

APPLICATIONS

Charge Variant Analysis of NIST mAb using a bioZen™ 6 µm WCX Column with a pH Gradient and Native MS Detection

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Overview

Charge variants of proteins commonly result from post translational modifications (PTMs) during recombinant production. These PTMs, including C-terminal lysine clipping, deamidation and glycosylation, give rise to changes in the acidic and basic charged residues relative to the native protein. The most common method to detect and assess these acidic and basic variants is cation-exchange chromatography (CEX), typically weak cation-exchange (WCX). Ion-exchange, unlike most other interactive chromatography mechanisms, is an on / off process and relies on the electrostatic interaction of analyte with stationary phase. The retention of an ionic compound depends on the number of charges as well as their location on the molecule. Elution is accomplished by increasing the concentration of salt when using a salt gradient or increasing the pH of the mobile phase when using a pH gradient. Optimization of the elution conditions is important to ensuring a good, reproducible method that gives consistent results. Ion exchange chromatography is a powerful technique for identifying charge heterogeneity of a monoclonal antibody. However, the presence of a previously undetected isoform typically leads to further analysis requirements which in themselves can be problematic. For example, the process used to identify unknown isoforms may create further post translational modifications. However, using native conditions for cation-exchange chromatography and high resolution mass spectrometry eliminates these issues. The utility of combining these techniques is reported in this application using the NIST monoclonal antibody reference standard which is a common mAb used for system evaluation.

In this application we show the separation of several charge variants of the NIST mAb under native ion exchange and mass spectrometry (high resolution MS) conditions using a pH gradient formed from volatile buffers. The novel buffers used for the pH gradient are stable for at least one week at room temperature when an inlet air filter is used (ex. SecurityCAP™) on the reservoir. The deconvoluted mass spectrum of the main peak is shown with the primary glycoforms identified. The main peak spectrum is also compared to the two known lysine variants.

LC Conditions

Column: bioZen 6 µm WCX

Dimension: 250 x 2.1 mm

Part No.: 00G-4777-AN

Recommended Guard: SecurityGuard™ ULTRA

Guard Cartridge Part No.: [AJ0-9850](#)

Guard Holder Part No.: [AJ0-9000](#)

Mobile Phase: A: 20 mM Ammonium Acetate, pH 5.2
B: 5 mM Ammonium Acetate, pH 10.2

Gradient: 60-100 % in 25 min

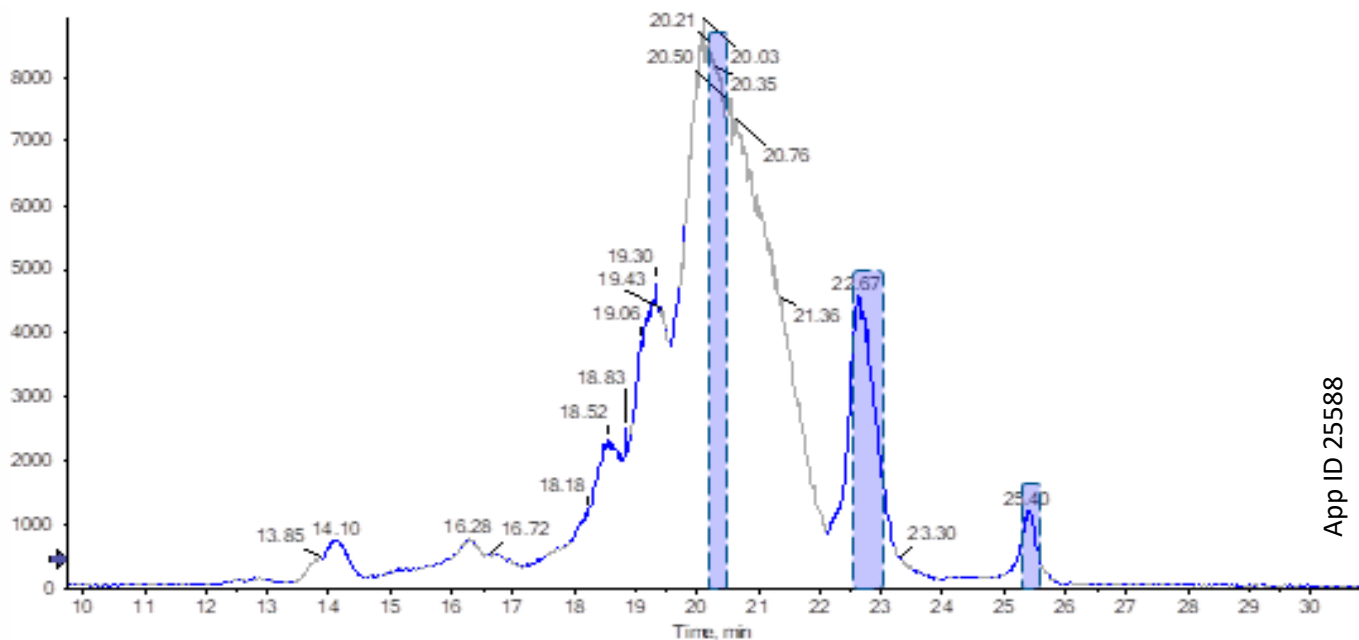
Flow Rate: 0.2 mL/min

Temperature: 30 °C

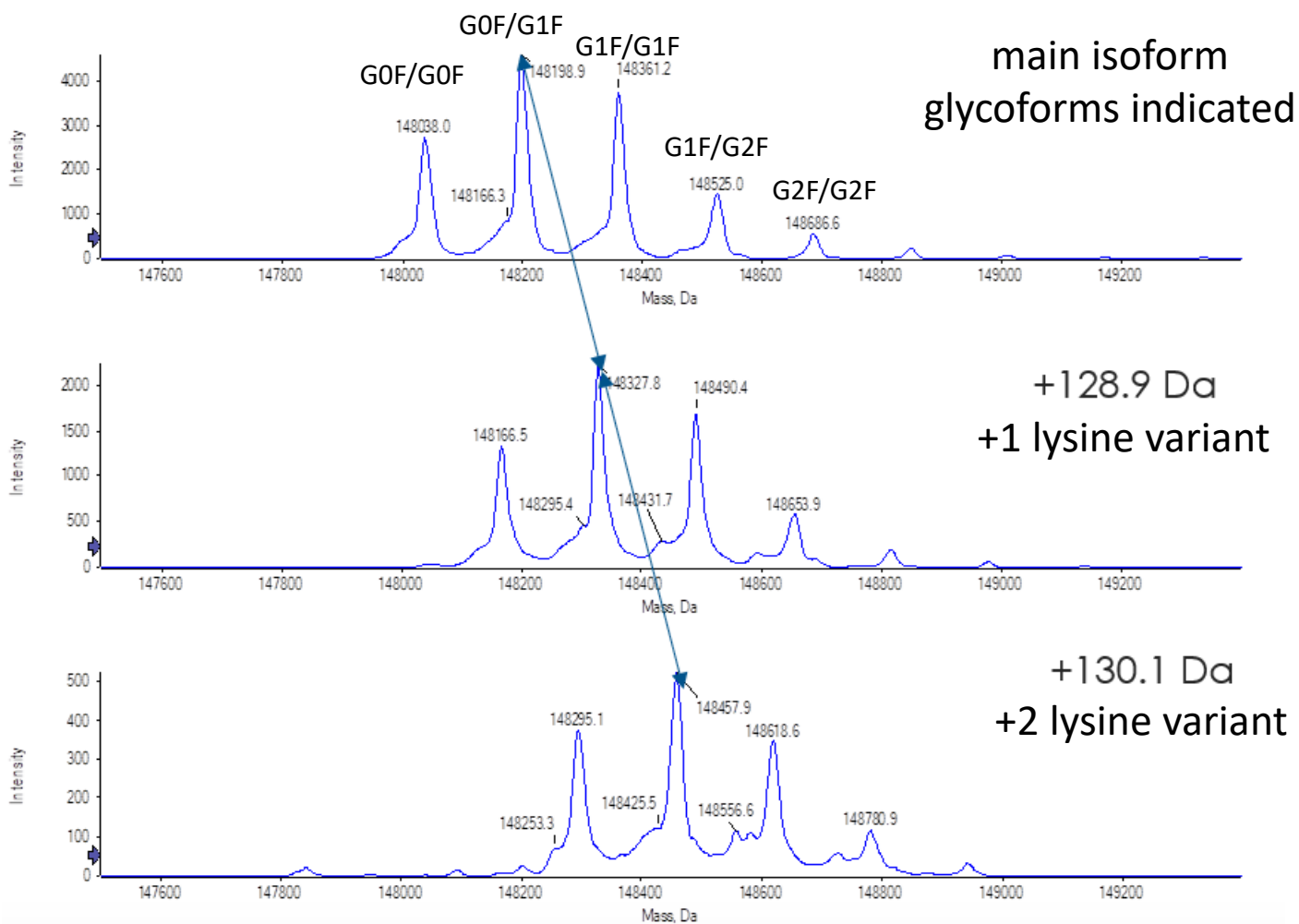
Detector: QTOF (SCIEX® X500B)

Sample: NIST mAb, 100 µg

Separation of NIST mAb, Lysine Variants



App ID 25588



APPLICATIONS

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