



A Geno Technology, Inc. (USA) brand name

Cobalt Chelating Resin

A Co-IDA IMAC resin for 6X-His Tagged Protein Purification

(Cat. # 786-286, 786-402, 786-403, 786-600)



INTRODUCTION	3
ITEM(S) SUPPLIED	3
STORAGE CONDITIONS	3
SPECIFICATIONS	3
IMPORTANT INFORMATION	4
ADDITONAL ITEMS REQUIRED	4
PREPARATION BEFORE USE	4
PROTOCOL FOR NATIVE PROTEINS	5
PROTOCOL FOR DENATURING PROTEINS	6
COLUMN REGENERATION	7
COLUMN RECHARGING PROTOCOL	7
TROUBLESHOOTING	8
CHEMICAL COMPATIBILITIES	9
RELATED PRODUCTS	11

INTRODUCTION

Immobilized Metal Ion Affinity Chromatography (IMAC), developed by Porath (1975), is based on the interaction of certain protein residues (histidines, cysteines, and to some extent tryptophans) with cations of transition metals. The Cobalt Chelating Resin is specifically designed for the purification of recombinant proteins fused to the 6 x histidine (6XHis) tag.

The Cobalt Chelating Resin is specifically designed for the purification of recombinant proteins fused to the 6 x histidine (6XHis) tag expressed in bacteria, insects, and mammalian cells. The resin is high affinity and selectivity for recombinant fusion proteins that are tagged with six tandem histidine residues. Although 6 X His tagged proteins bind with a slightly lower efficiency compared to Nickel Chelating Resin there is a significant reduction in non-specific binding.

The Cobalt Chelating Resin can be used to purify 6X His tagged proteins under native and denaturing conditions. Proteins bound to the resin can be eluted with low pH buffer or competition with imidazole or histidine.

The Cobalt Chelating Resin uses IDA (iminodiacetic acid) as its functional ligand. The tertiary amine and carboxylic acid side chains of IDA serve as the chelating ligands for dior trivalent metal ions. The structure offers selective binding of recombinant His-tagged proteins when this resin is charged with transition metals. As a result, the desired proteins can often be purified close to homogeneity in a single step.

ITEM(S) SUPPLIED

(-)		
Cat. #	Description	Size
786-286	Cobalt Chelating Resin*	10ml
786-402	Cobalt Chelating Resin*	100ml
786-403	Cobalt Chelating Resin*	500ml
786-600	Cobalt Chelating Resin*	2 x 500ml

^{*}Cobalt Chelating Resin is supplied as a 50% slurry in 20% ethanol

STORAGE CONDITIONS

It is shipped at ambient temperature. Upon arrival, store it refrigerated at 4° C, <u>DO NOT FREEZE</u>. This product is stable for 1 year at 4° C.

SPECIFICATIONS

- Ligand Density: 20-40μmoles Co²⁺/ ml resin
- Binding Capacity: >50mg/ml resin. We have demonstrated binding of >100mg of a
 50kDa 6X His tagged proteins to a ml of resin

Bead Structure: 6% cross-linked agarose

IMPORTANT INFORMATION

- The purity and yield of the recombinant fusion protein is dependent of the
 protein's confirmation, solubility and expression levels. We recommend optimizing
 and performing small scale preparations to estimate expression and solubility
 levels.
- Avoid EDTA containing protease inhibitor cocktails, we recommend our Recom ProteaseArrest[™] (Cat. # 786-376, 786-436) for inhibiting proteases during the purification of recombinant proteins.
- For recombinant proteins that are sequestered to inclusion bodies we recommend out IBS™ Buffer (Cat. # 786-183)

ADDITONAL ITEMS REQUIRED

- Disposable columns
- Binding Buffer and Elution Buffer, see protocol for details.

PREPARATION BEFORE USE

Sample preparation: Refer to manufacturer's protocols for optimal conditions for growth, induction and lysis of recombinant His-tagged clones. To avoid clogging of the resin filter the sample through a 0.45μm filter. The preferred buffers that improve binding affinity are 50mM acetate or 10-150mM phosphate buffers with pH 7-8, although this can fluctuate between pH 5.5-8.5. Avoid buffers with primary amines (Tris, Glycine) as these weaken binding affinity and can even strip metal ions. The buffer should be supplemented with 0.15-0.5M NaCl to suppress secondary ionic interactions and proteins/protein interactions.

PROTOCOL FOR NATIVE PROTEINS

- Add an appropriate amount of Cobalt Chelating Resin to a suitable column (suitable to hold 7 columns volumes (CV)). Allow the storage buffer to flow through the column or centrifuge at 800xg for 1 minute. Discard the flow-through.
 NOTE: If using tubes, as opposed to column, pellet the resin by centrifugation at 500xq for 2-5 minutes and carefully decant the storage buffer.
- 2. Add 5CV of distilled water and resuspend the resin. Remove water as above
- Wash the resin by resuspending in 1CV suitable binding buffer (i.e. 50mM Na₂HPO₄, 300mM NaCl pH8.0) supplemented with 10mM imidazole). Remove wash buffer as above and repeat this wash step once.

NOTE: The imidazole and sodium chloride is present to reduce non-specific interactions, however these may interfere with the binding of some 6X His tagged proteins. Optimize with 0-20mM imidazole and 100-500mM NaCl.

- 4. Add sample to the Cobalt Chelating Resin and incubate with mechanical rotation for 15-20 minutes at room temperature or 4°C for 60 minutes.
- Collect the sample lysate by gravity flow or centrifuge at 800xg for 1 minute. Save the supernatant to analyze by SDS-PAGE

NOTE: If using tubes, as opposed to column, pellet the resin by centrifugation at 500xg for 2-5 minutes and carefully decant the storage buffer.

6. Wash the resin with 5CV of wash buffer (i.e. $50 \text{mM} \text{ Na}_2 \text{HPO}_4$, 300 mM NaCl pH8.0) supplemented with 20 mM imidazole). Mix with mechanical rotation for 5 minutes and remove the wash buffer as before. Repeat step twice more.

NOTE: The imidazole and sodium chloride is present to reduce non-specific interactions, however these may interfere with the binding of some 6X His tagged proteins. Optimize with 10-50mM imidazole and 100-500mM NaCl.

- Add 2CV of elution buffer (50mM Na₂HPO₄, 300mM NaCl, 250mM Imidazole, pH 8) and mix with mechanical rotation for 5 minutes.
- 8. Collect the eluate as above and repeat the elution four more times. Examine the eluates by SDS-PAGE and pool together the samples of interest.

NOTE: To remove imidazole for downstream applications use gel filtration (G-Biosciences SpinOUT $^{\text{\tiny M}}$ Desalting Columns) or dialysis (G-Biosciences Tube-O-DIALYZER $^{\text{\tiny M}}$).

PROTOCOL FOR DENATURING PROTEINS

- Add an appropriate amount of Cobalt Chelating Resin to a suitable column (suitable to hold 7 columns volumes (CV)). Allow the storage buffer to flow through the column or centrifuge at 800xg for 1 minute. Discard the flow-through.
 NOTE: If using tubes, as opposed to column, pellet the resin by centrifugation at 500xq for 2-5 minutes and carefully decant the storage buffer.
- 2. Add 5CV of distilled water and resuspend the resin. Remove water as above
- 3. Wash the resin by resuspending in 1CV suitable binding buffer (i.e. 50mM Na₂HPO₄, 6M guanidine·HCl, 300mM NaCl pH8.0) supplemented with 10mM imidazole). Remove wash buffer as above and repeat this wash step once.
 NOTE: The imidazole and sodium chloride is present to reduce non-specific interactions, however these may interfere with the binding of some 6X His tagged proteins. Optimize with 0-20mM imidazole and 100-500mM NaCl.
 8M urea can be used as an alternative to the 6M guanidine·HCl.
- 4. Add sample to the Cobalt Chelating Resin and incubate with mechanical rotation for 15-20 minutes at room temperature or 4°C for 60 minutes.
- 5. Collect the sample lysate by gravity flow or centrifuge at 800xg for 1 minute. Save the supernatant to analyze by SDS-PAGE
 NOTE: If using tubes, as opposed to column, pellet the resin by centrifugation at 500xg for 2-5 minutes and carefully decant the storage buffer.
- Wash the resin with 5CV of wash buffer (i.e. 50mM Na₂HPO₄, 6M guanidine·HCl, 300mM NaCl pH8.0) supplemented with 20mM imidazole). Mix with mechanical rotation for 5 minutes and remove the wash buffer as before. Repeat step twice more.
 - **NOTE:** The imidazole and sodium chloride is present to reduce non-specific interactions, however these may interfere with the binding of some 6X His tagged proteins. Optimize with 10-50mM imidazole and 100-500mM NaCl. 8M urea can be used as an alternative to the 6M quanidine·HCl.
- Add 2CV of elution buffer (50mM Na₂HPO₄, 6M guanidine·HCl, 300mM NaCl, 250mM Imidazole, pH 8) and mix with mechanical rotation for 5 minutes.
 NOTE: 8M urea can be used as an alternative to the 6M quanidine·HCl.
- Collect the eluate as above and repeat the elution four more times. Examine the eluates by SDS-PAGE and pool together the samples of interest.
 NOTE: To remove imidazole for downstream applications use gel filtration (G-Biosciences SpinOUT[™] Desalting Columns) or dialysis (G-Biosciences Tube-O-DIALYZER[™]). Samples containing6M guanidine·HCl can be cleaned with G-

Biosciences PAGE-Perfect $^{\text{TM}}$ or must be dialyzed against a buffer containing 8M urea prior to SDS PAGE analysis.

COLUMN REGENERATION

- 1. Wash resin with 10 bed volumes of 20mM MES buffer, pH 5.0
- 2. Wash resin with 10 bed volumes of distilled water.
- 3. Wash resin with 10 bed volumes of 20% ethanol.
- 4. Store resin at 4°C in 20% ethanol.

COLUMN RECHARGING PROTOCOL

Column regeneration should be performed when a different protein is being isolated or when there is a significant loss in the yield of protein. If the Cobalt Chelating Resin loses its blue color the column needs recharging.

- 1. Wash the resin with 5 column volumes of a solution 20mM sodium phosphate supplemented with 0.5M NaCl, 50mM EDTA at pH 7.0.
- 2. Wash with 5 column volumes of distilled water to remove EDTA.
 NOTE: If the loss in yield is suspected to be due to denatured proteins or lipids a more drastic regeneration protocol should be followed. After step 2:
 - A. Elimination of ionic interactions: Wash in batch for approximately 20 minutes in a solution with 1.5M NaCl, follow with a wash with 10 column volumes of distilled water.
 - B. Elimination of precipitated proteins. Wash in batch for at least 2 hours with a solution 1M NaOH, follow with a wash with 10 column volumes of distilled water.
 - C. Elimination of strong hydrophobic interactions: Resuspend the resin in batch with 30% isopropanol and wash for approximately 20 minutes, follow with a wash with 10 column volumes of distilled water.
 - D. Elimination of lipids: Wash in batch for 2 hours with a solution 0.5% of nonionic detergent in 0.1 M acetic acid. Rinse away the detergent with approximately 10 column volumes of 70% ethanol, follow with a wash with 10 column volumes of distilled water.
- 3. Add 5 volumes of 0.1M cobalt(II) sulfate heptahydrate.
- 4. Wash with 5 column volumes of distilled water.
- 5. Add 5 column volumes of the binding buffer. The column is now ready for use. **NOTE:** If storing the column for a while store at 4°C in 20% ethanol.

TROUBLESHOOTING

Issue	Possible Reason	Suggested Solution
	High levels of nucleic	Treat sample with nuclease.
	acids in lysate	LongLife [™] Nuclease, 786-039
Viscous sample	Too little lysis/	
	homogenization buffer	Dilute sample with more buffer
<u> </u>	used	
Column becomes		
clogged after	Sample poorly clarified	Centrifuge the sample at higher speed
sample application	before loading	or filter the sample
аррисацоп	Low protein expression of	Check protein expression levels.
	target protein	Apply larger volume
	Recombinant protein	Increase intensity/ duration of lysis
	targeted to inclusion	Use denaturing conditions (6M
	bodies or possible	guanidine HCl or 8M urea) is protein
	insufficient lysis	is insoluble
	-	Reduce imidazole concentration in
		binding and wash buffers.
No protein found	Target protein in flow-through	Check pH levels of sample and adjust
in elution		to pH7-8
in ciution		Histidine tag may not be accessible.
		Use denaturing conditions or reclone
		with tag at opposite terminus
		Proteolytic cleavage during extraction
		has removed the tag, include protease
		inhibitors (Recom ProteaseARREST [™] , 786-436)
	Elution conditions are too	
	mild	Elute with acidic pH or imidazole step-elution
	Temperature too low	Perform at room temperature
Protein	Temperature too low	Add solubilization agents, such as
precipitates	Aggregate formation	non-ionic detergents, glycerol or
		β-mercaptoethanol
Poor recovery of target protein	Binding capacity of	Increase column size or reduce sample
	column has been exceeded	load
	Strong non-specific	Reduce interactions by including
	interactions of target	detergents, organic solvents or by
	protein on resin	increasing NaCl concentration

	Contaminants in elute	Increase number of binding and wash steps and include 10-20mM imidazole in buffers Prolong wash steps containing imidazole Column too large, reduce amount of resin used
	Strongly bound contaminants elute	Reduce the amount of imidazole in the elution buffer
Poor protein	Contaminants bind target protein through disulfide bounds	Include β-mercaptoethanol, avoid DTT
purity	Contaminants bind target protein through hydrophobic interactions	Add non-ionic detergents or alcohol
	Contaminants bind target protein through electrostatic interactions	Increase the concentration of NaCl
	Recombinant protein degraded	Include protease inhibitors (Recom ProteaseARREST [™] , 786-436)
	Contaminants have similar affinity to target protein	Explore additional chromatography step (Ion exchange, gel filtration)

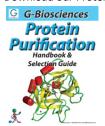
CHEMICAL COMPATIBILITIES

Reagent Effect Comments				
Reagent		Comments		
BUFFER REAGENT	BUFFER REAGENTS			
Tris, HEPES, MOPS	Buffers with secondary and tertiary amines will reduce metal ions	≤50mM secondary and tertiary amines		
Sodium or potassium phosphate	No interference	50mM sodium or potassium phosphate are recommended		
CHELATING REAGENTS				
EDTA, EGTA	Strip metal ions from resin	≤1mM has been used, but care must be taken >1mM causes significant reduction in binding capacity		
REDUCING (SULFHYDRYL) REAGENTS				
B-mercaptoethanol	Reduces disulfide cross- linkages Can reduce metal ions	≤20mM		
DTT, DTE, TCEP	Low concentrations will reduce metal ions	1mM maximum, but recommend β-mercaptoethanol		

DETERGENTS		
Non-ionic detergents (Triton, Tween, NP-40, etc.)	Removes background proteins and nucleic acids	≤2%
Cationic detergents (CTAB)	Improves membrane and lipid associating proteins or hydrophobic proteins solubility	≤1%, be carefully of protein precipitation
Zwitterionic detergents (CHAPS, CHAPSO)	Solubilizes membrane proteins	≤1%
Anionic detergents (SDS, Sarkosyl)	Strips metal ions Selective solubilization membrane proteins	Not recommended
DENATURANTS		
Guanidine·HCl	Solubilize proteins	≤6M
Urea	Solubnize proteins	≤8M
AMINO ACIDS		
Glycine, Glutamine, Arginine		Not recommended
Histidine	Binds resin and competes with 6X His tag histidines	Low (20mM) concentrations can block non specific binding and high (>100mM) concentrations will elute His tagged proteins.
OTHER ADDITIVE		
Sodium chloride (NaCl)	Reduces non-specific protein binding through ionic interactions	≤2M, at least 300mM NaCl should be included in buffers
Magnesium chloride (MgCl ₂)	Required for purification of Ca ²⁺ binding proteins	≤4M
Calcium chloride (CaCl ₂)	Essential metal cofactor for nucleases	≤5mM
Glycerol	Prevents hydrophobic	≤40%
Ethanol	interactions between proteins	<i>≤</i> 20%
Imidazole	Binds resin and competes with 6X His tag proteins for metal ions	Low (<25mM) concentrations can block non specific binding and high (>100mM to ≤500mM) concentrations will elute His tagged proteins
Citrate	Carboxylic side chains may potentially act as chelation site for metal ions, causing metal leakage	≤60mM

RELATED PRODUCTS

Download our Protein Purification Handbook.



http://info.gbiosciences.com/complete-protein-purification-handbook/

For other related products, visit our website at www.GBiosciences.com or contact us.

Last saved: 5/19/2015 CMH



www.GBiosciences.com