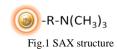
Instruction Manual

Strong Anion Exchange (SAX) Magnetic Beads

Magnetic Beads fractionate proteins or nucleic acids using beads-adsorbent technology as a chromatographic matrix. Ion exchange chromatography is widely used to separate or purify a target molecule from crude biological materials. The molecules are separated based on variations in their accessible surface charges utilizing very light binding and eluting conditions for intact biological activity,

BcMagTM Strong Anion Exchange (SAX) Magnetic Beads (Fig.1) are uniform magnetic resins grafted with a high density of quaternary amine functional group (Strong Anion Exchange) on the surface. The strong 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. Additionally, Strong ion exchangers can be effective separation tools when weak ion exchangers fail because the selectivity of weak and strong ion exchangers frequently differ.

Strong 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.

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.

Strong anion exchange beads feature and benefits:

- Fast and simple— 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—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

Strong anion exchange 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

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Specificities				
Composition	Magnetic beads grafted with quaternary amine functional group groups.			
Magnetization	~45 EMU/g			
Type of Magnetization	Superparamagnetic			
Effective Density	2.0 g/ml			
Stability	Most organic solvents			
	1 μm beads: ~3 mg BSA/ ml of Beads			
Strong anion exchange beads	5 μm beads: ~2 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™ SAX 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 Strong Anion Exchange 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. It is necessary to optimize 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.

Ion-exchanger	Weak anion	Strong anion	Weak cation	Strong cation
Functional groups charge	Positive		Negative	
Biomolecule net charge	Negative		Positive	
pH range	5-9	0-14	5-9	0-14
Running buffer pH	0.5-1 unit higher than buffer pH		than 0.5-1 unit lower than buffer pH	

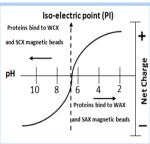


Fig.2

A. Equipment

Magnetic rack (for manual operation)

Based on sample volume, the user can choose one of the following magnetic racks: BcMagTM magnetic rack-2 for holding two individual 1.5 ml centrifuge tubes (Cat. # MS-01); BcMagTM magnetic rack-6 for holding six individual 1.5 ml centrifuge tubes (Cat. # MS-02); BcMagTM magnetic rack-24 for holding twenty-four individual 1.5-2.0 ml centrifuge tubes (Cat. # MS-03); BcMagTM 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)



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- 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

B. Buffer

- Binding/Wash Buffer: 25 mM Tris•HCl, pH 8.0. The pH of the Binding/Wash Buffer should be at least one pH unit away from the pI of the target protein or peptide. (Note: For best results, the salt should be ≤ 25mM in the sample).
- Elution Buffer: 25 mM Tris-HCl pH 8.0. To elute the target protein or peptide from the magnetic beads, the user should optimize elution conditions for individual application by stepwise elution using solutions with increasing salt concentration, i. e., increase stepwise to a final salt concentration of 2.5 M).

General Protocol for using the strong anion exchange magnetic beads.

a. BcMagTM Strong Anion Exchange 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.
- 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 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

- Add the equilibrated beads (Step a (4) to the sample and incubate on Mini BlotBoy 3D Rocker with continuous rotation for 5-10 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 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.



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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
Poor separation	Proteins or peptides with similar pI to the target protein	Optimize NaCl concentration and/or pH of the Elution Buffer

Related Products				
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				