# **MODERNIZING BIOTHERAPEUTIC COMPENDIAL METHODS WITH A NEXT-GENERATION HPLC SYSTEM**

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### INTRODUCTION

As the field of analytics continues to evolve, methods guided by regulatory general chapters are increasingly seen as outdated due to their limited resolving power and speed. These compendial methods were originally written when larger particle sizes and systems with larger system dispersion were the norm. In response, regulatory agencies have lifted restrictions on method parameters such as flow rate, column dimensions, and particle size. This allows scientists using compendial methods to leverage modern instruments and particle sizes for a significant enhancement in performance.

In this poster, an Alliance™ iS Bio HPLC System with several columns containing MaxPeak<sup>™</sup> High Performance Surfaces (HPS) Technology were used to modernize compendial biotherapeutic methods. To assess the benefits of method modernization by utilizing this technology, the modernized methods were compared to the regulatory general chapter methods analysed on a legacy HPLC system, evaluating them based on speed and recovery.

### **METHODS**

The compendial methods evaluated on both systems are from USP General Chapters <129> and <121.1>. For USP <129>, size exclusion chromatography (SEC) was used for size variant analysis of USP System Suitability and USP mAb reference standards.<sup>1</sup> For USP <121.1>, a reversed phase chromatography (RPLC) gradient was used to evaluate a peptide map of insulin.<sup>2</sup>

#### Monoclonal Antibody Analysis<sup>3</sup>

USP mAb reference standards were injected at a concentration of 10 mg/mL in formulation buffer onto both systems.

#### Compendial Method Conditions:<sup>1</sup>

System:	Legacy HPLC system
Column:	BioSuite™ Diol (OH) Column, 250Å,
Injection volume:	5 μm, 7.8 mm x 300 mm (p/n: 186002165)
injection volume.	20 μL
Flow Rate:	0.500 mL/min
Run Time:	30 minutes, isocratic

#### **Modernized Method Conditions:**

System:	Alliance iS Bio HPLC System
Column:	XBridge™ Premier Protein SEC Column 250Å,
	2.5 µm, 4.6 x 150 mm (p/n: 186009959RF)
Injection volume:	3.5 µL
Flow Rate:	0.350 mL/min
Run Time:	7.5 minutes, isocratic

#### USP <129> 0.0030compendial SEC method was 0.0025 Start p/v analyzed on a ⊋0.0020 1.13 Analytes legacy HPLC 0.0015 system and the 0.0010-Metal Alliance iS Bio Ti Fe 0.0005 HPLC System. 0.0000 Without any 10.0 12.0 20.0 14.0 16.0 18.0 modifications, the Minute Figure 1: The Alliance iS Bio Alliance iS Bio Alliance iS 0.0040 HPLC System is a bio-inert HPLC System system with MaxPeak HPS **Bio HPLC** 0.0035 showed Technology that was designed System 0.0030 improvements in for separating biomolecules. resolution (peak 0.0025 This advanced surface Start p/v height to valley technology eliminates risk when ₽0.0020 ratio) for more 1.74 analyzing biotherapeutics which 0.0015 accurate may have the potential to adsorb 0.0010 quantification of to metal surfaces. 0.0005 mAb impurities. 20.0 10.0 12.0 18.0 14.0 16.0 Minutes Legacy HPLC System 0.008 USP mAb 003, USP mAb 002. XBridge Premier Protein SEC 0.007 0.007 Monoclonal IgG1 Monoclonal IgG1 0.006 0.006 7.8 x 300 mm Column 250Å, 2.5 µm, 7.8 x 150 mm 7.8 x 300 mm 0.005 0.005 5 µm 5 µm 4X Reduced runtime: ₽ 0.004 Q 0.004 0.003 30 min — 7.5 min 0.003 0.002 0.002 LMWS2 0.001 0.00 Improved LMWS Start p/v 0.000 0.000 resolution: 14.0 12.0 16.0 Minutes 16.0 Minutes 20.0 18.0 Chromatogram LMWS1 LMWS2 Alliance iS Bio HPLC System 0.001 0.002 0.001 0.001 0.001 0.001 0.001 N.D. 1.13 USP mAb 003, USP mAb 002, ∧LMWS1 0.007 1.22 3.59 Monoclonal IgG1 Monoclonal IgG1 0.006 0.005 7.8 x 150 mm 4.6 x 150 mm ₹ 0.004 XBridge Premier Protein SEC 2.5 µm 2.5 µm 0.003 Column 250Å, 2.5 µm, 4.6 x 150 mm 0.002 LMWS2 LMWS3 0.001 6-fold reduction 3.0 3.5 Minutes 2.5 3.0 3.5 Minutes in solvent and sample use

Figure 3: Following the scaling guidance in USP <621>, the compendial SEC method was modernized using XBridge Premier Protein SEC columns. Two columns were selected that maintained the same resolving power, or length-to-particle size ratio, as the original method. The 7.8 mm ID column reduced the method runtime 4-fold and provided the best resolution for the low molecular weight species (LMWS) whereas the 4.6 mm ID column provided

## **RESULTS AND DISCUSSION**

0.0035

MaxPeak HPS Technology

0.0040 Legacy HPLC

System

Column:	XBridge Premier Protein SEC Column 250Å, 2.5 µm, 7.8 x 150 mm (p/n: 186009961)		
Injection volume:	10 µL		
Flow Rate:	1.000 mL/min		
Run Time:	7.5 minutes, isocratic		

#### Shared Conditions:<sup>1</sup>

Mobile Phase:	0.20 M potassium phosphate and 0.25 M potassium chloride, pH 6.2
Column temp.:	30 °C
Wavelength:	280 nm

#### **Insulin Peptide Analysis**

Insulin analogs were digested with Glu-C following the digestion procedure outlined in USP <121.1>.<sup>2</sup> Digesting samples were incubated at 25°C for 6 hours and quenched with sulfate buffer prior to injection.

#### Compendial Method Conditions:<sup>2</sup>

-	
System:	Legacy HPLC system
Column:	XSelect™ Peptide CSH™ C <sub>18</sub> Column 130Å,
	5 µm, 4.6 x 100 mm (p/n: 186005289)
Injection volume:	50 µL
Flow Rate:	1.000 mL/min
Run time:	50 minutes, gradient

Time (min)	Flow (mL/min)	%A	%B	Curve
Initial	1.000	95.0	5.0	Initial
3.00	1.000	95.0	5.0	6
30.00	1.000	41.0	59.0	6
35.00	1.000	20.0	80.0	6
40.00	1.000	95.0	5.0	6
50.00	1.000	95.0	5.0	6

#### **Modernized Method Conditions:**

System:	Alliance iS Bio HPLC System		
Column:	XSelect Premier Peptide CSH C <sub>18</sub> Column 130Å		
	2.5 µm, 4.6 x 50 mm (p/n: 186009907)		
Injection volume:	25 µL		
Flow Rate:	2.000 mL/min		
Run Time:	12.5 minutes, gradient		

Time (min)	Flow (mL/min)	%A	%B	Curve
Initial	2.000	95.0	5.0	Initial
0.75	2.000	95.0	5.0	6
7.50	2.000	41.0	59.0	6
8.75	2.000	20.0	80.0	6
10.00	2.000	95.0	5.0	6
12.50	2.000	95.0	5.0	6

#### Shared Conditions:<sup>2</sup>

700:100:200 H <sub>2</sub> O:ACN:(NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub> buffer
400:400:200 H <sub>2</sub> O:ACN:(NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub> buffer
40 °C
214 nm

benefits in reducing laboratory consumption by reducing solvent and sample use 6-fold.







Figure 4: Following digestion with Glu-C, insulin human produces four peptide fragments that can be used in identification assays. The digested product was analyzed on both the legacy HPLC system and the Alliance is Bio HPLC System and provided comparable performance for area and retention time.

### CONCLUSION

- Compendial SEC and insulin peptide mapping methods were scaled to 2.5 µm columns and provided a significant reduction in analysis time and mobile phase consumption.
- The Alliance iS Bio HPLC System is capable of migrating and modernizing compendial methods to accommodate both present and future biopharmaceutical workflows in QC environments.



Chromatogram MP

per injection

15 mL

Sample

20 µL

Waters™ alliance is bio

Figure 2: The

#### XSelect Peptide CSH C<sub>18</sub> Column 130Å, <u>5 µm, 4.6 x 100 mm</u>



#### XSelect Premier Peptide CSH C<sub>18</sub> Column 130Å, 2.5 µm, 4.6 x 50 mm



Figure 5: Taking advantage of the separation efficiency gains of smaller particles, the compendial method was scaled to a 50 mm column packed with 2.5 µm sized particles as the stationary phase on the Alliance iS Bio HPLC System. The selectivity of the separation was preserved and the compendial method requirements for resolution ( $\geq$  3.4) and peak tailing ( $\leq$  1.5) of the encircled critical pair were met. The Alliance iS Bio HPLC System supports 2.5 µm column technology due to higher system pressure tolerance and lower system dispersion, thereby significantly reducing operating costs such as analysis time, solvent, and sample use.

#### References

- 1. USP. Chromatography <621>. In: USP-NF. Rockville, MD: USP; Dec 1, 2022.
- USP. Physicochemical Analytical Procedures for Insulins <121.1>. In: USP-NF. Rockville, 2. MD: USP; Dec 1, 2016.
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