Flowthrough Histidine tag not accessible • Purify protein under denaturing conditions to expose histidine tag • Reclone histidine tag not opposite terminus (N- or C- terminus) Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting • Elute with imidazole gradient (10–500 mM) rather than step elution • Elute with imidazole gradient (10–500 mM) rather than step elution • Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target • Add protease inhibitors to protein lysate • Purify at 4°C or under denaturing conditions	Possible Causes	Possible Solutions
Add nuclease to lysate to degrade DNA. Centrifuge and filter lysate again No Target Protein in Eluate Low level of target Check expression level of protein in starting SDS-PAGE material **Check levels of target protein in lysate, flowthrough, wash fractions, and eluted fractions **and eluted fractions** Check for presence of histidine tag with antihistidine antibody **Target Protein in Flowthrough** Histidine tag not accessible **Purify protein under denaturing conditions to expose histidine tag **Reclone histidine tag onto opposite terminus (N- or C- terminus) Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold **Precipitation during Purification Binding capacity of column exceeded Protein aggregating **Include low amount of detergent (0.1% Triton X-100, Tween 20) **Include glycerol up to 10% Elute with imidazole gradient **Elute With imidazole gradient (10–500 mM) rather than step elution **Increase imidazole in the wash to increase wash stringency, but keep below 40 mM **Target Protein Is Degraded Proteolysis of target **Add protease inhibitors to protein lysate		
No Target Protein in Eluate Low level of target Check expression level of protein in starting SDS-PAGE material • Check levels of target protein in lysate, flowthrough, wash fractions, and eluted fractions • Check for presence of histidine tag with antihistidine antibody Target Protein in Flowthrough Histidine tag not accessible • Purify protein under denaturing conditions to expose histidine tag • Reclone histidine tag onto opposite terminus (N- or C- terminus) Proteolysis and removal Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating • Include low amount of detergent (0.1% Triton X-100, Tween 20) • Include glycerol up to 10% Fluted Protein Is Impure Contaminants coeluting • Elute with imidazole gradient (10–500 mM) rather than step elution • Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target • Add protease inhibitors to protein lysate	Particulates in sample	Filter all samples and buffers through 0.2 µm filter prior to application
Check expression level of protein in starting SDS-PAGE material Target protein not binding Check levels of target protein in lysate, flowthrough, wash fractions, and eluted fractions Check for presence of histidine tag with antihistidine antibody Target Protein in Flowthrough Histidine tag not accessible Purify protein under denaturing conditions to expose histidine tag Reclone histidine tag onto opposite terminus (N- or C- terminus) Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating Include low amount of detergent (0.1% Triton X-100, Tween 20) Include glycerol up to 10% Elute with imidazole gradient Eluted Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate	Sample too viscous	Add nuclease to lysate to degrade DNA. Centrifuge and filter lysate again
Target protein not binding ■ Check levels of target protein in lysate, flowthrough, wash fractions, and eluted fractions ■ Check for presence of histidine tag with antihistidine antibody Target Protein in Flowthrough Histidine tag not accessible ■ Purify protein under denaturing conditions to expose histidine tag ■ Reclone histidine tag onto opposite terminus (N- or C- terminus) Proteolysis and removal Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating ■ Include low amount of detergent (0.1% Triton X-100, Tween 20) ■ Include glycerol up to 10% Frotein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting ■ Elute with imidazole gradient (10–500 mM) rather than step elution ■ Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target ■ Add protease inhibitors to protein lysate	No Target Protein in Eluate	
and eluted fractions Check for presence of histidine tag with antihistidine antibody Target Protein in Flowthrough Histidine tag not accessible Purify protein under denaturing conditions to expose histidine tag Reclone histidine tag onto opposite terminus (N- or C- terminus) Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating Include low amount of detergent (0.1% Triton X-100, Tween 20) Include glycerol up to 10% Protein too concentrated in step elution Elute Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate	Low level of target	Check expression level of protein in starting SDS-PAGE material
Target Protein in Flowthrough Histidine tag not accessible Proteolysis and removal Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating Include low amount of detergent (0.1% Triton X-100, Tween 20) Include glycerol up to 10% Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate	Target protein not binding	
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 ■ Reclone histidine tag onto opposite terminus (N- or C- terminus) Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating	Target Protein in Flowthrough	
Proteolysis and removal Include protease inhibitors in histidine-tagged lysis buffer or purify in the cold Precipitation during Purification Binding capacity of column exceeded Protein aggregating Include low amount of detergent (0.1% Triton X-100, Tween 20) Include glycerol up to 10% Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate	Histidine tag not accessible	 Purify protein under denaturing conditions to expose histidine tag
Precipitation during Purification Binding capacity of column exceeded Protein aggregating Include low amount of detergent (0.1% Triton X-100, Tween 20) Include glycerol up to 10% Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate		 Reclone histidine tag onto opposite terminus (N- or C- terminus)
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 ■ Include glycerol up to 10% Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting ■ Elute with imidazole gradient (10–500 mM) rather than step elution ■ Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target ■ Add protease inhibitors to protein lysate 		Load less sample
Protein too concentrated in step elution Eluted Protein Is Impure Contaminants coeluting • Elute with imidazole gradient (10–500 mM) rather than step elution • Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target • Add protease inhibitors to protein lysate	Protein aggregating	■ Include low amount of detergent (0.1% Triton X-100, Tween 20)
Eluted Protein Is Impure Contaminants coeluting Elute with imidazole gradient (10–500 mM) rather than step elution Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate		Include glycerol up to 10%
Contaminants coeluting • Elute with imidazole gradient (10–500 mM) rather than step elution • Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target • Add protease inhibitors to protein lysate		Elute with imidazole gradient
 Increase imidazole in the wash to increase wash stringency, but keep below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate 	Eluted Protein Is Impure	
below 40 mM Target Protein Is Degraded Proteolysis of target Add protease inhibitors to protein lysate	Contaminants coeluting	■ Elute with imidazole gradient (10-500 mM) rather than step elution
Proteolysis of target Add protease inhibitors to protein lysate		· · · · · · · · · · · · · · · · · · ·
	Target Protein Is Degraded	
 Purify at 4°C or under denaturing conditions 	Proteolysis of target	 Add protease inhibitors to protein lysate
		 Purify at 4°C or under denaturing conditions

Ordering Information

Catalog # Description

EconoFit Nuvia IMAC Ni-Charged Columns

EconoFit Nuvia IMAC Ni-Charged Column, 1 x 1 ml column 12009288 12009285 EconoFit Nuvia IMAC Ni-Charged Columns, 5 x 1 ml columns 12009286 EconoFit Nuvia IMAC Ni-Charged Column, 1 x 5 ml column EconoFit Nuvia IMAC Ni-Charged Columns, 5 x 5 ml columns 12009287

Nuvia IMAC Ni-Charged Resins

7800800 Nuvia IMAC Ni-Charged Resin, 25 ml bottle 7800801 Nuvia IMAC Ni-Charged Resin, 100 ml bottle 780-0802 Nuvia IMAC Ni-Charged Resin, 500 ml bottle 780-0802 12003233 Nuvia IMAC Ni-Charged Resin, 5 L bottle 12002782 Nuvia IMAC Ni-Charged Resin, 10 L bottle

Section 11

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