

MabSelect SuRe™ LX

MabSelect SuRe LX is an affinity chromatography medium for capturing high titer monoclonal antibodies from large volumes of feed by packed bed chromatography.

- High dynamic binding capacity (DBC) for high titer cultures reduces process time and amount of medium used.
- Alkali-tolerant protein A-derived ligand allows the use of 0.1–0.5 M sodium hydroxide for Cleaning-In-Place (CIP). Eliminates the need for expensive and corrosive CIP reagents.
- High-flow agarose matrix allows processing of large volumes of feed.
- Improved performance and reduction in overall costs.
- Simple scale-up to production-sized AxiChrom™ columns.



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1 Description

The protein A-derived MabSelect SuRe LX ligand is produced in *Escherichia coli*. Fermentation and subsequent purification are performed in the absence of animal products. The ligand has been specially engineered to create an affinity medium with enhanced alkali and protease stability. The specificity of binding to the Fc region of IgG is similar to that of conventional Protein A and provides excellent purification in one step. MabSelect SuRe LX has very high dynamic binding capacities at extended residence times, and is developed for high titer antibody processes, see Figure 1. Alkali tolerance, high capacity and low ligand leakage plus the rigid base matrix, make MabSelect SuRe LX ideal for the purification of monoclonal antibodies for clinical applications.

Figure 1 shows the relation between dynamic binding capacity and residence time for MabSelect SuRe LX.

Figure 2 shows stability in alkaline conditions of MabSelect SuRe LX compared with MabSelect SuRe in terms of dynamic binding capacity.

The characteristics of the medium are summarized in Table 1.

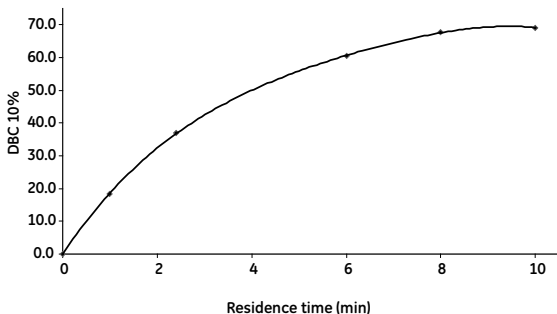


Fig 1. Relation between dynamic binding capacity and residence time for HiScreen MabSelect SuRe LX for polyclonal IgG.

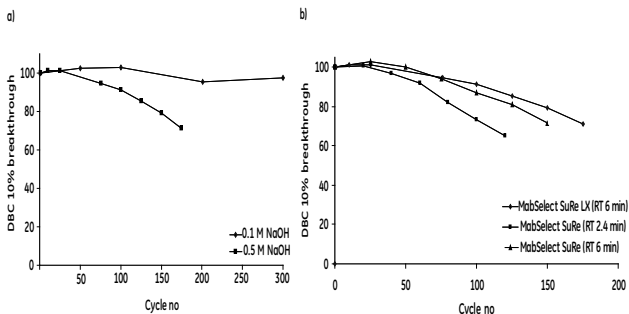


Fig 2. Cycling with buffers

a) Dynamic binding capacity (residence time 6 min) for MabSelect SuRe LX after CIP with 0.1 and 0.5 M NaOH for 0-300 cycles.

b) Dynamic binding capacity for MabSelect SuRe LX and MabSelect SuRe after CIP with 0.5 M NaOH for 0-175 cycles (residence time 2.4 and 6 min).

Each cycle in Figure 2 consisted of:

- 5 column volumes binding buffer pH 7.4
- 5 column volumes 0.1 M acetic acid pH 3.0
- 0.1 or 0.5 M NaOH, 15 minutes contact time
- 5 column volumes binding buffer pH 7.4

The dynamic binding capacity for polyclonal IgG (DBC 10% breakthrough) was measured regularly during the study.

Table 1. Characteristics of MabSelect SuRe LX

Composition	Rigid, highly cross-linked agarose
Average particle size (d_{50v}) ¹	85 μm
Ligand	MabSelect SuRe ligand (alkali-tolerant, protein A-derived (from <i>E. coli</i>))
Coupling chemistry	Epoxy
Dynamic binding capacity ²	Approx. 60 mg human IgG/ml medium
Chemical stability	Stable in all aqueous buffers commonly used in Protein A chromatography.
pH working range	3 to 12
Cleaning-in-place stability ³	0.1 to 0.5 M NaOH
Maximum operational velocity ⁴	500 cm/h
Temperature stability ⁵	2°C to 40°C
Delivery conditions	20% ethanol
Regulatory support	Regulatory support file is available. No material of animal origin is used in the manufacturing process

¹ d_{50v} is the medium particle size of the cumulative volume distribution.

² Determined at 10% breakthrough by frontal analysis at a mobile phase velocity of 100 cm/h in a column with a bed height of 20 cm, i.e. residence time is 6.0 min. Residence time is equal to bed height (cm) divided by nominal fluid velocity (cm/h) during sample loading. Nominal fluid velocity is equal to volumetric flow rate (ml/h) divided by column cross-sectional area (cm²).

³ See Figure 2 and section 7 Cleaning-In-Place (CIP).

⁴ In AxiChrom 300 column, bed height 20 cm, operating pressure < 2 bar, water at 20°C.

⁵ Recommended long-term storage conditions: +2°C to +8°C, 20% ethanol.

2 Process development

For initial studies on MabSelect SuRe LX, PreDicator™ plates is preferentially used. The PreDicator plates are 96-well plates pre-filled with chromatography media, which can be used for rapid screening of chromatographic conditions in small scale. For further optimization in small-scale columns, we recommend prepacked HiScreen™ columns or HiScale™ columns.

Choose a residence time (see Table 1, footnote 2, and Figure 1) that fulfills your demand on dynamic binding capacity and nominal fluid velocity according to Figure 4. Ancillary cycle operations including wash, elute and equilibration steps can be run at maximum operational velocities, see Table 1. Example of a pressure/flow curve in water is seen in Figure 3.

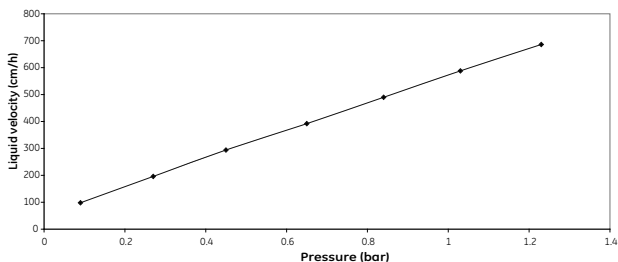


Fig 3. Example of a pressure/flow curve in water 20°C for 20 cm packed bed of MabSelect SuRe LX in AxiChrom 300, Packing Factor 1.15. The additional pressure from test system and tubing is subtracted.

Make sure the chosen bed height and velocity do not conflict with the large-scale pressure/flow limitations (Figure 4).

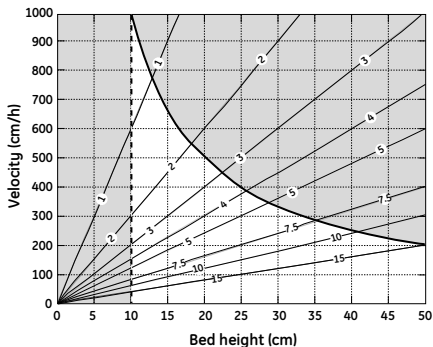


Fig 4. Operating window for MabSelect SuRe LX (white area). Choosing bed height and operating velocity in terms of residence time, pressure restrictions and large-scale column packing challenges.

Figure 4 shows the possible combinations of bed height and operational nominal fluid velocity for MabSelect SuRe LX. The figure also displays the residence time in the interval 1–15 minutes for any bed height and velocity. Included are also pressure drop limitations and packing limitations at large scale. The solid curved line shows the calculated large-scale column pressure restriction which is 2 bar according to specification (500 cm/h at 2 bar and 20 cm bed height). The dashed vertical line indicates that operating at below 10 cm bed height is not favorable. The reason for this is that large diameter columns have a very different aspect ratio, and that packing short wide beds is a greater challenge.

Figure 4 can be used as a guide when determining suitable bed height and operating velocity in terms of residence time and thus capacity and pressure drop.

3 Recommended screening conditions

Examples of suitable buffers:

- Buffer A: 20 mM sodium phosphate, 150 mM NaCl, pH 7.2
- Buffer B: 0.1 M sodium citrate, pH 3.0–3.6.

Experimental conditions:

- Equilibrate the column with 5 column volumes (CV) of buffer A.
- Apply a small sample of antibody at residence time ≥ 6 mins.
- Wash the column with 5 CV of buffer A.
- Elute the column with a 10 CV linear gradient from 0% to 100% buffer B.
- Collect fractions into titrating diluent (e.g. 1.0 M Tris-HCl, pH 8.0 so that the diluent volume equals 5% of the programmed fraction volume).
- Regenerate the column with 5–10 CV of 100% buffer B.
- Wash the column with 3 CV of buffer A.
- Perform CIP with 5 CV of 0.1–0.5 M NaOH.
- Re-equilibrate the column with buffer A.

To minimize the use of buffer, however, we recommend optimizing the washing procedure with respect to residence time, volumes, pH and conductivity.

When optimizing elution conditions, determine the highest pH that allows efficient desorption of antibody from the column. This will prevent denaturing sensitive antibodies due to exposure to low pH. Step-wise elution (Figure 5) is often preferred in large-scale applications since it allows the target monoclonal antibody to be eluted in a more concentrated form, thus decreasing buffer consumption and shortening cycle times. It might be necessary to decrease the flow rate due to the high concentrations of protein in the eluate.

Figure 5 shows an example of purification of a monoclonal antibody from a clarified mammalian cell culture on MabSelect SuRe LX. The load was 21 mg antibody/ml column volume (CV), and the yield was 94% of highly purified antibody. A HiScale 16/20 column with a CV of 20 ml and a bed height of 10 cm was used.

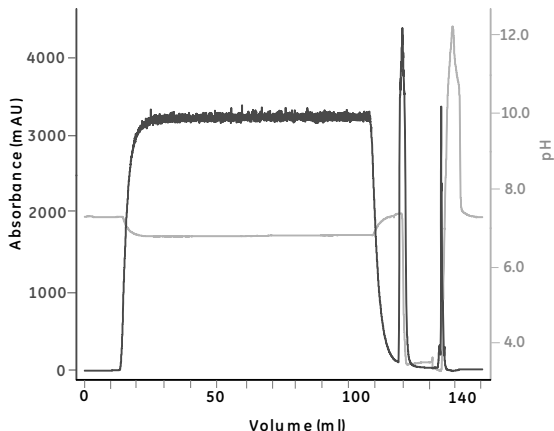


Fig 5. Purification of a monoclonal antibody from a mammalian cell culture on MabSelect SuRe LX.

The dynamic binding capacity for the target antibody should be determined by frontal analysis using real process feedstock. The dynamic binding capacity is a function of the sample residence time and should therefore be defined over a range of different sample residence times.

4 Removal of leached ligand from final product

The ligand leakage from MabSelect SuRe LX is generally low. For example, the eluate from the purification run shown in Figure 5 contained 11 ppm (ng ligand/mg antibody) of leached ligand. However, in many monoclonal antibody applications it is a requirement to eliminate leached ligand from the final product.

There are a number of chromatographic solutions, such as cation and anion exchange chromatography, or multimodal anion exchange chromatography, which can be used to remove leached ligand.

For more details about removal of leached ligand and antibody aggregates, see the application note *Two step purification of monoclonal IgG₁ from CHO cell culture supernatant* (28-9078-92). Methods used for removal of leached ligand from MabSelect SuRe is applicable also for removal of leached ligand from MabSelect SuRe LX.

5 Packing columns

Recommended columns

Table 2. Recommended columns for MabSelect SuRe LX

Column	Inner diameter (mm)	Bed volume ¹	Bed height (cm)
Lab scale			
HiScale 16/20	16	20–40 ml	max 20
HiScale 16/40	16	20–70 ml	max 35
HiScale 26/20	26	53–106 ml	max 20
HiScale 26/40	26	53–186 ml	max 35
HiScale 50/20	50	196–393 ml	max 20
HiScale 50/40	50	196–687 ml	max 35
Production scale			
AxiChrom™ ²	50–200	0.2–12.5 l	max 40
AxiChrom ²	300–1000	7–314 l	max 40
BPG™ ³	100–300	1–28 l	max 40
Chromaflo™ standard ^{4,5}	400–800	12–151 l	max 30 cm

¹ Bed volume range calculated from 10 cm bed height to maximum bed height.

² Intelligent packing method according to MabSelect SuRe can be used.

³ The pressure rating of BPG 450 is too low to use with MabSelect media.

⁴ Packing instructions for MabSelect in Chromaflo columns are described in Application Note 11-0007-52.

⁵ Larger pack stations might be required at larger diameters.

All large-scale columns can be supplied as variable bed height columns. Do not choose large diameter columns if the bed height is low.

For practical instructions in good packing techniques, see the CD-ROM *Column Packing - The Movie* (18-1165-33). For more details about packing HiScale columns, see instructions *HiScale™ columns (16, 26, 50) and accessories (28-9674-70)*. For information on packing of process scale columns, please contact your local GE Healthcare representative.

Packing HiScale columns

Packing preparations

Materials needed

MabSelect SuRe LX

HiScale column

HiScale packing tube (depending on bed height)

Plastic spoon or spatula

Glass filter G3

Vacuum suction equipment

Filter flask

Measuring cylinder

20% ethanol with 0.4 M NaCl

Equipment

ÄKTA™ systems, or a stand-alone pump such as Pump P-900, depending on the flow rate required, can be used for packing.

Equilibrate all materials to room temperature.

Definitions

The bed height of a gravity settled bed differs from the bed height of a bed settled at a given flow (consolidated). Therefore, the compression factor (CF) has to be separated from the packing factor (PF).

L_{settled}	Settled bed height/Gravity settled bed height. Bed height measured after settling by gravity.
L_{cons}	Consolidated bed height Bed height measured after settling the medium at a given flow velocity.
L_{packed}	Packed bed height
CF	Compression factor $CF = L_{\text{settled}}/L_{\text{packed}}$
PF	Packing factor $PF = L_{\text{cons}}/L_{\text{packed}}$
A_C	Cross sectional area of the column
V_C	Column volume $V_C = L_{\text{packed}} \times A_C$
C_{slurry}	Concentration of the slurry

Preparation of the slurry

To measure the slurry concentration, let the media settle in 20% ethanol at least overnight in a measuring cylinder or use the method for slurry concentration measurement described in application note 28-9259-32. This method can also be used for HiScale columns.

For calculating the amount of medium to fill into the column, use the following equation: $V = (A_C \times L_{\text{packed}} \times CF) / C_{\text{slurry}}$

CF for MabSelect SuRe LX is 1.1 in 20% ethanol

Washing the medium

Mount a glass filter funnel onto a filtering flask. Suspend the medium by shaking and pour into the funnel and wash according to the following instructions:

- 5 times with 5 ml 20% ethanol with 0.4 M NaCl/ml medium
- Gently stir with a spatula between additions.
- Move the washed medium from the funnel into a beaker and add 20% ethanol with 0.4 M NaCl to obtain a 50% slurry concentration.

Packing the column

Table 3. Main features of the packing method for HiScale 16/20 and HiScale 16/40

Column	HiScale 16/20	HiScale 16/40
Bed height (cm)	10	20
Slurry/ packing solution	20% ethanol with 0.4 M NaCl	
Slurry concentration (%)	50	50
Packing factor (PF)	1.10	1.10
Packing velocity (cm/h)	300	300
Packing flow rate (ml/min)	10	10
Flow condition (cm/h)	750	450
Flow condition (ml/min)	25	15

Table 4. Main features of the packing method for HiScale 26/20 and HiScale 26/40

Column	HiScale 26/20	HiScale 26/40
Bed height (cm)	10	20
Slurry/ packing solution	20% ethanol with 0.4 M NaCl	
Slurry concentration (%)	50	50
Packing factor (PF)	1.15	1.13
Packing velocity (cm/h)	300	300
Packing flow rate (ml/min)	27	27
Flow condition (cm/h)	750	450
Flow condition (ml/min)	66	40

Procedure

- 1 Assemble the column according to the column instructions (HiScale columns (16, 26, 50) and accessories, code no 28-9674-70).
- 2 Mount the column tube in a stand.
- 3 Connect the bottom adapter unit to the pump or a syringe and prime the bottom net with a slow flow of packing solution. This is easiest done if the nets are dry but if air is trapped under the net it can be removed by a light suction with a syringe.
- 4 Mount the bottom adapter unit in the bottom of the column tube and tighten the o-ring.
- 5 Fill the column with approximately 1 cm packing liquid using the pump/syringe. Disconnect the pump/syringe and put a stop plug on the outlet.
- 6 Mount the packing tube on top of the column tube.
- 7 Connect the top adapter to the pump and prime it with a slow down flow. The net needs to be facing the roof as this is done.
- 8 Fill the column with slurry suspended in packing solution. If needed, top up the slurry with extra packing solution so the top adapter dips into the slurry to avoid air under the net.
- 9 Mount the top adapter unit on top of the packing tube. Tighten the o-ring firmly and remove the bottom stop plug.
- 10 Start a downward flow, see Table 3 and Table 4.
- 11 Let the flow run until the bed has consolidated.
- 12 Use the scale on the column to measure the bed height. There might be a build up of media at the column wall after the bed is consolidated and to easier see where the top of the bed is, a light source can be used.
- 13 Calculate the final bed height by dividing the bed height with the desired packing factor.
$$L_{\text{packed}} = L_{\text{cons}} / \text{PF}$$
- 14 Turn off the flow and put a stop plug in the bottom.
- 15 Dismount the top adapter from the packing tube.
- 16 Over a beaker or a sink, detach the packing tube from the column.

- 17** Remount the top adapter in the column tube. Make sure no air is trapped under the net and lower the adapter down to 1 to 2 cm above the bed, making sure the surface is not disturbed.
- 18** Tighten the O-ring on the adapter. Remove the bottom stop plug and carefully start turning the end cap down. While spilling out liquid through the bottom, proceed turning until the calculated bed height is reached.
- 19** Make sure that the pressure peaks that occur during turning the end knob down do not exceed the pressure specifications of the media.
- 20** Start a downward flow to flow condition the bed. The flow rate is shown in Table 3 and Table 4.
- 21** Let the flow run for about 10 column volumes. The column is ready to be tested.

6 Evaluation of column packing

Test the column efficiency to check the quality of packing. Testing should be done immediately after packing and at regular intervals during the working life of the column and also when separation performance is seen to deteriorate. The best method of expressing the efficiency of a packed column is in terms of the height equivalent to a theoretical plate (HETP) and the asymmetry factor (A_s). These values are easily determined by applying a sample such as 1% acetone solution to the column. Sodium chloride can also be used as a test substance. Use a concentration of 0.8 M NaCl in water with 0.4 M NaCl in water as eluent.

Note: *The calculated plate number will vary according to the test conditions and it should only be used as a reference value. It is important that test conditions and equipment are kept constant so that results are comparable. Changes of solute, solvent, eluent, sample volume, flow velocity, liquid pathway, temperature, etc. will influence the results.*

For optimal results, the sample volume should be at maximum 2.5% of the column volume and the fluid velocity 30 cm/h. If an acceptance limit is defined in relation to column performance, the column plate number can be used as one of the acceptance criteria for column use.

Method for measuring HETP and A_s

Calculate HETP and A_s from the UV curve (or conductivity curve) as follows:

$$\text{HETP} = \frac{L}{N}$$
$$N = 5.54 \times \left(\frac{V_R}{W_h} \right)^2$$

L = bed height (cm)
 N = number of theoretical plates
 V_R = volume eluted from the start of sample application to the peak maximum
 W_h = peak width measured as the width of the recorded peak at half of the peak height
 V_R and W_h are in the same units

The concept of reduced plate height is often used for comparing column performance.

The reduced plate height, h , is calculated as follows:

$$h = \frac{\text{HETP}}{d_{50v}}$$

d_{50v} = mean diameter of the beads (cm)

As a guideline, a value of < 3 is very good.

The peak should be symmetrical, and the asymmetry factor as close to 1 as possible (values between 0.8 and 1.8 are usually acceptable).

A change in the shape of the peak is usually the first indication of bed deterioration due to excessive use.

Peak asymmetry factor calculation:

$$A_s = \frac{b}{a}$$

a = ascending part of the peak width at 10% of peak height
 b = descending part of the peak width at 10% of peak height

Figure 6 shows a UV trace for acetone in a typical test chromatogram from which the HETP and A_s values are calculated.

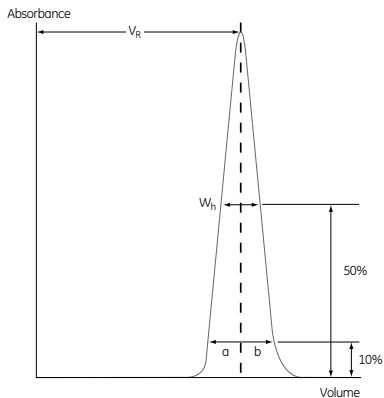


Fig 6. A typical test chromatogram showing the parameters used for HETP and A_s calculations.

7 Cleaning-In-Place (CIP)

Cleaning-In-Place (CIP) is the removal of very tightly bound, precipitated or denatured substances from the purification system. If such contaminants are allowed to accumulate, they may affect the chromatographic properties of the column, reduce the capacity of the column and, potentially, come off in subsequent runs. If the fouling is severe, it may block the column, increase back pressure and reduce flow rate.

Regular CIP prevents the build up of contaminants in the packed bed, and helps to maintain the capacity, flow properties and general performance of MabSelect SuRe LX. We recommend performing a blank run, including CIP, before the first run with antibody feed.

MabSelect SuRe LX is an alkali-tolerant medium allowing the use of NaOH as CIP agent. NaOH is widely accepted for cleaning due to the low cost and the ability to dissolve proteins and saponify fats. For difficult cases where CIP with NaOH is not sufficient to restore the column performance an extended protocol including wash with 100 mM thioglycerol pH 8.5 followed by CIP with 0.1 - 0.5 M NaOH is recommended. For more details, see the application note *High-throughput process development for design of cleaning-in-place protocols* (28-9845-64).

CIP protocol

- 1 Wash the column with 3 CV of binding buffer.
- 2 Wash with at least 2 CV of 0.1–0.5 M NaOH. Contact time 10–15 minutes.
- 3 Wash immediately with at least 5 CV of sterile and filtered binding buffer at pH 7–8.

CIP is usually performed immediately after the elution. Before applying the alkaline NaOH CIP solution, we recommend equilibrating the column with a solution of neutral pH in order to avoid the direct contact between low-pH elution buffer and high-pH NaOH solution on the column. Mixing acid and alkaline solutions might cause a rise in temperature in the column.

NaOH concentration, contact time and frequency are typically the main parameters to vary during the optimization of the CIP. The nature of the feed material will ultimately determine the final CIP. However, the general recommendation is to clean the column at least every 5 cycles during normal use. Depending on the nature of the contaminants, different protocols may have to be combined, for example 0.1 M NaOH every cycle and 0.5 M NaOH every 10 cycles.

8 Sanitization

Sanitization reduces microbial contamination of the chromatographic bed to a minimum. MabSelect SuRe LX is alkali-tolerant allowing the use of NaOH as sanitizing agent. NaOH is very effective for inactivating viruses, bacteria, yeasts, and endotoxins. In addition, NaOH is inexpensive compared with other sanitizing agents.

Sanitization protocol

- 1 Wash the column with 3 CV of binding buffer.
- 2 Equilibrate the column with at least 2 CV of 0.1–0.5 M NaOH.
- 3 Use a contact time of at least 15 minutes for 0.5 M NaOH or 30 minutes for 0.1 M NaOH (see also the note below).
- 4 Wash immediately with at least 5 CV of sterile and filtered binding buffer at pH 7–8.

For more challenging microbial contamination, a mixture of 30% to 40% 1- or 2-propanol in 0.5 M NaOH could be used for sanitization.

Note: *Higher concentrations of NaOH and/or longer contact time inactivates microorganisms more effectively. However, these conditions might also lead to a decrease in the dynamic binding capacity. The conditions for sanitization should therefore be evaluated to maximize microbial killing and to minimize loss of capacity.*

9 Storage

Store unused media in its container at a temperature of +2°C to +8°C. Ensure that the screw top is fully tightened.

Equilibrate packed columns in buffer containing 20% ethanol or 2% benzyl alcohol to prevent microbial growth.

After storage, equilibrate with starting buffer and perform a blank run, including CIP, before use.

10 Scaling up

After optimizing the antibody fractionation at laboratory scale, the process can be scaled up to pilot and process scales.

- Keep the residence time constant in order to maintain the dynamic binding capacity.
- Select bed volume according to required binding capacity. Verify the purification step with the new bed height, if it is changed.
- Select column diameter according to your volume throughput requirements. Then determine the bed height to give the desired residence time. Bed heights of 10–25 cm are generally considered appropriate. Note that the backpressure increases proportionally with increasing bed height at constant nominal velocity.
- Keep sample concentration and elution conditions constant.

See also Figure 4 for appropriate windows of operation for MabSelect SuRe LX.

11 Troubleshooting

The list describes faults observed from the monitor curves.

Fault	Possible cause/corrective action
High backpressure during the run	<ul style="list-style-type: none">• Change the in-line filter.• The column is clogged. Perform CIP.• The adapter net/filter is clogged. Clean or replace the net/filter.
Unstable pressure curve during sample application	<ul style="list-style-type: none">• Remove air bubbles that might have been trapped in the sample pump.• Degas the sample using a vacuum degasser or an air trap.
Gradual broadening of the eluate peak	<ul style="list-style-type: none">• Might be due to insufficient elution and CIP caused by contaminants accumulating in the column. Optimize the elution conditions, the CIP protocol and/or perform CIP more frequently.
Gradual decrease in yield	<ul style="list-style-type: none">• Too high sample load. Decrease the sample load.• Precipitation during elution. Optimize the elution conditions.• Might be due to insufficient elution and CIP. Optimize the elution conditions, the CIP protocol and/or perform CIP more frequently.
Gradual increase in CIP peaks	<ul style="list-style-type: none">• Might be due to insufficient elution or CIP. Optimize the elution conditions, the CIP protocol and/or perform CIP more frequently.
High ligand leakage during the first purification cycle	<ul style="list-style-type: none">• Perform a blank run, including CIP, before the first purification cycle on a new column.

12 Ordering information

Product	Quantity	Code No
MabSelect SuRe LX	25 ml	17-5474-01
	200 ml	17-5474-02
	1 l	17-5474-03
	5 l	17-5474-04
	10 l	17-5474-05

Related product	Quantity	Code No
HiScreen MabSelect SuRe LX	1 × 4.7 ml	17-5474-15
PreDictor MabSelect SuRe LX, 6 µL	4 × 96-well plates	17-5474-30
PreDictor MabSelect SuRe LX, 20 µL	4 × 96-well plates	17-5474-31
PreDictor MabSelect SuRe LX, 50 µL	4 × 96-well plates	17-5474-32
HiScale 16/20	1	28-9644-41
HiScale 16/40	1	28-9644-24
HiScale 26/20	1	28-9645-14
HiScale 26/40	1	28-9645-13
HiScale 50/20	1	28-9644-45
HiScale 50/40	1	28-9644-44

Related literature		Code No.
Data Files	AxiChrom Columns	28-9290-41
	BPG columns	18-1115-23
	Chromaflo columns	18-1138-92
	MabSelect SuRe LX	28-9870-62
Application notes	MabSelect SuRe – Leakage and Toxicity	11-0011-64
	MabSelect – Column packing	11-0007-52
	MabSelect SuRe LX - Life time performance study	28-9872-96
	MabSelect SuRe LX - High capacity affinity medium for high titer monoclonal antibodies	28-9875-25
	Two step purification of monoclonal IgG ₁ from CHO cell culture supernatant	28-9078-92
	High-throughput process development for design of cleaning-in-place protocols	28-9845-64
Movie	Column Packing - The Movie	18-1165-33
Instructions	HiScale columns (16, 26, 50) and accessories	28-9674-70

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