



## Nucleus Protocols

# Exchange Buffers and Concentrate by Spin Filtration

- Dilute your sample ~10x with your new buffer (e.g., add 54 mL Protein Buffer to 6 mL eluted protein).
- Load your sample onto a centrifugal filter column.
- Centrifuge samples at either 4000 rcf (4 mL or 15 mL filters) or 14 000 rcf (0.5 mL filters) at 4°C until you've reached your target volume. Check on your sample volume in the first 10 min, then again as needed.
- Continue to dilute and spin your sample, noting your sample volume and dilution volume at each step, until you've reached your target dilution factor (we recommend  $\geq 300x$ ).
- Add an equal volume of Protein Buffer (60% glycerol) to your sample to bring the final glycerol concentration to 30%. Freeze samples at  $-80^{\circ}\text{C}$  for storage.
- Store columns for later use.
  - Wash columns by loading with ddH<sub>2</sub>O and spinning.
  - Load columns with EtOH 20% (v/v) and store at room temp.