

Sephadex™ G-25 media and prepacked formats

Gel filtration has successfully been employed for size based separations of macromolecules since the late 1950s by the introduction of the first commercially available medium (resin), Sephadex, in 1959. Gel filtration, or size exclusion chromatography, may be performed mainly in two principally different modes, depending on the size differences of the solutes to be separated: group separation and fractionation.

The main usage for Sephadex product family is group separation, such as desalting, buffer exchange, and sample cleanup. Sephadex media are members of the BioProcess™ media family and carry comprehensive technical and regulatory support

Sephadex G-25 media and prepacked columns offer:

- Quick desalting, removal of contaminants, and buffer exchange in one single step
- Excellent recovery and minimum sample dilution
- Well-established media for industrial applications

Separation principle

"Group separation" separates molecules according to their relative sizes: small molecules such as salts, free labels, and other impurities are efficiently separated from high molecular weight molecules of interest. In Sephadex, the degree of cross-linking of the dextran determines the extent to which macromolecules can permeate the beads. Large molecules are excluded while smaller sized molecules enter the beads to varying extents according to their different sizes. Large molecules thus leave the column first, followed by smaller molecules.



Fig 1. Prepacked formats such as HiTrap[™], HiPrep[™], PD-10 Desalting Columns, PD MiniTrap[™] G-25, PD MidiTrap[™] G-25, PD SpinTrap[™] G-25, and PD MultiTrap[™] G-25 desalting columns or 96-well plate give fast and convenient group separations between high and low molecular weight substances.

Sephadex G-25 has a fractionation range for globular proteins of molecular weights ($M_{
m p}$) 1000 to 5000, with an exclusion limit of approximately $M_{
m p}$ 5000. Proteins and peptides larger than $M_{
m p}$ 5000 are therefore easily separated from molecules with $M_{
m p}$ of less than 1000. Separations are conveniently performed using different prepacked column and 96-well plate formats, either manually with gravity flow The principles of gel filtration and the structures of Sephadex and other gel filtration media are explained in more detail in the handbook Gel filtration, principles and methods, see Related literature under "Ordering information".

Chromatography media characteristics

Sephadex media are prepared by cross-linking dextran with epichlorohydrin. The media are supplied dry, except when prepacked in columns and plate format. The media swells in aqueous solutions (Table 1). Sephadex G-25 is one of six different Sephadex G-types. The different types of Sephadex media differ in their degree of cross-linking and hence in their degree of swelling and molecular fractionation range.

The effect of particle size

Sephadex G-25 is available in four different particle size grades (Coarse, Medium, Fine, and Superfine). These four grades form the basis of a range of group separation products. Coarse and Medium grades are preferred for group separations at process scale where high flow rates and low operating pressures are required. The Fine and Superfine grades are for routine laboratory work including preparative separations. The smaller particle sizes of the Fine and Superfine grades give shorter diffusion distances and allow for highly efficient separations at high flow rates. Examples of different uses are shown under Applications and illustrate several of these properties.

The particle sizes of Sephadex media are usually reported as dry diameter in µm. However, as the media are swollen before use, it is the wet bead diameter that is of practical importance when choosing the correct type of equipment to use. The dry particle diameters as well as the hydrated-to-dry diameter ratios for Sephadex-25 media are shown in Table 1. Note that the degree of swelling in organic solvents is not the same as in aqueous solutions and must be determined by experiment for each solvent to be used. For routine use in organic solvents, lipophilic Sephadex LH-20 is recommended.

Table 1. Particle sizes for Sephadex G-25 media

Sephadex G-25	Dry particle diameter (µm)	Diameter ratio hydrated / dry
Superfine	20 to 50	~1.7
Fine	20 to 80	~1.7
Medium	50 to 150	~1.7
Coarse	100 to 300	~1.7

Stability

The mechanical strength and pH stability of Sephadex gel filtration media depend on the degree of cross-linking.

Sephadex G-25 is one of the more rigid members of the family and has a working pH range of 2 to 13. In trials, Sephadex G-25 has been exposed to 0.1 M HCl for 1 to 2 hours and to 0.02 M HCl for 6 months without impact on its chromatographic properties. Leakage studies have shown that Sephadex G-25 also withstands long term exposure to NaOH, and can be stored in 0.01 M NaOH without alterations in its performance.

For storage, 20% ethanol can also be used. Cleaning and sanitization by 60 to 90 min exposure to 0.2 M NaOH, followed by flushing with water or buffer, is recommended and can be repeated for hundreds of cleaning cycles.

Sephadex media can be autoclaved in their wet form (pH 7.0)

Batch-to-batch reproducibility

for 30 min at 120°C.

Consistent quality and performance from batch-to-batch is important for all separation media, but is most significant when the media are routinely used in industrial processes. Quality control data gathered for more than 50 years show outstanding batch-to-batch consistency of the Sephadex media.

Working with Sephadex G-25 Column dimensions

In gel filtration, column length is important for resolution and column diameter determines loading capacity. For group separations at high sample loadings, short (5 to 30 cm) and wide columns give excellent results at sample loadings of about 30% of the column volume.

Column packing

As Sephadex is supplied as a dry powder, it must be swollen in buffer before packing in the column. Laboratory columns can be packed by pouring the swollen G-25 slurry into the column and using a pump to pack the medium bed. For packing flow rates, see the instructions for detailed information. HiTrap and HiPrep, PD-10 Desalting Columns, PD MiniTrap G-25, PD MidiTrap G-25, PD SpinTrap G-25, and PD MultiTrap G-25 prepacked formats give convenience and reproducible results. Packing process scale columns requires modified techniques but gives excellent results and consistent performance.

Parameters affecting desalting

The most critical parameter affecting resolution in desalting applications is the sample-to-medium volume ratio. To minimize dilution and still retain good separation, sample volumes of up to approximately 30% of the total bed volume are recommended. Figure 2 illustrates how the resolution of a group separation run on HiPrep 26/10 Desalting is affected by sample volume. For base line separation, resolution should be at least 1.5. As flow rate has a minor impact on resolution, desalting can be performed at high flow rates (Fig 3).

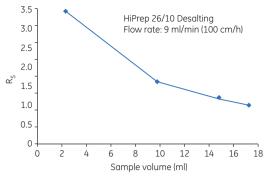


Fig 2. Influence of sample volume on the resolution (R₂).

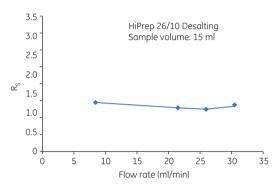


Fig 3. Influence of flow rate on the resolution (R_s) .

Cleaning packed columns

For columns that have been in use for some time, it can be necessary to remove precipitated proteins or other contaminants. Columns packed with Sephadex G-25 may be cleaned with 2 column volumes of 0.2 M NaOH or a nonionic detergent solution (60 to 90 min exposure). The frequency of cleaning will depend on the nature of the sample material and should be worked out on a case-by-case basis.

Scale-up

Scaling up a separation based on Sephadex G-25 from small-scale to large-scale routine production is a straightforward procedure. Well known examples of successful commercial applications include buffer exchange in processes for removing endotoxins from albumin (see "Applications") and as a preparative step during the production of vaccines.

Applications

As noted earlier, group separations such as buffer exchange and desalting are the most widely used applications for Sephadex G-25. Buffer exchange is often necessary when the buffer composition of the sample needs to be changed between chromatography steps. Desalting of a sample is often performed before ion exchange, after hydrophobic interaction or affinity chromatography, and prior to analysis. Cleanup of small contaminants are, for example, needed after labeling to remove free labeling molecules prior to analysis. At small scales, gel filtration with Sephadex G-25 has replaced much slower and more cumbersome dialysis procedures.

Figure 4 shows process data from a buffer exchange step using Sephadex G-25 Coarse in a process for removal of endotoxins and/or ethanol from albumin. The high throughput of Sephadex G-25 Coarse grade resulted in a rapid and efficient separation, despite the high concentration of protein in the feedstock.

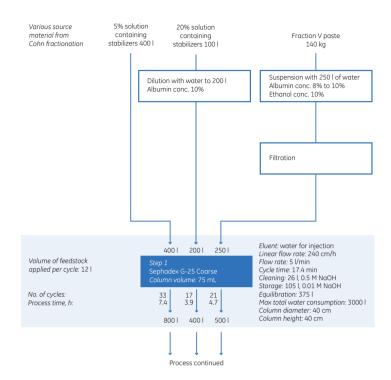


Fig 4. Process data from a buffer exchange step using Sephadex G-25 Coarse in a production scale process to remove endotoxins and/or ethanol from albumin.

Other desalting applications at larger scale can be exemplified by intermediate group separation of allergen extract from low molecular weight impurities using a BPG 450 column packed with Sephadex G-25 Superfine. The bed height was 15 cm, giving a bed volume of 25 l. The sample volume was up to 6 l (24% load) and the flow velocity 1.1 cm/min, giving a cycle time of 17 min. The dilution factor was 1.4.

A classic example of desalting is given by the de-ethanolization of human serum albumin (1). In this example, 12 l of a 9% protein solution was purified at a cycle time of 17.4 min using a 60×40 cm i.d. column packed with Sephadex G-25 Coarse, corresponding to a productivity of 50 g/h and l.

Horton describes the application of a 2500 l column of Sephadex G-25 Coarse for desalting a crude enzyme preparation (2). The column used was a stainless steel GF 18–10 (100 × 180 cm i.d.) column, the sample volume was 875 l (i.e., 35% of the column volume) and the flow rate was 62.5 l/min, giving a cycle time of 1 h. Horton also provides a comparison between the productivity of Sephadex G-25 Coarse and Sephadex G-25 Medium (Table 2). In conclusion, Sephadex G-25 Coarse provides the highest productivity of these two media.

Gel filtration can offer a very robust purification step as noted for the use of Sephadex G-25 Coarse for initial buffer exchange of raw plasma in a large scale albumin fractionation plant (3). The column, containing 75 l of medium, was used for more than six years and processed 70 000 l of material.

Table 2. Comparison between the productivity of Sephadex G-25 Coarse with Medium for desalting of albumin from ammonium sulfate

Parameter	Sephadex G-25 Coarse	Sephadex G-25 Medium
Sample volume (I)	31	37
Relative sample volume (%)	25	30
Sample capacity (I/h)	70	18
Salt concentration in product (%)	0.4	0.04
Salt removed (%)	98.0	99.9
Albumin concentration in product (%)	4.0	5.0
Dilution factor of albumin	1.5	1.2
Albumin processed (g/h)	4125	1170
Productivity (g/h, l)	33	9
Cost per kg of albumin (USD)	0.64	2.20

Note: Albumin concentration in feed is 6%, salt concentration in feed is 24%. Column: 100×40 cm i.d., giving a bed volume of 125 l.

Prepacked columns with Sephadex G-25

Prepacked column and 96-well plate formats can be used when removal of low molecular weight components is needed. Examples of applications are:

- Preparation of samples prior to or after ion exchange, affinity, or hydrophobic interaction chromatography
- Removal of free low molecular weight labels or contaminants from proteins/peptides
- Removal of substrates, inhibitors, or co-factors from enzymes
- Preparation of samples for concentration, freeze-drying, or storage
- Termination of a reaction between a macromolecule and a low molecular weight reagent

HiTrap Desalting

HiTrap Desalting column is packed with Sephadex G-25 Superfine. The column is made of biocompatible polypropylene with polyethylene frits. The column is delivered with a stopper on the inlet and a twist-off end on the outlet. The column cannot be opened or refilled. Operation is easy, using a syringe, a peristaltic pump, or a liquid chromatography system. Larger sample volumes or, if required, better resolution can easily be achieved by connecting up to five columns in series.

Table 3 shows the main characteristics of HiTrap Desalting.

Desalting in a fraction of a minute

Because of the rigidity of the Sephadex G-25 matrix and an optimized column packing method, HiTrap Desalting column has a very low back pressure (0.25 bar [0.025 MPa, 3.7 psi] at 10 ml/min). The low back pressure makes it possible to run separations in a fraction of a minute using a syringe or a pump. The separation of bovine serum albumin (BSA) and NaCl is shown in Figure 4. The whole separation took only 45 s and the protein was eluted in less than 30 s.

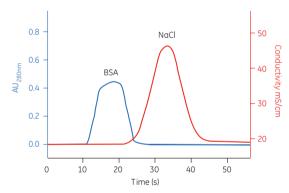
Column: HiTrap Desalting

Sample: 1.4 ml BSA (2 mg/ml) in 50 mM sodium phosphate,

0.5 M sodium chloride, pH 7.0

Buffer: 50 mM sodium phosphate, 0.15 M sodium chloride, pH 7.0

Flow rate: 10 ml/min



 $\textbf{Fig 5.} \ \textbf{Highly efficient desalting in half a minute using HiTrap Desalting}.$

Scaling up by connecting HiTrap Desalting columns in series

Larger sample volumes or, if required, better resolution can easily be achieved by connecting up to five columns in series. Figures 6 A to C show the results obtained when one, three, and five HiTrap Desalting columns were connected in series. The sample volumes were 1.4, 4.3, and 7.1 ml, respectively.

Column: HiTrap Desalting, 1×5 ml, 3×5 ml, 5×5 ml Sample: 2 mg/ml BSA in 50 mM sodium phosphate,

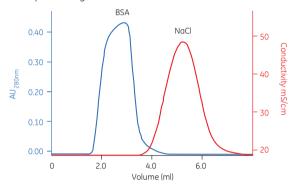
0.5 M sodium chloride, pH 7.0

Sample volume: 28% of column volume (1.4, 4.3 and 7.1 ml, respectively)

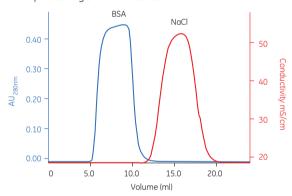
Buffer: 50 mM sodium phosphate, 0.15 M sodium chloride, pH 7.0

Flow rate: 5 ml/min

(A) HiTrap Desalting 1×5 ml



(B) HiTrap Desalting 3×5 ml in series



(C) HiTrap Desalting 5×5 ml in series

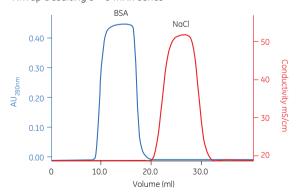


Fig 6. Separation of BSA from NaCl on HiTrap columns connected in series.

Online buffer exchange

HiTrap Desalting columns can easily be attached to other columns for online buffer exchange prior to or after a chromatographic step. An example is shown in Figure 7, where a HiTrap Desalting column is connected serially after a HiTrap Protein G HP column in order to adjust the pH of the eluted material. The HiTrap Desalting column was equilibrated with 20 mM sodium phosphate, pH 7.0.

A mouse monoclonal antibody (IgG1) from a serum-free cell culture supernatant was purified using a HiTrap Protein G HP 1 ml column. After loading of 5 ml sample and washing with 5 ml binding buffer (20 mM sodium phosphate, pH 7.0), bound material was eluted with 5 ml of 0.1 M glycine, pH 2.7. The first 1.3 ml eluted from the HiTrap Protein G HP column was discarded before the column was connected to the inlet of the HiTrap Desalting column. The eluate from the columns was monitored for UV absorbance and pH was measured in the collected 0.5 ml fractions.

Figure 7A shows the original separation without the extra desalting step. As shown in Figure 7B, the obtained yield was 18% higher when the HiTrap Desalting column was used for online desalting.

Columns: (A) HiTrap Protein G HP, 1 ml

(B) HiTrap Desalting, 5 ml

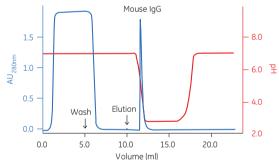
Sample: Eluted mouse monoclonal IgG from HiTrap Protein G HP

Binding buffer: 20 mM sodium phosphate, pH 7.0

Elution buffer: 0.1 M glycine, pH 2.7

Flow rate: 2 ml/min

(A) Original separation without the desalting step



(B) Separation of mouse IgG including a desalting step

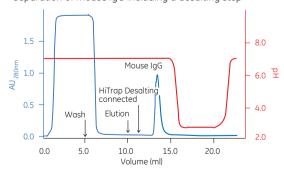


Fig 7. Online buffer exchange with HiTrap Desalting improves yield at a mouse IgG purification. (A) Original separation without the desalting step. (B) Separation including a desalting step.

HiPrep 26/10 Desalting

HiPrep 26/10 Desalting is packed with Sephadex G-25 Fine. The column has an internal diameter of 2.6 cm and a bed height of 10 cm. The medium bed volume is approximately 53 ml.

The HiPrep column is made of biocompatible polypropylene with nylon frits. The supplied set of connectors makes it easy to connect the column to different chromatography systems. The column is not designed to be opened or repacked. The characteristics of HiPrep 26/10 Desalting column are summarized in Table 3.

Table 3. Characteristics of HiTrap Desalting and HiPrep 26/10 Desalting

	HiTrap Desalting	HiPrep 26/10 Desalting
Matrix	Sephadex G-25 Superfine, cross- linked dextran	Sephadex G-25 Fine, cross-linked dextran
Bed volume	5 ml	53 ml
Bed dimension	1.6 × 2.5 cm	2.6 × 10 cm
Void volume	1.5 ml	15 ml
Recommended sample volume	0.1 to 1.5 ml	≤ 15 ml
Sample dilution, syringe operation	1.3 to 4.0 fold	1.2 to 3 fold
Exclusion limit, globular proteins	M _r 5000	M _r 5000
Average particle size	90 µm	90 μm
Maximum flow rate*	15 ml/min	40 ml/min
Recommended flow rate*	1 to 10 ml/min	9 to 31 ml/min
Column hardware pressure limit	5 bar (0.5 MPa, 70 psi)	5 bar (0.5 MPa, 70 psi)
Chemical stability	All commonly used buffers	All commonly used buffers
pH stability, cleaning and working [†]	2 to 13	2 to 13
Avoid	Oxidizing agents	Oxidizing agents
Storage	8°C to 30°C in 20% ethanol	8°C to 30°C in 20% ethanol

^{*} Room temperature, aqueous buffers.

Buffer exchange of mouse plasma

Buffer was exchanged on mouse plasma within 1.5 min on HiPrep 26/10 Desalting column. A 10 ml sample was applied at a flow rate of 22 ml/min (250 cm/h). Figure 8 shows the chromatogram of the separation. The protein was eluted in a volume of 19 ml.

Column: HiPrep 26/10 Desalting

Sample: Mouse plasma, centrifuged at $10\,000 \times g$ for $10\,min$

Sample volume: 10 ml

Buffer: 25 mM sodium acetate, pH 7.0

Flow rate: 22 ml/min (250 cm/h)

Instrument: ÄKTAexplorer 100 (1 mm i.d. tubing installed)

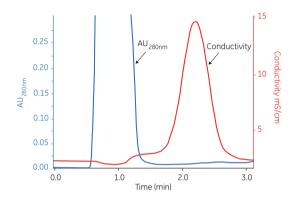


Fig 8. Buffer exchange of mouse plasma on HiPrep 26/10 Desalting.

Reproducible removal of N-Hydroxysuccinimide from BSA

Figure 9 shows the removal of N-Hydroxysuccinimide (M_r 115) from BSA (M_r 67 000) in three parallel runs. Highly efficient and reproducible desalting was achieved in all three runs.

Column: HiPrep 26/10 Desalting

Sample: 2 mg/ml BSA, 0.07 mg/ml N-Hydroxysuccinimide (NHS)

in 50 mM sodium phosphate, 0.15 M NaCl, pH 7.0.

Filtered through a 0.45 µm filter

Sample volume: 13 ml

Buffer: 50 mM sodium phosphate, 0.15 M NaCl, pH 7.0

Flow rate: 31 ml/min (350 cm/h)

Instrumentation: ÄKTAexplorer 100 (1 mm i.d. tubing installed)

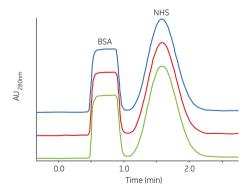


Fig 9. Reproducible removal of N-Hydroxysuccinimide from BSA.

[†] Cleaning refers to the pH interval for regeneration. Working refers to the pH interval where the medium is stable over a long period of time without adverse effects on its performance.

Scaling up sample volumes

Samples of 15 ml (30% of the total bed volume) or less can be applied to a single column, and by coupling of up to four columns in series (Fig 10), a maximum sample volume of 60 ml can be run. Even with four columns in series, high flow rates can be maintained without back pressure problems, resulting in fast separations. In fact, up to 60 ml of sample can be desalted or buffer exchanged in 20 to 30 min.



Fig 10. Four HiPrep 26/10 Desalting columns connected in series.

Table 4 shows run data and results for HiPrep 26/10 Desalting columns in series. Samples consisted of either 30 or 60 ml of a fungal culture supernatant containing a secreted recombinant protein. Samples were run on two (30 ml sample) or four (60 ml sample) columns connected in series.

Table 4. Run data for buffer exchange of 30 ml and 60 ml samples on HiPrep 26/10 Desalting columns connected in series

Columns

2 × HiPrep 26/10 Desalting in series

Columns	$(VT = 106 \text{ ml})$ for 30 ml sample or $4 \times \text{HiPrep } 26/10$
	Desalting in series (VT = 212 ml) for 60 ml sample
Sample	30 ml or 60 ml <i>Pichia pastoris</i> culture supernatant containing a secreted recombinant protein
Sample preparation	Filter through 0.45 µm filter
Sample loop	Superloop™ 150 ml
Sample elution	
30 ml loading	35 ml (dilution factor 1.2)
60 ml loading	70 ml (dilution factor 1.2)
Buffer	0.1 M Tris, 0.15 M NaCl, 0.05% Tween™ 20, pH 7.6
Flow rates (30 ml)	
sample loading elution	12 ml/min 17 ml/min
Flow rates (60 ml)	
sample loading	8 ml/min
elution	11 ml/min

Disposable PD-10 Desalting Columns, PD MidiTrap G-25, PD MiniTrap G-25, PD SpinTrap G-25, and PD MultiTrap G-25

Disposable PD-10 Desalting Columns, PD MidiTrap G-25. PD MiniTrap G-25, PD SpinTrap G-25, and PD MultiTrap G-25 are prepacked, single-use column and 96-well filter plate formats packed with Sephadex G-25 Medium or Sephadex G-25 Superfine for buffer exchange, desalting, and cleanup of biological samples, for example proteins and carbohydrates (Figure 11). These columns and 96-well filter plates provide convenient small-scale preparation of protein samples prior to downstream analytical techniques such as gel electrophoresis, liquid chromatography, and liquid chromatography-mass spectrometry (LC-MS). This range of columns and plates cover sample volumes up to 2.5 ml (Table 5). The columns and 96-well plates are manufactured from biocompatible polypropylene. Special column and plate frits protect the medium from running dry during use. Recommended volumes for sample loading and elution with subsequent dilution factors are listed in Table 6.



Fig 11. Simple desalting & buffer exchange with Disposable PD-10 Desalting Columns, PD MidiTrap G-25 and PD MiniTrap G-25 gravity flow columns, PD SpinTrap G-25 microspin column, and PD MultiTrap G-25 96-well filter plate.

	PD MultiTrap G-25	PD SpinTrap G-25	PD MiniTrap G-25	PD MidiTrap G-25	PD-10 Desalting Columns
Volume of prepacked medium	500 µl/well	~600 µl/column	2.1 ml	2.1 ml	8.3 ml
Packed bed dimensions	N/A	N/A	0.97 × 2.8 cm	1.3 × 2.6 cm	1.45 × 5.0 cm
Well/column volume	800 µl	1000 μΙ	5 ml	8.5 ml	13. 5 ml
Void volume	~ 150 µl	~ 150 µl	~ 0.5 ml	~ 1.0 ml	2.5 ml
Maximum sample volume	130 µl	180 μΙ	0.5 ml	1.0 ml	2.5 ml
Volume of eluted sample (gravity)	-	_	1.0 ml	1.5 ml	3.5 ml
Volume of eluted sample (spin)*	130 µl	180 μΙ	0.5 ml	1.0 ml	2.5 ml
Recovery [†]	70% to 90%	70% to 90%	70% to 95%	70% to 95%	70% to > 95%
Desalting capacity	> 85%	> 85%	> 90%	> 90%	> 90%
Plate/column material	polypropylene and polyethylene				
Storage solution	20% ethanol	0.15% Kathon™	0.15% Kathon	0.15% Kathon	0.15% Kathon
Storage temperature	4°C to 30°C				

^{*} Applied volume = eluted volume

Table 6. Recommended sample and elution volumes on desalting/buffer exchange columns and plate

Column	Loaded volume (ml)	Eluted volume (ml)	Dilution factor	Operation
HiPrep 26/10 Desalting	10 15 (max)	10 to 15 15 to 20	1 to 1.5 1 to 1.3	pump pump
2 × HiPrep 26/10 Desalting	30 (max)	30 to 40	1 to 1.3	pump
3 × HiPrep 26/10 Desalting	45 (max)	45 to 55	1 to 1.2	pump
4 × HiPrep 26/10 Desalting	60 (max)	60 to 70	1 to 1.2	pump
HiTrap Desalting	0.25 0.5 1.0 1.5 (max)	1.0 1.0 2.0 2.0	4 3 2 1.3	syringe/pump syringe/pump syringe/pump syringe/pump
2 × HiTrap Desalting	3.0 (max)	4 to 5	1.3 to 1.7	syringe/pump
3 × HiTrap Desalting	4.5 (max)	6 to 7	1.3 to 1.7	syringe/pump
PD-10	1.5 2.0 2.5 (max)	3.5 3.5 3.5	2.3 1.8 1.4	gravity gravity gravity
PD MiniTrap G-25	0.1 to 0.5 0.2 to 0.5	1.0 Up to 0.5	2	gravity spin
PD MidiTrap G-25	0.5 to 1.0 0.75 to 1.0	1.5 Up to 1.0	2	gravity Spin
PD SpinTrap G-25	0.1 to 0.18	0.14 to 0.18	0 to 1.4	Centrifugation
PD MultiTrap G-25	0.07 to 0.13	0.1 to 0.13	0 to 1.4	Vacuum/centrifugation

Small-scale cleanup

PD SpinTrap G-25 and PD MultiTrap G-25 are designed for small-scale cleanup and are valuable tools for screening purposes and high-throughput applications. SpinTrap columns only require a standard microcentrifuge (Fig 12A). MultiTrap 96-well filter plates allow cleanup by centrifugation, either manually or automated with robotics (Fig 12B).

Typical desalting capacity is above 85% with recoveries between 70% and 90% (biomolecule dependent). In a run of 96 parallel wells with BSA in 1 M NaCl, the salt removal capacity varied by 1% (Fig 13) and the recovery of total loaded material varied by 3%.

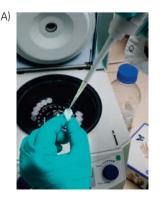




Fig 12. (A) PD SpinTrap G-25 sample preparation. (B) PD MultiTrap G-25 sample automated preparation in a robotic system.

[†] Biomolecule dependent

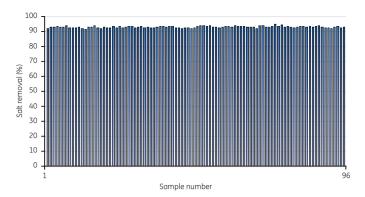


Fig 13. Removal of NaCl from BSA on a PD MultiTrap G-25 96-well plate showed highly reproducible results. The average desalting capacity was 93% and the well-to-well variation was 1% (relative standard deviation).

Columns for both gravity and spin protocols

PD-10 Desalting Columns, PD MidiTrap G-25, and PD MiniTrap G-25 provide two possible application protocols, gravity and spin. The typical desalting capacity, for both protocols, is above 90% with recoveries between 70% and 95% (biomolecule dependent).

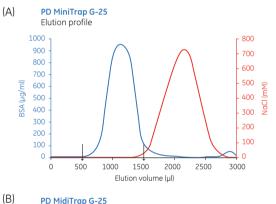
Gravity protocol

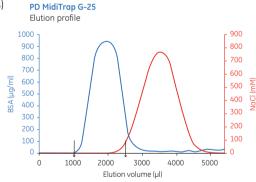
By using the gravity protocol, a simple cleanup of the sample is performed on the lab bench with one or several columns in parallel without any need for a purification system. To simplify the use of PD-10 Desalting Columns with the gravity protocol, the LabMate PD-10 Buffer Reservoir may be used (see "Ordering information"). Using buffer reservoir, wash and equilibration buffers can be applied in one step. Elution profiles for BSA using the gravity protocol are shown in Figure 14.

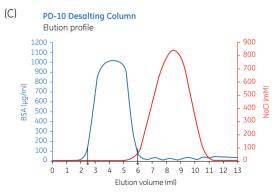
Trap products: (A) PD MiniTrap G-25, (B) PD MidiTrap G-25, and (C) PD-10 Desalting Columns

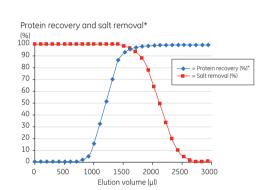
Sample: 1000 μ g/ml BSA in 1 M NaCl Sample volumes: (A) 500 μ l, (B) 1000 μ l, and (C) 2.5 ml

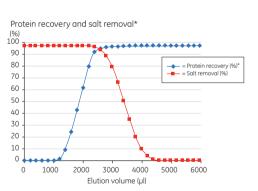
Equilibration buffer: Milli-Q™ water











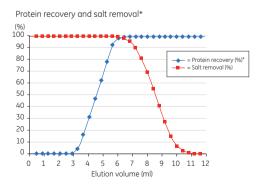


Fig 14. Removal of NaCl from BSA using the gravity protocol. The protein recovery (between arrows) was 95% for PD MiniTrap G-25 and PD MidiTrap G-25 (A and B) and > 95% for PD-10 Desalting Columns (C). The desalting capacity was > 98% for PD MidiTrap G-25 (B) and > 99% for PD MiniTrap G-25 and PD-10 Desalting Columns (A and C). The images to the right illustrate the protein recovery and salt removal versus the total elution volume for the used column.

normalized against the total amount of loaded sample.

Spin protocol

By using the spin protocol, the samples are run in parallel in a standard centrifuge. The spin protocol gives minimal dilution of the eluted sample. Four adapters are included in each product package to enable easy use of a standard centrifuge (Fig 15). To facilitate increased throughput, a package of 10 adapters can be ordered separately.





Fig 15. Spin adapters are used together with PD-10 Desalting Columns, PD MidiTrap G-25, and PD MiniTrap G-25 to enable use in a standard centrifuge.

Efficient cleanup of carbohydrates before enzymatic cleavage

The PD products prepacked with Sephadex G-25 Medium are excellent for desalting biomolecules other than proteins, for example carbohydrates. To demonstrate this, bovine intestinal 3H-labeled heparan sulfate was eluted from a DEAE column using a high salt concentration. The eluate was run on a PD MidiTrap G-25 column before enzymatic cleavage. Due to high peak resolution, the sample was collected with high recovery in a very low concentration of salt. The elution profile is shown in Figure 16.

Trap product: PD MidiTrap G-25

Sample: 18 600 cpm bovine intestinal ³H-labeled heparan

sulfate ([3H]HS) in 1.5 M NaCl

Sample volume: 0.5 ml Equilibration buffer: Distilled water

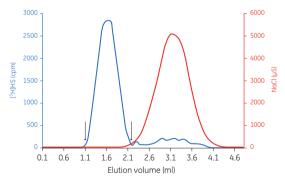


Fig 16. Removal of NaCl from [3H]HS on a PD MidiTrap G-25 column. The fractions were analyzed with regards to conductivity and radioactive content. Fractions 11 through 21 (between arrows) contained 87% of the total recovery of [3H]HS. The recovery volume was 1.1 ml with a NaCl concentration of 20 mM, corresponding to > 98% salt removal.

Conclusions

Sephadex G-25 media are excellent for desalting of both protein and DNA preparations. Sephadex G-25 Coarse will generally be the gel filtration medium of choice for industrial scale. It is especially suitable for rapid processing of large volumes of feed where the requirement for productivity is high. In cases where sample dilution needs to be minimized, the higher resolution of a smaller particle size, for example Sephadex G-25 Medium, might be required.

For laboratory scale desalting where less effort needs to be spent on optimization, it is common to use more conservative sample volumes (i.e., a load below 20% of the column volume). For these applications, Sephadex G-25 Medium or Sephadex G-25 Fine can be recommended to minimize the dilution factor. For micropreparative work using very small sample volumes, Sephadex G-25 Superfine will be the medium of choice.

For Sephadex G-25 media, several different prepacked column and plate formats are available, and for which the choice depends on sample volume and existing laboratory equipment.

References

- Hagel L. and Janson J.-C., Size-exclusion chromatography, in Chromatography, 5th edition (E. Heftmann, ed.), Elsevier, Amsterdam, pp. A267–AA307 (1992)
- 2. Horton T., Large-scale gel filtration for purification of natural products, *Amer Lab*, May (1972)
- 3. Marrs S. B., Large scale albumin fractionation by chromatography, *Biotech Blood Prot* 227:169–173 (1993)

Ordering information

Prepacked columns	Quantity	Code number
HiTrap Desalting	1 × 5 ml 5 × 5 ml 100 × 5 ml	29-0486-84 17-1408-01 11-0003-29
HiPrep 26/10 Desalting	1 × 53 ml 4 × 53 ml	17-5087-01 17-5087-02
Bulk media		
Sephadex G-25 Coarse	100 g 500 g 5 kg 40 kg	17-0034-01 17-0034-02 17-0034-03 17-0034-07
Sephadex G-25 Medium	100 g 500 g 5 kg 40 kg	17-0033-01 17-0033-02 17-0033-03 17-0033-07
Sephadex G-25 Fine	100 g 500 g 5 kg 40 kg	17-0032-01 17-0032-02 17-0032-03 17-0032-07
Sephadex G-25 Superfine	100 g 500 g 5 kg	17-0031-01 17-0031-02 17-0031-03
Prepacked disposable products		
PD-10 Desalting Columns	30 columns	17-0851-01
PD SpinTrap G-25	50 columns	28-9180-04
PD MultiTrap G-25	4 × 96-well plates	28-9180-06
PD MiniTrap G-25	50 columns	28-9180-07
PD MidiTrap G-25	50 columns	28-9180-08
MiniSpin Adapter	10	28-9232-43
MidiSpin Adapter	10	28-9232-44
PD-10 Spin Adapter	10	28-9232-45
Collection plate 500 µl V-bottom (for collection of fractions from MultiTrap)	5 × 96-well plates	28-4039-43
LabMate PD-10 Buffer Reservoir	10	18-3216-03

Accessories	Quantity	Code number
1/16" male/Luer female*	2	18-1112-51
Tubing connector flangeless/M6 female	2	18-1003-68
Tubing connector flangeless/M6 male	2	18-1017-98
Union 1/16" female/M6 male	6	18-1112-57
Union M6 female/1/16" male	5	18-3858-01
Union luerlock female/M6 female	2	18-1027-12
HiTrap/HiPrep, 1/16" male connector for ÄKTAdesign	8	28-4010-81
Stop plug female, 1/16"†	5	11-0004-64
Fingertight stop plug, 1/16"‡	5	11-0003-55

^{*} One connector included in each HiTrap package.

Related literature

Handbook. Gel Filtration, Principles and Methods	18-1022-18
Selection Guide: Gel Filtration Column and Media	18-1124-19
Selection guide: Sample preparation for analysis of proteins, peptides and carbohydratesDesalting, Buffer Exchange, Cleanup, Concentration	18-1128-62

 $^{^\}dagger$ Two, five, or seven stop plugs female included in HiTrap packages depending on product.

 $^{^{\}dagger}$ One fingertight stop plug is connected to the top of each HiTrap column at delivery.

For local office contact information, visit www.gelifesciences.com/contact

www.gelifesciences.com/protein-purification

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