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(FOR RESEARCH USE ONLY. DO NOT USE IT IN CLINICAL DIAGNOSIS !)

Co Focurose 6FF (IMAC) (Co-6FF (IMAC))

Catalog No: E-CM-AF09

This manual must be read attentively and completely before using this product.

May you have any problems, please contact our Technical Service Center for help.

Phone: 240-252-7368(USA) 240-252-7376(USA) Email: <u>techsupport@elabscience.com</u> Website: <u>www.elabscience.com</u>

Please kindly provide us the lot number (on the outside of the box) of the kit for more efficient service.

Please read this manual carefully before use to ensure the performance and successful operation. If you have any questions, please contact our Technical Support.

Product introduction

Co-6FF (IMAC) can be used for separation and purification with the interactions between Co^{2+} and some amino acids (mainly include histidine, cysteine, tryptophan) on the side chain of protein, and it is suitable for separation and purification of His-tagged protein and biological molecules which interact with Co^{2+} .

Advantages

- 1. Quick and easy (one-step purification).
- 2. Wide application, simple-operation. Suitable for gravity column and prepacked column.
- 3. Lower Co²⁺ abscission probability and wider compatibility with reagents (Table 2) when compared with Ni-6FF (IDA).

Matrix	Highly cross-linked 6% agarose	
Particle size range	45-165 μm	
Average particle size	90 µm	
Binding capacity	45 mg (His-tagged protein)/mL (media)	
pH stability*	3-12 (long-term)	
	2-14 (short term)	
Chemical stability*	0.01M HCl, 0.01M NaOH (1 week)	
	1M NaOH, 70% Ethanol (12 hours)	
	2% SDS (1 hour)	
	30% Isopropyl alcohol (30 minutes)	
Flow rate	300-600 cm/h	
	(0.3MPa, XK16/40. Column bed height: 30 cm)	
Pressure	≤ 0.3 MPa	
Storage buffer	20% Ethanol	
Storage temperature	4~8℃	

Table 1: Performance index

*: The stability of media when unchelated with metal ions.

companying with common reagents			
	0.05M sodium phosphate, pH 7.4		
Buffer	0.1M Tris-HCl, pH 7.4		
	0.1M Tris-acetate, pH 7.4		
	0.1M HEPES, pH 7.4		
	0.1M MOPS, pH 7.4		
	0.1M sodium acetate, pH 4		
Denaturant	8M Urea		
	6M Gua-HCl		
Detergent	2% Triton X-100		
	2% Tween 20		
	2% NP-40		
	2% Cholate		
	1% CHAPS		
	0.005M DTE		
	0.005M DTT		
Reductant*	$0.02M \beta$ -mercaptoethanol		
	0.005M TCEP		
	0.01M reduced glutathione		
	0.5M Imidazole		
	20% Ethanol		
	50% Glycerol		
Other additives	0.1M Na ₂ SO ₄		
	1.5M NaCl		
	0.001M EDTA**		
	0.06M Citrate		

 Table 2: Compatibility with common reagents

* During the operation with Co-6FF (IMAC), it is allowed to add low concentration of reductant, but it must be avoided of being immersed or stored for a long time with solution with reductant.

** During the operation with Co-6FF(IMAC), it is allowed to add nominal concentration of metal ions chelating agent (e.g. 0.0001M EDTA) into samples of small volume, but it must be avoided of adding or loading a large volume of samples containing chelating agent into the purified solution.

Operation (take 1 mL column and 5 mL column for example)

1. Wash (water)

Wash the media with 5~10 CV (column volume) of purified water with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL).

Note: This operation is used to remove the 20% ethanol in media.

2. Equilibration

Balance the media with 5~10 CV of equilibrium liquid with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL) until the baseline turns stable and then set to zero.

Note: This procedure is used to balance the media. Make sure that the pH and component in media are in accordance with sample.

3. Sample application

Apply the sample with flow rate of 0.2 mL/min (1 mL) or 1.0 mL/min (5 mL) after centrifugation and filtration (0.45 μ m). Wash with the equilibrium liquid until the baseline tends to zero.

Note: The binding capacity of proteins varies with the type of lysates, the properties of target proteins, flow rate, temperature, and pH. Low flow rates often increase the binding efficiency of samples.

4. Wash

Wash out the impurity substance with 5~10 CV of washing solution with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL) and collect the washed solution.

Note: The washing solution is used to wash out some non-specific adsorbed impurity proteins.

5. Elution

Elute with 5~10 CV of eluent with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL) and collect the eluted solution.

Note: Low flow rate may increase the concentration of target protein in the eluted solution.

6. Wash with water

Wash the media with 5~10 CV of purified water with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL).

Note: This procedure is used to remove the eluent in media.

7. Storage

Wash the media with 5~10 CV of 20% ethanol with a flow rate of 0.5 mL/min (1 mL) or 2.0 mL/min (5 mL) and the store the media.

Note: 20% ethanol can prevent the growth of microorganism. Media preserved with 20% ethanol can be stored at $4 \sim 30^{\circ}$ C ($4 \sim 8^{\circ}$ C is preferred).

8. Preparation of buffer

Add 8M urea or 6M guanidine hydrochloride to the following equilibrium liquid, washing buffer and eluent if the sample is inclusion body.

Equilibrium liquid: 0.02M PB, 0.5M NaCl, adjust the pH to 7.4. Store the prepared equilibrium liquid at room temperature.

Note: NaCl in equilibrium liquid is used to inhibit the ion exchange of media.

Washing buffer: 0.02M PB, 0.5M NaCl, 0.02-0.04M iminazole, adjust the pH to 7.4. Store the prepared washing buffer at room temperature.

Note: Add 0.02-0.04M iminazole to the washing buffer (purity preferred) or to the equilibrium liquid directly (yield preferred) according to the final application demands.

Eluent: 0.02M PB, 0.5M NaCl, 0.5M iminazole, adjust the pH to 7.4. Store the prepared eluent at room temperature.

Note: Concentration of iminazole in eluent between the range of 0.05-0.25M is enough to elute the target protein in general condition.

Cleaning

The excellent performance of media (e.g. loading ability, mobility, column efficiency, etc.) can be recovered after cleaning the strong coupling substance (e.g. some strong coupling protein, denatured protein, lipids, etc.).

It is recommended to wash the media after used for each 5 times. The exact washing frequency should be adjusted according to the cleanliness of the purified sample.

1. Wash the media with $5 \sim 10$ CV of purified water.

Note: This procedure is used to remove the eluent (wash after using) or 20% ethanol (wash before using).

 Wash the media with 5~10 CV of 0.02M Tris-HCl, 0.1M EDTA (pH 8.0), then wash the media with 5~10 CV of purified water immediately. Note: This procedure is used to remove the Co²⁺.

3. Wash the media with 5~10 CV of 1.0M NaOH, Wash the media until the pH to neutral with purified water after standing for 1 hour.

Note: This procedure is used to remove the precipitates or lipids accumulated in the media.

4. Wash the media with 5~10 CV of 0.1M NiSO₄, then wash the media with 5~10 CV of purified water after standing for 0.5 hour.

Note: This procedure is used to couple Co^{2+} .

Store the media after washed with 5~10 CV of 20% ethanol.
 Note: 20% ethanol can prevent the growth of microorganism. Media preserved with 20% ethanol can be stored at 4~30°C (4~8°C is preferred).

Problem **Possible cause** Suggestion Overloading of sample volume. Decrease the sample volume. Speed of sample loading is too Reduce the flow speed of sample loading. fast. Protein or lipids accumulate in the Wash the media timely and media. effectively. The target compound The expression condition is too An empty vector is suggested to does not combine with severe that the His-tag was set as control of expression and the media or the enwrapped and cannot combine purification to check whether the combining amount is low with the media. expression condition is suitable. The initial sample does not when purifying Verify through the gene sequence contain His-tagged protein. or His-tag antibody. The target protein has not been expressed successfully, or the pH The target protein occurs in the and components in sample and flowthrough. equilibrium liquid were incorrect. The target compound does not Confirm whether the target combine with the media or the combine with media or not. combining amount is low. Increase the iminazole Unsuitable elution condition. concentration in eluent. Decrease the flow rate and Insufficient elution time. prolong the retention time of No target compound was eluent. collected or only a small Insufficient elution volume. Increase the elution volume. amount of target Decrease the iminazole Target protein was washed out compound was collected during the washing. concentration in washing buffer. Check the stability of target compound in the elution buffer The target compound accumulates (salt concentration, pH, etc.) Add in the elution buffer. some additives into the eluent. such as 0.2% Triton X-100 or 0.5% Tween 20.

Trouble shootings

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	Sample has not been pretreated.	Samples must be centrifuged or filtered before loading.
-		Dilute the sample properly with
	High viscosity of sample.	equilibrium liquid to decrease the
	inger (isobility of sumple.	viscosity.
-		Increase the washing volume until
	Insufficient washing.	the baseline smooth and keep
]]]		consistence with equilibrium
		liquid.
	Impurity protein or lipids	Wash the media timely and
	accumulate in the media.	effective.
		Purify with other type of media
	Impurity substances have	(e.g. ion–sieve or
1 7 0	strong-affinity with Co ²⁺ .	molecular-sieve).
compound		Determine the stability of target
	The target compound degrades.	compound
	Bad loading effect of column	
	resin.	Reload or re-purchase.
	Non-specificity absorption of	Add appropriate additive to
		reduce the non-specific
		absorption, such as 0.5% Triton
1	impurity materials.	X-100, 1.0% Tween 20 or 50%
		glycerol.
, r	The top of separation column has	Reload the column or reduce the
2	a large volume of sample.	volume of the sample
	There is microbial grow in the	Correctly store the media after
	media.	used.
	Speed of sample loading is too	Reduce the flow speed of sample
f	fast.	loading.
	Protein or lipids accumulate in the	Wash the modie timely
	media.	Wash the media timely.
Decrease of loading	Ligand dropped off due to	Re-couple with new media.
volume	excessive use.	
	The expression condition is too	An empty vector is suggested to
5	severe that the His-tag was	set as control of expression and
	severe that the His-tag was enwrapped and cannot combine	set as control of expression and purification to check whether the

The chromatographic peak rises slowly	The media was loaded too tight.	Reload the column.
The chromatographic peak trails	The media was loaded too loose.	Reload the column.
The column bed cracks or being dry	Leakage occurred or a large volume of bubbles was introduced.	Check whether there is leakage or bubble, reload the column.
Flow of the column is exceedingly slow	Protein or lipids accumulate in the media.	Wash the media or filter membrane timely.
	Protein precipitates in the media.	Adjust the content of equilibrium liquid and wash buffer to maintain the stability of target compound and combining efficiency of media.
	There is microbial grow in the media.	Filter and degas all the reagents. Samples must be centrifuged or filtered before loading.