

Consistent Size-based Separations Across LC Systems with SEC 125 Å MaxPeak Premier Columns

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Published on September 24 2025

Application Brief

This is an Application Brief and does not contain a detailed Experimental section.

Abstract

In this Application Brief, consistent size-based separations of peptides and small proteins when transferring methods from UHPLC to HPLC have been demonstrated with SEC 125 Å, MaxPeak™ Premier Columns.

Benefits

- Reliable, high-resolution size-based separations of peptide-based therapeutics (*e.g.*, insulin, GLP-1 receptor agonists) and small proteins.
 - Excellent SEC method transferability from UHPLC to HPLC.
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Introduction

Waters ACQUITY™ Premier Systems in combination with ACQUITY 125 Å SEC, MaxPeak Premier 1.7 µm Columns, enable highly efficient size-based separations, reducing run times as well as sample and eluent consumption.¹ The speed and efficiency achieved with UHPLC provide clear advantages in R&D and analytical development labs where sample volumes may be limited, and method parameters are being evaluated. However, in some organizations traditional HPLC continues to be commonplace for routine QC analyses, meaning methods should be designed with transferability in mind. The ability to move analytical methods and product knowledge from R&D through development to manufacturing QC without compromising accuracy, precision, or regulatory compliance is critical, and hinges upon proper column selection.

Sub-2 µm size-exclusion chromatography (SEC) columns with a 4.6 mm ID lose their efficiency benefit with increased system dispersion volume because of their inherently small peak volumes (see Waters application note [720006337](#) for a systematic in-depth demonstration of LC dispersion volume impact on protein separations for different SEC column configurations).² Therefore, HPLC applications require a larger diameter column with subsequently larger peak volumes to mitigate the impact of system dispersion volume on separation efficiency. Waters 7.8 mm ID SEC Columns demonstrate consistent chromatographic efficiency across a range of higher system dispersion volumes typical of HPLC instruments,² albeit with increased time and material consumption compared to the smaller UHPLC column.

The ability to achieve highly similar chromatographic resolution across two LC systems having different system dispersion volumes has been demonstrated here by appropriately scaling analytical methods from a 4.6 mm ID ACQUITY 125 Å SEC, MaxPeak Premier, 1.7 µm Column to a 7.8 mm ID XBridge™ 125 Å SEC, MaxPeak Premier, 2.5 µm Column. Flow rates were scaled to achieve a constant reduced linear velocity across both the ACQUITY and XBridge Columns, and injection volumes were selected to maintain a consistent sample to column volume ratio.³⁻⁴ The USP High Molecular Weight Insulin Standard and the Waters BEH125 SEC Protein Standard Mix (p/n: [186006519](#) <<https://www.waters.com/nextgen/global/shop/standards--reagents/186006519-beh125-protein-standard-mix.html>>) were analyzed as representative peptide and small protein samples.

$$F_2 = \frac{F_1 \times d_{p,1} \times D_{c,2}^2}{d_{p,2} \times D_{c,1}^2}$$

In Equation (1), flow rate (F) for the second column is determined for constant reduced linear velocity by

considering the differences in both the column diameter (D_c) and particle diameter (d_p). Adjusting flow rate to maintain constant reduced linear velocity is prescribed by USP <621> for chromatography method adjustment.

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Results and Discussion

SEC analysis of human insulin was performed under denaturing eluent conditions to separate insulin monomer from covalent high molecular weight products (HMWPs) using four different SEC 125 Å, MaxPeak Premier Columns (see Table 1 for summary of LC conditions). Figure 1 shows the insulin chromatograms for the four different column configurations, along with relative peak areas for the HMWPs and the USP monomer half-height (HH) resolutions. The resolution values and HMWPs relative peak areas were comparable for the two 150 mm columns and for the two 300 mm columns, despite a 32% reduction in L/d_p ratio from the ACQUITY Column to the XBridge Column. The insulin dimer analysis example highlights the potential of matching ACQUITY Column performance on an LC system with higher dispersion volume by utilizing a 7.8 mm ID Waters column leveraging the same BEH particle technology. However, in some cases, it may be necessary to maintain or increase L/d_p to achieve the desired resolution for the transferred method, as seen in the next example.

125 Å SEC MaxPeak Premier Column Configuration	L/d_p	LC System	LC System Dispersion Volume (μ L)	Flow Rate, Insulin (mL/min)	Column Back Pressure, Insulin (psi)	Flow Rate, Protein Mix (mL/min)	Column Back Pressure, Protein Mix (psi)
4.6 x 150 mm 1.7 μ m	88,235	ACQUITY Premier System	<9	0.4	4,892	0.3	2,587
4.6 x 300 mm 1.7 μ m	176,471	ACQUITY Premier System	<9	0.4	10,650	0.3	5,837
7.8 x 150 mm 2.5 μ m	60,000	Arc Premier System	<30	0.78	734	0.59	325
7.8 x 300 mm 2.5 μ m	120,000	Arc Premier System	<30	0.78	1,872	0.59	950

Table 1. Summary of LC conditions (unless otherwise noted).

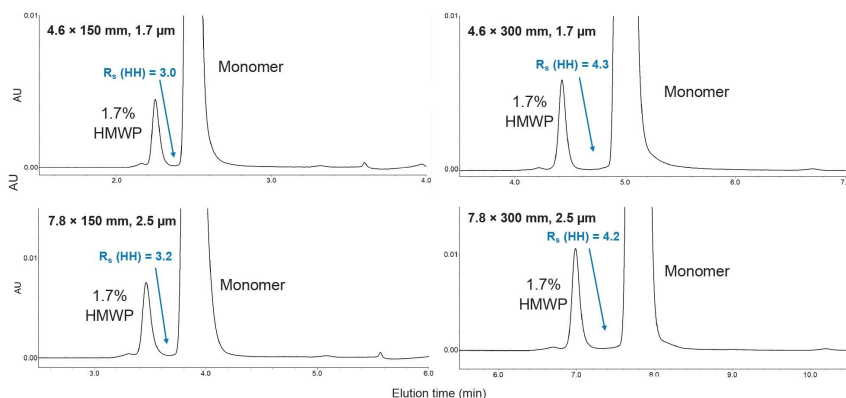


Figure 1. USP HMW Insulin Standard analyzed with ACQUITY 125 Å SEC, MaxPeak Premier, 1.7 μm, 4.6 x 150 mm and 300 mm Columns on an ACQUITY Premier System, and XBridge 125 Å SEC, MaxPeak Premier, 2.5 μm, 7.8 x 150 mm, and 300 mm Columns on an Arc™ Premier System. Elution condition: 65%/20%/15% 1 g/L arginine / acetonitrile / glacial acetic acid (USP <121.1>). The flow rates were 0.4 mL/min for the 4.6 mm ID columns and 0.78 mL/min for the 7.8 mm ID columns.

Analysis of Waters BEH125 SEC Protein Standard Mix (p/n: [186006519 <https://www.waters.com/nextgen/global/shop/standards--reagents/186006519-beh125-protein-standard-mix.html>](https://www.waters.com/nextgen/global/shop/standards--reagents/186006519-beh125-protein-standard-mix.html)) under native SEC eluent conditions (Figure 2) showed similar chromatograms for both the ACQUITY and XBridge 125 Å SEC MaxPeak Premier Columns. Average peak capacity and reduced plate height (RPH) values are shown in Table 2 for each of the four columns evaluated. The ACQUITY 125 Å SEC, MaxPeak Premier Columns showed a higher average peak capacity compared to XBridge 125 Å SEC, MaxPeak Premier Columns of the same length. The RPH values, which allow for a comparison of peak efficiency across columns of differing particle sizes, were highly similar for each column length.⁶

When transferring LC methods, the ratio of column length to particle diameter (L/d_p), which directly corresponds to column efficiency, would ideally be held constant. However, it is not always possible to easily match the L/d_p value when transferring from UHPLC to HPLC due to limited availability of column configurations. In cases where any reduction in separation efficiency is not acceptable for the method transfer,

an increase in L/d_p should be considered. USP <621> allows for up to a 50% increase in L/d_p for monograph methods.⁵ For example, a method initially developed using a 4.6 x 150 mm ACQUITY 125 Å SEC, MaxPeak Premier Column ($L/d_p = 88,235$) could be transferred to a 7.8 x 300 mm XBridge 125 Å SEC, MaxPeak Premier Column ($L/d_p = 120,000$), representing a 36% increase in L/d_p . Based on the peak capacity results shown in Table 2, the 300 mm XBridge Column had increased chromatographic resolution for the protein mix sample at the same reduced linear velocity. Increasing the linear velocity (*i.e.*, flow rate) for the analysis with the 300 mm XBridge Column will decrease the observed peak capacity. To demonstrate this effect, an additional flow rate of 0.86 mL/min was tested, resulting in an average peak capacity of 23.2, a reduction from the 25.2 value observed at 0.59 mL/min and closer to the average peak capacity of 21.4 achieved with the 4.6 x 150 mm ACQUITY Column (Table 2). When transferring a method to a column with a higher L/d_p ratio, the potential exists to increase the flow rate above the value calculated with Equation (1) and still achieve the desired chromatographic resolution.

In this Application Brief, SEC analysis of representative peptide and protein samples with Waters ACQUITY and XBridge 125 Å SEC, MaxPeak Premier Columns demonstrated the ability to transfer methods across different LC systems while maintaining consistent separation efficiency. While the 7.8 mm XBridge 125 Å SEC, MaxPeak Premier, 2.5 µm Columns require larger volumes and longer run times, they provide comparable separations on LC systems with higher dispersion volumes at lower operating pressures.

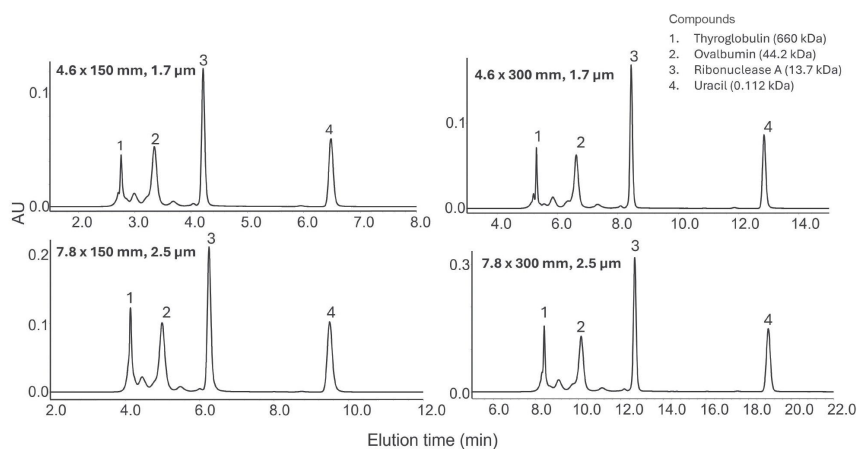


Figure 2. Waters BEH125 SEC Protein Standard Mix analyzed with ACQUITY 125 Å SEC, MaxPeak Premier, 1.7 μm, 4.6 x 150 mm and 300 mm Columns on an ACQUITY Premier System, and with XBridge 125 Å SEC, MaxPeak Premier, 2.5 μm, 7.8 x 150 mm, and 300 mm, Columns on an Arc Premier System. Elution condition: 50 mM sodium phosphate (pH 7), 200 mM potassium chloride. The flow rates were 0.3 mL/min for the 4.6 mm ID columns and 0.59 mL/min for the 7.8 mm ID columns.

Column Configuration	Average Peak Capacity,* P _{0.4σ}	Reduced Plate Height (USP)			
		Peak 1	Peak 2	Peak 3	Peak 4
4.6 x 150 mm 1.7 μm	21.4	3.8	13.8	3.2	2.7
7.8 x 150 mm 2.5 μm	19.2	3.6	12.6	3.1	2.8
4.6 x 300 mm 1.7 μm	30.1	2.9	17.0	3.2	2.6
7.8 x 300 mm 2.5 μm	25.2	2.6	13.2	2.9	2.8

*Peak capacities calculated⁷ based on average peak width (4σ) for Peaks 2 and 3, and the difference in retention times between Peaks 1 and 4.

Table 2. Peak resolution and reduced plate height for chromatograms of SEC125 Protein Standard Mix (shown in Figure 2) on 125 Å SEC, MaxPeak Premier Columns.

Conclusion

By incorporating the same QC batch-tested BEH particle chemistry across multiple column configurations and particle sizes, SEC 125 Å, MaxPeak Premier Columns enable analytical method development and QC transferability for size-based separations of peptides and small proteins. Points of note when transferring methods include:

- Scale flow rates to maintain constant reduced linear velocity.
- Matching L/d_p is ideal for achieving consistent separation efficiency but is not always viable.
- Optimizing flow rate empirically may be necessary to match resolution when constant L/d_p cannot be maintained.

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720009011, September 2025



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