The Integration of Mass Spectrometry to the Process Development of Fusion Protein Therapeutics

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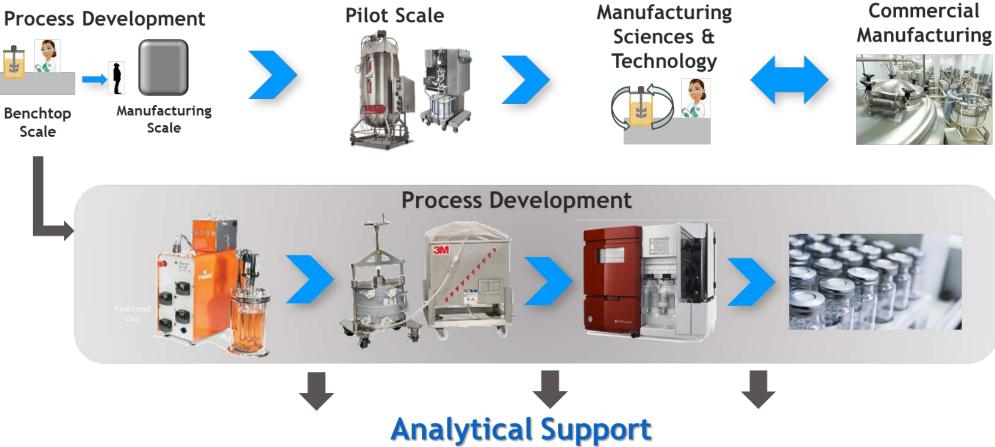
2021 CASSS Northeast Regional Forum



Outline

- Introduction
- Site-Specific Glycosylation Monitoring with LC-MS
- Summary
- Acknowledgement

Bioprocess Development Analytical Support



- High-throughput Platform Assays
- Sample Automation
- PAT for Real time Analytics
- Mass Spectrometry for In-Process Support

Fc-fusion Protein Therapeutics

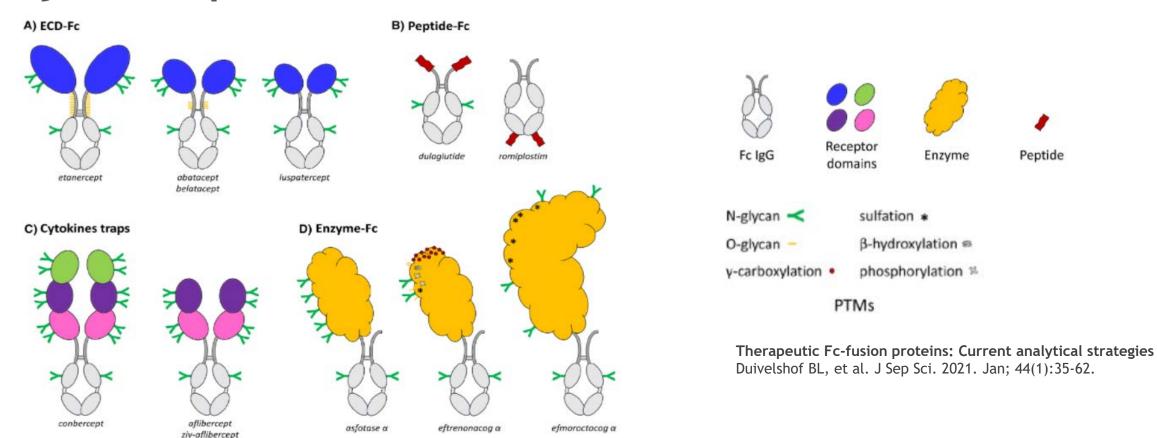
• Fc-fusion protein therapeutics are one of the most successful classes of IgG-based products;

Product	Sales in 2019 (billion, USD)
Eylea	7.5
Enbrel	7.2
Trulicity	4.3
Orencia	3.2
Elocta	1.2

- Fc-fusion protein combine the pharmacological properties of biological ligands with the additional stability and inherent properties of IgG Fc domain;
- Fc-fusion protein can significantly improve the clinical potential of active protein drugs such as extend the plasma half life as well as engage immune-mediated effector functions;
- To date, ~37 therapeutic fusion proteins are in clinical development and 13 products have been approved by the FDA, CFDA, and EMA;
 - Enbrel (TNFR-Fc fusion, FDA approval in 1998) and Orencia (CTLA4-Fc fusion, FDA approval in 2005) for treatment of rheumatoid arthritis;

Duivelshof BL, et al. J Sep Sci. 2021. Jan; 44(1):35-62.

Major Groups of Fc-fusion Proteins



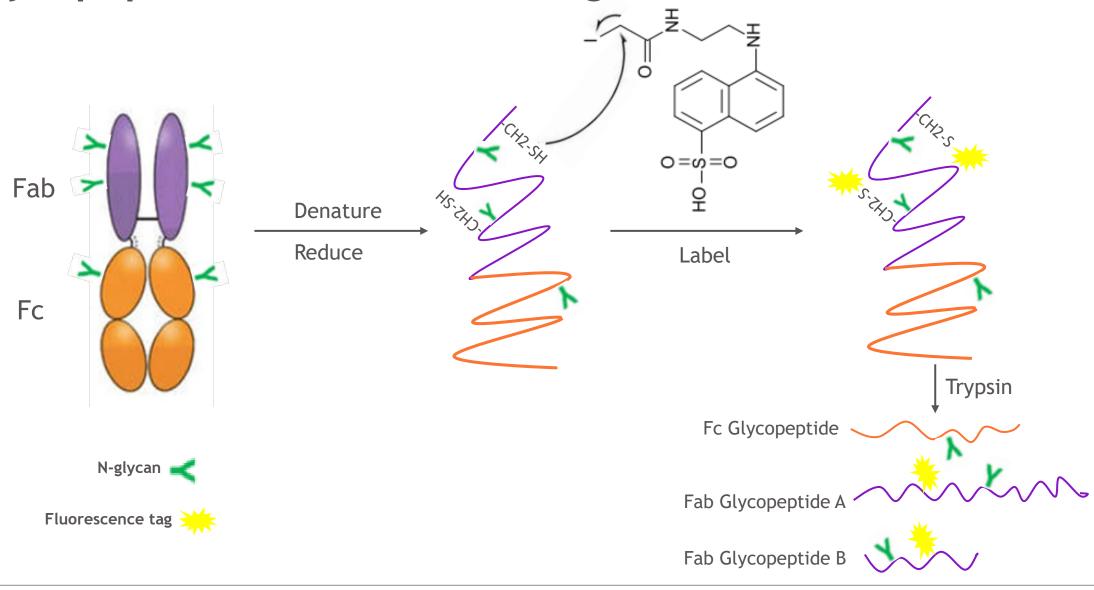
- Due to highly **heterogenous** structure (the presence of sialic acid, complex glycan structure, etc.), the analysis of Fc-fusion proteins is more challenging and complex than monoclonal antibodies;
- **Product-specific** methods over conventional generic or platform methods are often desirable to support process development;

Peptide

Fab Site-Specific Glycosylation Monitoring

- A Fc-fusion protein with **four** *N*-glycosylation sites in Fab region and **two** *N*-glycosylation sites in the Fc region;
- A LC method was developed to monitor <u>Fab site-specific glycoforms</u> (G0F, G1F, G2F, G2FS1, and G2FS2) as the understanding about site-specific glycosylation as pCQA is continuously evolving during product development lifecycle;
- Mass spectrometry is coupled with LC for the peak identification of the complex chromatogram and the optimization of LC method parameters;
 - Mobile phase screening
 - Fluorescence tag screening
 - LC gradient optimization

Glycopeptide Chemical Labeling



Workflow Overview

Label

- Denature
- Reduce
- Label Cys with fluorescence tag (Alkylation)

Desalt

• Sample desalt with SEC spin column

Proteolysis

Trypsin digestion

LC Separation

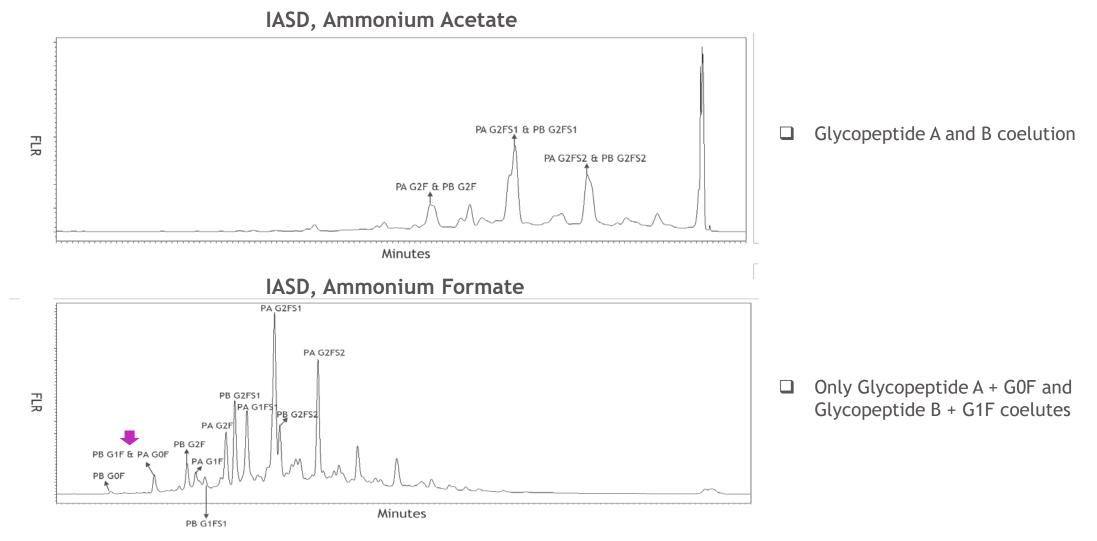
HILIC for glycopeptide separation

MS Identification & FLR Quantitation

- MS for peak identification
- FLR for Fab glycopeptide relative quantitation



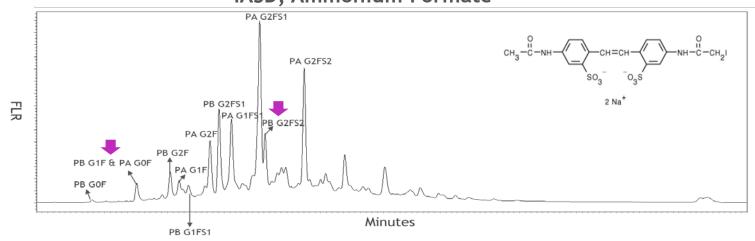
Ammonium Acetate vs Ammonium Formate as Mobile Phase



• Significant improvement in separation resolution with ammonium formate for IASD FLR tag;

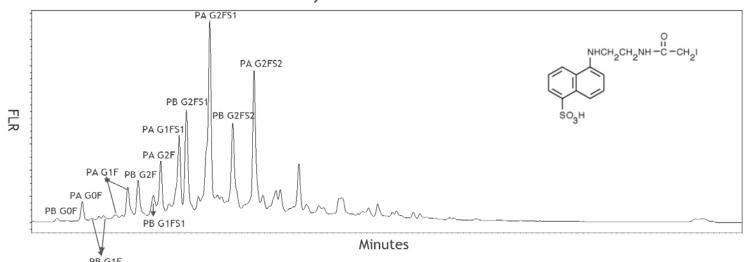
IASD vs IAEDANS as Fluorescence Tag

IASD, Ammonium Formate



Only Glycopeptide A + G0F and Glycopeptide B + G1F coelutes

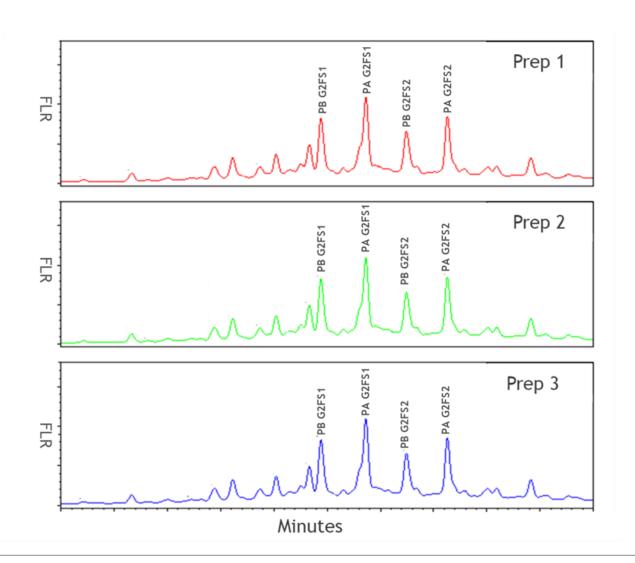
IAEDANS, Ammonium Formate

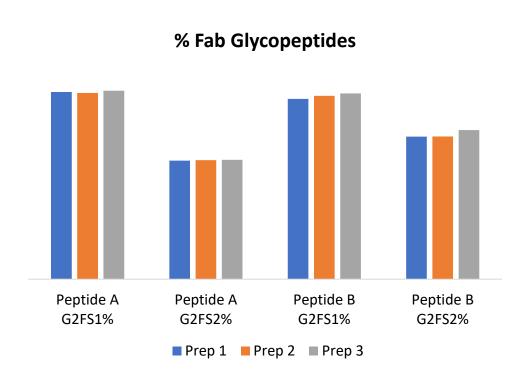


- Glycopeptide A + G0F and Glycopeptide B + G1F were separated
- Better resolution between GlycopeptideA+ G2FS1 and Glycopeptide B+ G2FS2

• Different glycoforms of Fab glycopeptide A and B are fully resolved with IAEDANS FLR tag;

Quantitation of Fab Site-Specific Glycopeptides through Fluorescence Peak Intensity





• Reproducible quantitation for glycopeptides;

Summary

- A HILIC LC-FLR-MS method was established to separate glycopeptides and quantify Fab site-specific glycosylation for a Fc-fusion protein;
- Molecule specific methods are often required to support the development of complex protein therapeutics;
- Creativity and collaboration cross functional areas is essential to develop innovative methodology for problem solving;

Acknowledgement

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Thank You for Your Attention!



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