

His-tag Protein Purification Gel, 45-135 µm-NTA-Cu PRODUCT DATA SHEET

His-tag Protein Purification Gel, 45-135 µm-NTA-Cu

Cat No: AG-His-4500-NTA-Cu

Description

His-tag protein agarose gel (NTA-Cu) can be used for the purification of histidine labeled (6xHis-tag) proteins from various expression sources, such as E. coli, yeast, insect cells, and mammalian cells. It is made from a highly cross-linked 4% agarose gel as a matrix chemically coupled to tetra-coordinated nitrogen triacetic acid (NTA). After chelating copper ions (Cu²⁺), a relatively stable planar quadrilateral structure can be formed, so that more sites can continue to coordinate with the imidazole ring on the histidine label to achieve the effect of binding the target protein.

Matrix	Highly crosslinked 4% agarose gel
Ligand	Triacetic acid (NTA)
Chelating metal ions	Cu ²⁺
Particle size	45~135 μm
Concentration	The gel volume accounts for 50% of the suspension volume
Pressure flow rate	80 ~ 150 cm/h (0.3 MPa, 3 bar)
Carrying capacity	> 60 mg 6xHis protein
Storage buffer	1× PBS containing 20% ethanol
Shelf life	Stable storage at 2 ~ 8°C for two years

Characteristics

Operation process

1. Buffer preparation

Binding/balancing buffer, washing buffer and eluting buffer, please prepare or purchase our His-tag protein purification kit.

2. Sample preparation

Take E. coli expression system, 500 mL induced bacterial solution as an example.

1) Centrifuge at 4°C for 30 min (4000x g) to collect bacteria and discard supernatant.



2) Re-suspend the bacteria with a pre-cooled **binding/balancing buffer**, adding appropriate inhibitors, such as protease inhibitors (PMSF) or other protease inhibitors, if needed.

Note: The added inhibitors should not affect the performance of **His-tag protein agargo gel (NTA-Cu)**. The crushing solution should not contain chelating agents such as EDTA and EGTA, reducing agents such as DTT and mercaptoethanol, and denaturants such as urea and guanidine hydrochloride.

3) Use ultrasonic crushing method to break the bacteria on the ice until the sample is broken completely.

Optional: If the lysate is too thick, RNase A (final concentration 10 μ g/mL) and DNase I (final concentration 5 μ g/mL) can be added and incubated on ice for 10 ~ 15 min.

4) Centrifuge at 4°C for 20 min (12,000x g), separate supernatant and precipitation, and filter for impurity removal. Retain samples of supernatant and precipitation for subsequent testing.

3. Purification of recombinant His-tag fusion protein

1) Gently resuspension His-tag protein agarose gel (NTA-Cu).

2) Absorb 2 mL of **His-tag protein agarose gel (NTA-Cu)** and add it to the chromatographic column, and balance **His-tag protein agarose gel (NTA-Cu)** with 10 mL **binding/balancing buffer**. Repeat the preceding steps one more time.

3) Close the bottom outlet of the chromatographic column, add the prepared supernatant containing His-tag protein to the chromatographic column, and then cover the upper inlet of the chromatographic column tightly, and it is recommended to seal with a sealing film. Place on the mixer and incubate at room temperature for $1 \sim 2$ h. (Can also be incubated at $2 \sim 8^{\circ}$ C $2 \sim 4$ h or overnight)

4) After the end of incubation, open the upper and lower inlet and outlet of the chromatographic column, and collect the supernatant after all the supernatant flows out of the chromatographic column, and put it at $2 \sim 8^{\circ}$ C as a flow through for subsequent detection.

5) Immediately add 10 mL **washing buffer** to the chromatographic column, collect the washing liquid and place it at 2 ~ 8°C for subsequent testing. Repeat the preceding steps four times.

6) **Routine elution:** Add 1 mL **eluent buffer** and collect eluent with 1.5 mL Ep tube. Collect 5 to 10 tubes respectively.

Note: Gradient elution can also be used if the elution procedure is optimized.

Gradient elution: Different concentrations of imidazole were used for elution and eluent was collected.

7) SDS-PAGE detection



The resulting sample (including run-off, washing solution and eluent) and the original sample were tested for purification using SDS-PAGE. Add an appropriate amount of **protein fast dyeing solution** to immerse the PAGE glue, and then put it on the shaking table, dyeing $10 \sim 30$ min to observe the results.

Note: The target protein should be dialysis or ultrafiltration to remove imidazole and other impurities before storage, and then subpackaged and frozen to -80°C.

(Optional) Gel regeneration and storage

Gel regeneration steps please refer to or directly buy our company His-tag protein purification regeneration kit.

After the gel is regenerated, it can be used immediately, if not used immediately, it is necessary to suspend the gel in an equal volume of 20% ethanol and store it at $2 \sim 8^{\circ}$ C.

Problem solving

There was no target protein	The protein may be an	The lysate can be detected
in the eluent	inclusion body, but the	by electrophoresis to
	supernatant has no protein	determine whether the
		supernatant contains the
		target protein, and the
		inclusion body protein needs
		to be purified according to
		the inclusion body protein.
	Underexpression	Optimize expression
		conditions.
	The target protein is weakly	Reduce imidazole
	bound and washed off	concentration.
	during the scrubbing step	
	The target protein is	Add appropriate protease
	degraded by protease	inhibitors, such as PMSF, to
		the cleavage step or washing
		step.
	The target protein cannot be	Increase the imidazole



	effectively eluted from the	concentration.
	gel	Use a 10-100 mM EDTA
		solution to strip copper ions
		and obtain the target
		protein.
The purified target protein is	The laundry is not thorough	Increase the number of
not pure		washes.
	The sample contained other	The cleaning conditions
	histidine label proteins	were optimized by adjusting
		the concentration of
		imidazole. The elution
		components are then
		further purified by using
		other purification methods
		(e.g., ion exchange,
		hydrophobic, etc.).
Protein precipitation occurs	Concentration is too large	Dilute protein appropriately.
during binding	Protein aggregation	Add a stabilizer, such as
		0.1% Triton X-100 or Tween-
		20, to the sample and all
		buffers.
	Operating temperature is	Operate at 2-8°C.
	too high	

Ordering Information

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