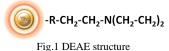


DEAE Magnetic Beads

Magnetic spheres segregate proteins or nucleic acids with the use of beads-adsorbent technique as a chromatographic matrix. Ion exchange chromatography is extensively employed to segregate or refine a targeted molecule from raw biological materials. The molecules are discriminated based on differences in their available surface charges utilizing extremely gentle binding and eluting conditions to preserve their biological activity intact.

BcMagTM DEAE Magnetic Beads are uniform magnetic resins grafted with a high density of diethyl aminoethyl (DEAE) functional groups on the surface (Fig.1). The weak anion exchange magnetic bead-based format enables rapid high-yield processing of 96 samples in about 20 minutes. It can quickly fraction proteins or nucleic acids from complex biological samples (such as serum, plasma, etc.) manually or automatically. The purified protein can be used in downstream applications such as sample fractionation for 1D and 2D SDS-PAGE, X-ray crystallization, and NMR spectroscopy. The Weak Ion Exchange resins allow the rapid release of very strong ions that may be retained irreversibly on Strong Ion Exchange beads. Additionally, weak ion exchangers can be effective separation tools when strong ion exchangers fail because the selectivity of weak and strong ion exchangers frequently differ.



6

Weak anion exchange magnetic resins are used to replace time-consuming, complex, and costly chromatographic procedures such as agarose, cellulose, Sepharose, and Sephadex-based columns or resins. In column-based procedures, the lysate is centrifuged or cleared, the supernatant is added to the column, the membrane or resin is washed with buffer through centrifugation or vacuum manifold, and the required biomolecules are eluted in an adequate volume of buffer. When using column-based technologies, processing multiple samples in academic research labs may necessitate a significant quantity of hand pipetting. This pipetting can discourage differences in target biomolecule yield between experiments and people. Staff and students may require extensive training and practice to produce constant protein yields.

Weak anion exchange magnetic resins have significant advantages over non-magnetic resin technologies. It is due to the numerous benefits of magnetic resins, such as their ease of use, rapid experimental protocols, suitability, and convenience for high throughput automated and miniaturized processing. They thus see increasing use in various areas of life-sciences research and development, including drug discovery, biomedicine, bioassay development, diagnostics, genomics, and proteomics.

Weak anion exchange beads feature and benefits:

- Fast and simple— DEAE magnetic beads-based format eliminates columns or filters or a laborious repeat of pipetting or centrifugation.
- **Convenient and expandable**—Magnetic format enables high-throughput processing of multiple samples in parallel with many different automated liquid handling systems.
- Robust— DEAE Magnetic beads do not crack or run dry.
- Low bed volume—Working with small magnetic bead volumes allows for minimal buffer volumes, resulting in concentrated elution fractions.

DEAE magnetic beads Applications:

- Protein pre-fractionation in cell lysates
- · Optimizing purification conditions for new protein preparation protocols
- Protein purification and concentration
- Antibody purification from serum, ascites, or tissue culture supernatant
- Preparation of samples before 1D or 2D PAGE
- Phosphopeptide purification before MS analysis

1



Specificities			
Composition	Magnetic beads grafted with diethylaminoethyl (DEAE) groups.		
Magnetization	~45 EMU/g		
Type of Magnetization	Superparamagnetic		
Effective Density	2.0 g/ml		
Stability	Most organic solvents		
	1 µm beads: ~2 mg BSA/ ml of Beads		
DEAE beads	5 µm beads: ~1.5 mg BSA/ ml of Beads		
Storage	Store at 4°C upon receipt		

Protocol

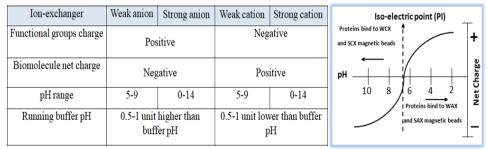
Note: The following protocol is an example of fractionating a protein or peptide sample with BcMag[™] DEAE magnetic beads. Users are encouraged to choose alternative binding, washing, or elution buffers to get the best results and determine the optimal working conditions based on the protocol and suggestions described in the troubleshooting section. It is critical to match the amount of the beads to the amount of protein in the starting material in all protein purification experiments. It is not only for financial reasons but also because insufficient DEAE resin results in inadequate protein binding in the solution. Too many affinity binding sites will result in the binding of other proteins, making the purification less selective and requiring extra purification steps to achieve pure protein. We recommend performing a titration to optimize the beads used for each application. Should scale volumes of elution to avoid unnecessary sample dilution.

Note: Select the appropriate buffer

• Based on the protein's pI, empirically calculate the appropriate buffer (pH and salt concentration) for purifying and eluting the protein of interest Fig.2). In a buffered solution above the protein's pI, the protein becomes negatively charged (deprotonated) and binds to the positively charged functional groups of an anion exchange resin. To choose the correct buffer for a selected pH, the following is a general rule for selecting a buffer pH:

Anion exchanger — 0.5–1.5 pH units higher than the protein's pI of interest.

Cation exchanger - 0.5-1.5 pH units lower than the protein's pI of interest.





A. Equipment

• Magnetic rack (for manual operation)

Based on sample volume, the user can choose one of the following magnetic racks: BcMag magnetic rack-2 for holding two individual 1.5 ml centrifuge tubes (Cat. # MS-01); BcMag magnetic rack-6 for holding six individual 1.5 ml centrifuge tubes (Cat. # MS-02); BcMag magnetic rack-24 for holding twenty-four individual 1.5-2.0 ml centrifuge tubes (Cat. # MS-03); BcMag magnetic rack-50 for holding one 50 ml centrifuge tube, one 15 ml centrifuge tube, and four individual 1.5 ml centrifuge tubes (Cat. # MS-04); BcMagTM magnetic rack-96 for holding a 96 ELISA plate or PCR plate (Cat. # MS-05). For larger scale purification, Ceramic magnets Block for large scale purification (6 in x 4 in x 1 in block ferrite magnet, Applied Magnets, Cat# CERAMIC-B8)

- Corning 430825 cell culture flask for large-scale purification (Cole-Parmer, Cat#EW-01936-22)
- Mini BlotBoy 3D Rocker, fixed speed, small 10" x 7.5" platform w/ flat mat (Benchmark Scientific, Inc. Cat# B3D1008) or compatible

2



B. Buffer

- Binding/Wash Buffer: Binding/Wash Buffer: 10 mM Tris-HCl, pH 8.0
- Elution Buffer: 50 mM Sodium phosphate pH 8.0, 0.1-1.0 M NaCl

General Protocol for using the Weak anion exchange magnetic beads.

a. BcMagTM DEAE magnetic beads preparation

1. Vigorously shake the bottle until the magnetic resins become homogeneous and transfer an appropriate volume of the magnetic resins from the bottle to a new tube or flask.

Note:

- Optimize the number of resins used for each application. Insufficient resins will lead to lower yields. Too many beads will cause higher background.
- Do not allow the resins to sit for more than 3 minutes before dispensing. Resuspend the magnetic beads every 3 minutes.
- Place the tube on the magnetic rack for 1-3 minutes. Remove the supernatant while the tube remains on the rack. Add ten bead-bed volumes of dH₂O and mix the beads by pipetting or vortex. Again, place the tube on the magnetic rack for 1-3 minutes and remove the supernatant while the tube remains on the rack.
- 3. Repeat step (2) one more time.
- 4. Equilibrate the beads by adding ten bead-bed volumes of Binding/Washing buffer and shake it to mix them. Incubate at room temperature with continuous rotation for 2 minutes. Place the tube on the magnetic rack for 1-3 minutes. Remove the supernatant while the tube remains on the rack. The resins are ready for purification.

b. Purification

- 1. Add the equilibrated beads (Step a (4) to the sample and incubate on Mini BlotBoy 3D Rocker with continuous rotation for 5-10 minutes.
- 2. Place the tube on the magnetic rack for 1-3 minutes. Remove the supernatant while the tube remains on the rack. Add ten bead-bed volumes of Binding/Washing buffer and shake it ten times to wash the beads. Again, place the tube on the magnetic rack for 1-3 minutes and remove the supernatant while the tube remains on the rack.
- 3. Repeat step (2) six times.

Note:

- This step is critical to get high pure protein. It may be necessary to wash the beads more than six times for some proteins to reduce the nonspecific binding.
- Optimize the washing buffer (pH and salt concentration)

Elute protein with an appropriate volume of elution buffer by pipetting up and down 10-15 times or vortex mixer for 5 minutes.

Note: Determine the optimum elution buffers (pH and salt concentration) and eluting the protein 2-3 times may be necessary.

4. Elute protein with an appropriate volume of elution buffer by pipetting up and down 10-15 times or vortex mixer for 5 minutes.

Note: Determine the optimum elution buffers (pH and salt concentration) and eluting the protein 2-3 times may be necessary.

5. Collect and transfer the supernatant to a new tube.

Troubleshooting

Problem Possible Causes		Suggestions	
Low yield	The sample's ionic strength is high.	• The sample should be dialyzed, desalted, or diluted in a salt ≤25 mM purification buffer.	
	The sample contains interfering detergents.		
The protein failed to	Ionic interaction is too strong.	Increase the NaCl concentration.	
elute.		• Decrease pH of the Elution Buffer.	
		Using Weak Anion Exchange Magnetic Beads	
	Carry-over between eluted fractions	Add more wash steps between each elution step	

3



Poor separation	Proteins or peptides with similar pI to the target protein	•	Optimize NaCl concentration ar Buffer	nd/or pH of the Elution
	Related P	Related Products		
	Product Name		Product Name	

Product Name	Product Name
DEAE Magnetic Beads	SCX Magnetic Beads
PSA Magnetic Beads	PEI Magnetic Beads
WCX Magnetic Beads	Hydroxyapatite Magnetic Beads
SAX Magnetic Beads	