

Anti-FLAG (DYKDDDDK) Tag agarose Beads

Properties

Ligand: Anti-FLAG (DYKDDDDK) Tag

Binding capacity: 12-15 μ g of recombinant 1xFlag tagged protein per 25 μ L bead slurry

Reactivity: Specifically binds to flag-tag sequence. Compatible with N-, C-terminal.

Bead size: 45-165 µm (cross-linked 4 % agarose beads)

Applications: IP, Protein purification.

Buffer compatibility: See Wash buffer compatibility table.

Stability: Stable storage at 2 to 8 °C for 1 year.

Required buffer solutions

RIPA buffer: 50 mM Tris-HCl pH 8.0, 150 mM NaCl, 0.1 % SDS, 1 % NP-40, 0.5 % deoxycholate

Dilution Buffer: 10 mM Tris-HCl pH 7.4, 150 mM NaCl, 0.5 mM EDTA

- Wash buffer: 10 mM Tris-HCl pH 7.4, 150 mM NaCl, 0.05 % Tween[™] 20, 0.5 mM EDTA.
- **2x SDS buffer**: 100mM Tris-HCl pH 6.8, 20 % glycerol, 4% SDS, 0.04 % bromophenol blue, 10 % β -mercaptoethanol (or 200mM DTT).

Acidic elution buffer: 200 mM glycine pH 2.5.

Peptide elution buffer: 150 µg/mL FLAG-peptide.

Neutralization buffer: 1 M Tris pH 10.4.



Wash buffer compatibility table

Buffer ingredients	Max. concentration
β-mercaptoethanol	10 mM
DTT	10 mM
NaCl	tested up to 1M
SDS	0.2 %
Urea	2 M
Triton-X100	tested up to 0.5 %
Tween™ 20	tested up to 5 %

Immunoprecipitation protocol

Take 300 μ g mammalian cell lysate as an example and start the immunoprecipitation experiment from the protein binding step.

Mammalian cell lysis

1. Selection of lysis buffer

- \cdot For cytoplasmic proteins, resuspend the cell pellet in 300 μL RIPA buffer containing protease inhibitors and 1mM PMSF.
- \cdot For nuclear/chromatin proteins, add DNaseI, MgCl2, protease inhibitor cocktail, and 1mM PMSF to RIPA buffer, then resuspend the cell pellet in 300 μL of RIPA buffer.
- 2. Place the tube on ice for 30 minutes and mix the suspension every 10 minutes.
- 3. Centrifuge the cell lysate at 15,000x g for 10 minutes at 4°C. Transfer the supernatant to a pre-chilled EP tube and add 300 μ L Dilution buffer. If required, save 50 μ L of the diluted lysate for further analysis (input control).

Equilibration & Protein binding

- 1. Resuspend beads by pipette. **Do not vortex the beads**.
- 2. Transfer 25 μL of bead slurry to a 1.5 mL centrifuge tube containing 600 μL pre-chilled dilution buffer.
- 3. Centrifuge at 3000xg for 5 minutes at 4°C to collect beads and discard the supernatant.
- 4. Add the diluted lysate to the centrifuged beads and rotate at 4°C for 1 hour.

Washing

- 1. Centrifuge at 3000x g for 5 minutes at 4°C to pellet the beads.
- 2. If required, retain a portion of the supernatant for further analysis and discard the



remainder.

- 3. Add 600 μL wash buffer to resuspend the beads.
- 4. Centrifuge at 3000x g for 5 minutes at 4°C to pellet the beads and discard the supernatant.
- 5. Repeat step 3 & 4 at least four times.
- 6. Last washing step, transfer the beads to a new tube.

Elution with 2x SDS buffer

- 1. Discard the remaining supernatant.
- 2. Add 20 μ L 2x SDS buffer to resuspend the beads.
- 3. Heat at 100°C for 5 minutes to separate the immune complexes from the beads.
- 4. Centrifuge at 3000x g for 2 minutes at 4°C to pellet the beads.
- 5. Analyze the supernatant by SDS-PAGE/WB.

Note: For Western blot detection.

Elution with Acidic elution buffer

1. Remove the remaining supernatant.

- 2. Add 40-80 μ L of acidic elution buffer and incubate for 5-10 min at room temperature.
- 3. Centrifuge at 3000 x g for 2 minutes at 4°C to pellet the beads.
- 4. Transfer the supernatant to a new tube.
- 5. Immediately neutralize the eluate with 4-8µL of neutralization buffer.
- 6. Repeat this step at least once to increase elution efficiency.

Elution with Peptide elution buffer

- 1. Prepare the 1 × FLAG (DYKDDDDK) peptide to 150μ g/mL in PBS.
- 2. Remove the remaining supernatant from the beads.
- 3. Add 100 μ L of the 1 × FLAG (DYKDDDDK) peptide and mix at room temperature for 20-30 minutes, or at 4°C for 30-60 minutes.
- 4. Centrifuge at 3000 x g for 2 minutes at 4°C to pellet the beads.
- 5. Transfer the supernatant to a new tube.
- 6. Repeat this step at least once to increase elution efficiency.