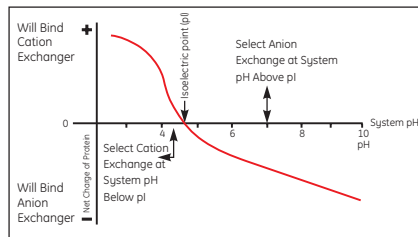


Selection Guide – Ion Exchange Media

Selecting an anion or cation exchanger

Ion exchange separates proteins on the basis of differences in their net surface charge in relation to pH of the surroundings. The figure here illustrates how the net charge of a protein can vary with pH. Every protein has its own charge/pH relationship.



If isoelectric point (pI) of the target protein is known:

- select an anion exchanger (Q, DEAE, ANX) with a buffer pH above the pI.
- select a cation exchanger (S, SP, CM) with a buffer pH below the pI.

If pI is unknown:

- test for selectivity using a strong ion exchanger, Q, S or SP. Strong ion exchangers maintain their charge over a wider pH range than weak ion exchangers and are suitable for most applications.

Polishing

Remove trace impurities or closely-related substances
Sample condition:
almost pure

Highest resolution
µg/run

Highest resolution
mg/run

High resolution
High throughput
Easy scale-up

High resolution
Easy scale-up

Easy scale-up
Broad choice of selectivity, including
alternatives to Q or S ion exchange media

High volume through-put and high capacity
Easy scale-up

High binding capacity for selected proteins
Easy scale-up

Large scale, viscous samples

Industrial scale, filtration and
capture in one step

Intermediate purification

Remove bulk impurities
Sample condition:
partially purified

Capture

Isolate, concentrate and
stabilize target protein(s)
Sample condition:
clarified or
non-clarified

Start here

MiniBeads
(Q or S)



Use for intermediate purification if column capacity is sufficient and no scale-up is required.

MonoBeads
(Q or S)



Use for intermediate purification if column capacity is sufficient and no scale-up is required. Can be used for capture steps if sample is free from particulate matter.

SOURCE 15
(Q or S)



Use SOURCE 15 when resolution is top priority.

SOURCE 30
(Q or S)



Use SOURCE 30 when speed is top priority.

Sepharose High Performance
(Q or SP)



Use HiTrap columns prepacked with Sepharose High Performance, Sepharose XL and Sepharose Fast Flow for media selection and pH scouting.

Sepharose Fast Flow
(Q, SP, DEAE, CM, ANX)



Try weak ion exchangers such as DEAE, CM or ANX if the selectivity of Q or S is unsatisfactory.

Capto
(Q, S, MMC)



Use high bed heights for increased productivity. Use MMC for high salt feed.

Sepharose XL
(Q or SP)



Use Sepharose Q XL virus licensed as an alternative to cesium chloride gradients for purification of viruses, including adenovirus, or viral vectors.

Sepharose Big Beads
(Q or SP)

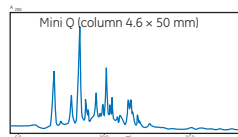


Use with step elution.

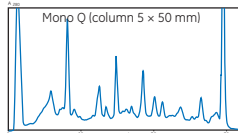
STREAMLINE™
(Q XL, SP XL, SP, DEAE, HST)



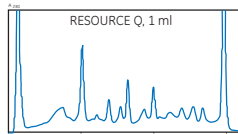
Use STREAMLINE for direct capture from unclarified feed-stock. Use HST, a salt tolerant adsorbent, to minimize dilution and reduce process time.



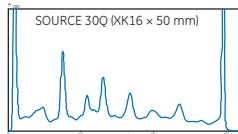
Sample: Pancreatin Gradient elution



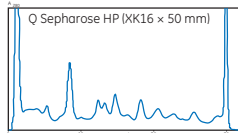
Sample: Pancreatin Gradient elution



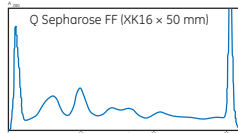
Sample: Pancreatin Gradient elution



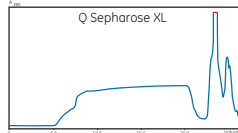
Sample: Pancreatin Gradient elution



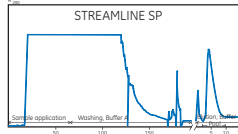
Sample: Pancreatin Gradient elution



Sample: Pancreatin Gradient elution



Sample: Recombinant α -amylase
Pilot scale: Gradient elution begins after 20 l



Sample: Recombinant antigen binding fragment
Pilot scale: Step elution

Resolution