

### Laser-Free Oxidation (Fox<sup>™</sup>) Hydroxyl Radical Protein Footprinting (HRPF) for mAb Drug Development

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- Global mAb therapy market is projected to grow from \$178.50 billion in 2021 to \$451.89 billion in  $2028^{1}$ 
  - Biosimilars
- In-depth analytical studies to understand physicochemical and functional characteristics
  - Epitope and Paratope Characterization
  - Intermolecular interactions/ Aggregation

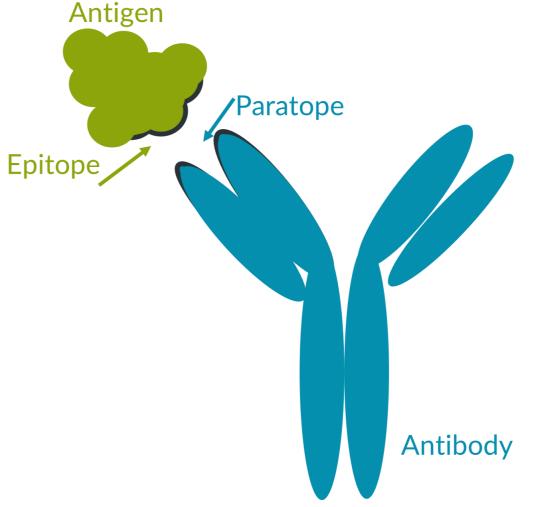
1. Fortune Business Insights. August 30, 2021



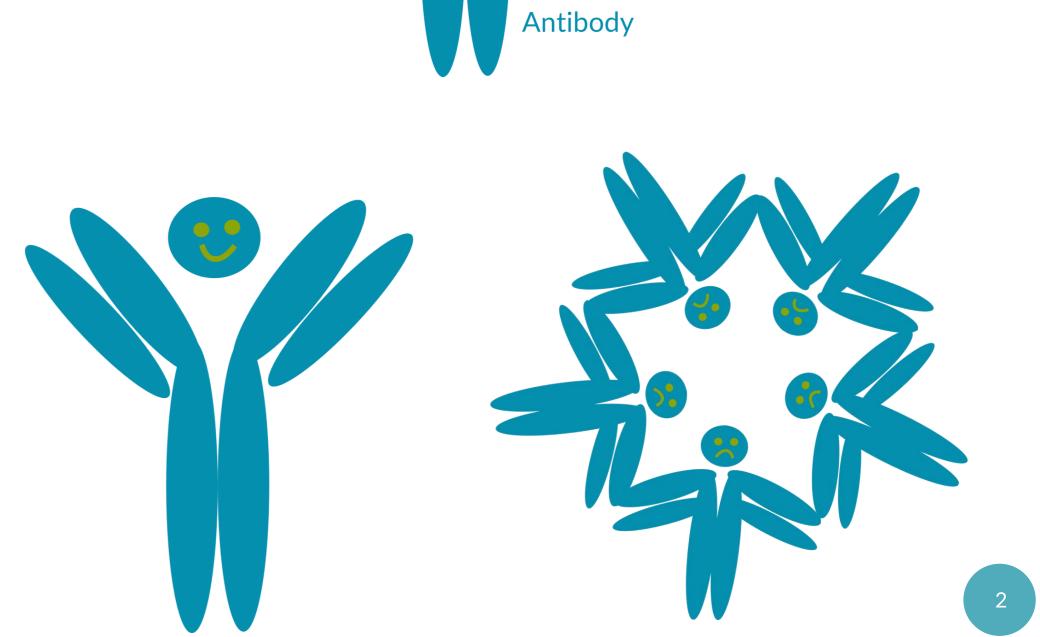


Epitope and Paratope Mapping









### Overview of HOS Approaches

Thermal shift assay, chromatography, circular dichroism (CD), microfluidic modulation spectroscopy (MMS), infrared (IR) spectroscopy, light scattering, surface plasmon resonance (SPR), isothermal titration calorimetry (ITC), etc.



#### HIGH RESOLUTION DATA HARDER TO PERFORM HIGHER COST

### LOW RESOLUTION DATA:

### • PROS

- More widely available
- Lower cost
- Easier to use

### • CONS

- Ambiguous and marginally actionable HOS data
  - Spatially averaged information over the entire protein population
  - Fails to inform on a residue-level
  - Examines a very limited number of specific moieties in the protein structure

### LOW RESOLUTION DATA EASIER TO PERFORM LOWER COST



### 3

### Overview of **HOS Approaches**

NMR, X-ray & CryoEM

Thermal shift assay, chromatography, circular dichroism (CD), microfluidic modulation spectroscopy (MMS), infrared (IR) spectroscopy, light scattering, surface plasmon resonance (SPR), isothermal titration calorimetry (ITC), etc.



#### **HIGH RESOLUTION DATA** HARDER TO PERFORM **HIGHER COST**



- PROS
  - Yields highest quality data

• CONS

- Demands deep expertise
- Demands highly trained operators
- Complex and lengthy workflows
- Very expensive equipment
- May use dangerous lasers or electron beam sources

LOW RESOLUTION DATA EASIER TO PERFORM LOWER COST





### Overview of HOS Approaches

NMR, X-ray & CryoEM

**MS-based** protein footprinting (HDX & HRPF)

Thermal shift assay, chromatography, circular dichroism (CD), microfluidic modulation spectroscopy (MMS), infrared (IR) spectroscopy, light scattering, surface plasmon resonance (SPR), isothermal titration calorimetry (ITC), etc.



#### **HIGH RESOLUTION DATA** HARDER TO PERFORM **HIGHER COST**

### **MID-RESOLUTION DATA:**

#### • PROS

- Not as difficult as high-resolution methods
- Less cost that high resolution methods
- Data quality in parity with high resolution methods

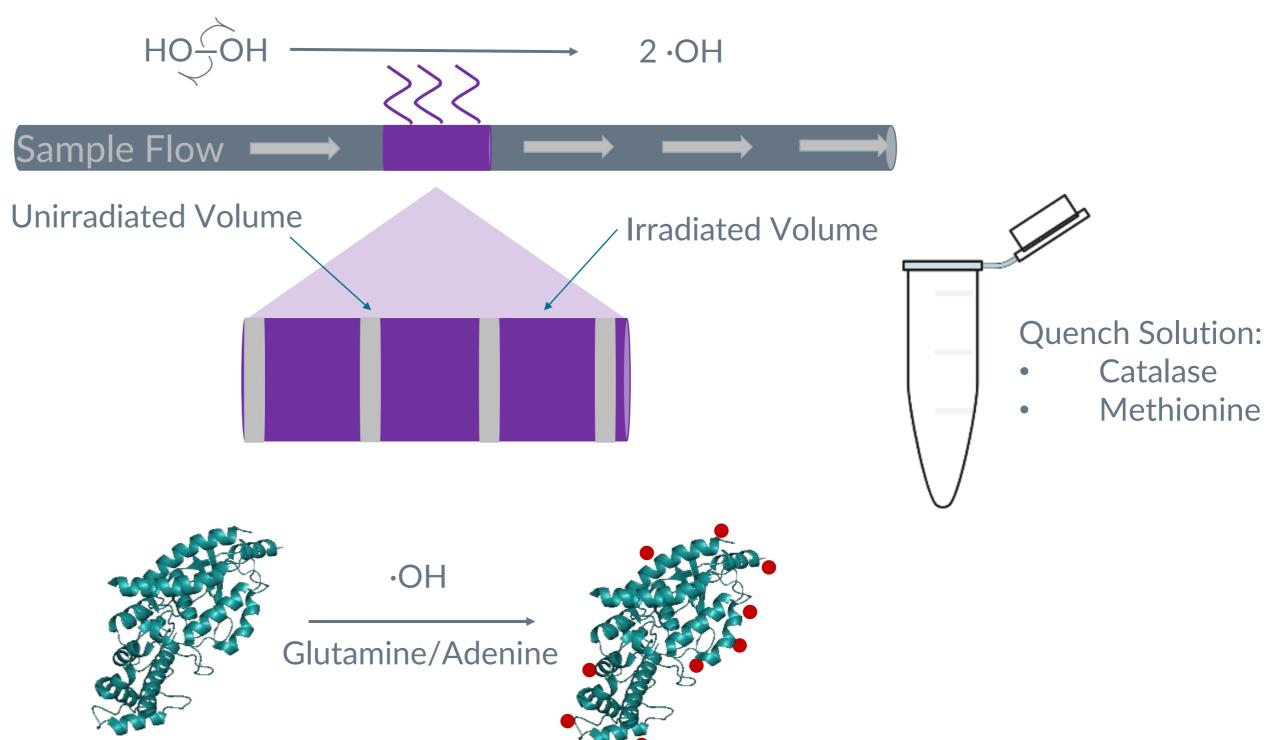
LOW RESOLUTION DATA EASIER TO PERFORM LOWER COST



### Fast Photochemical Oxidation of Protein (FPOP) HRPF HOS Analysis



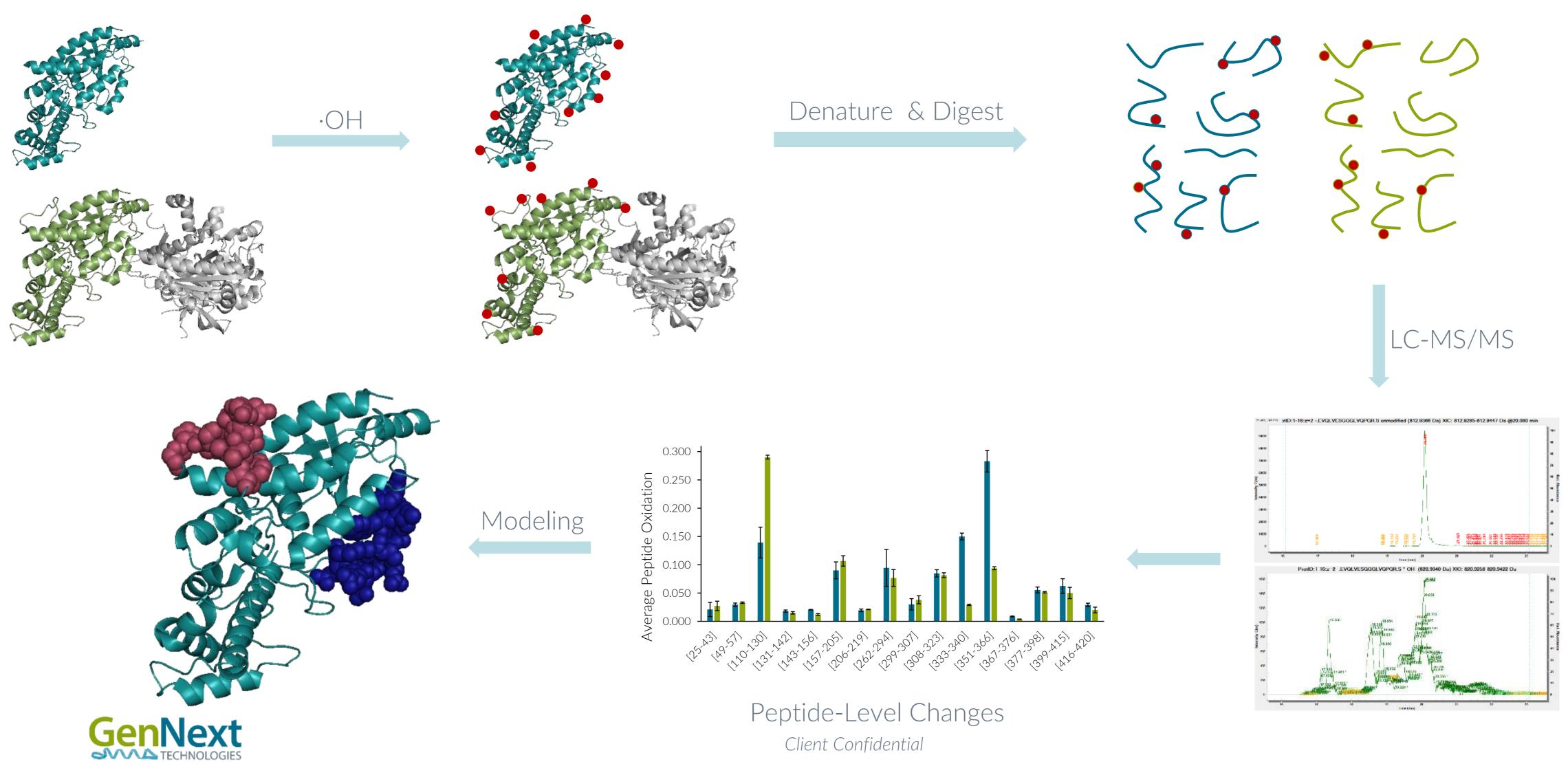
- OH generated by flash photolysis of hydrogen peroxide
- Modifies exposed side chains
- Measures protein topography at peptide to amino acid resolution







### Hydroxyl Radical Protein Footprinting Typical Workflow

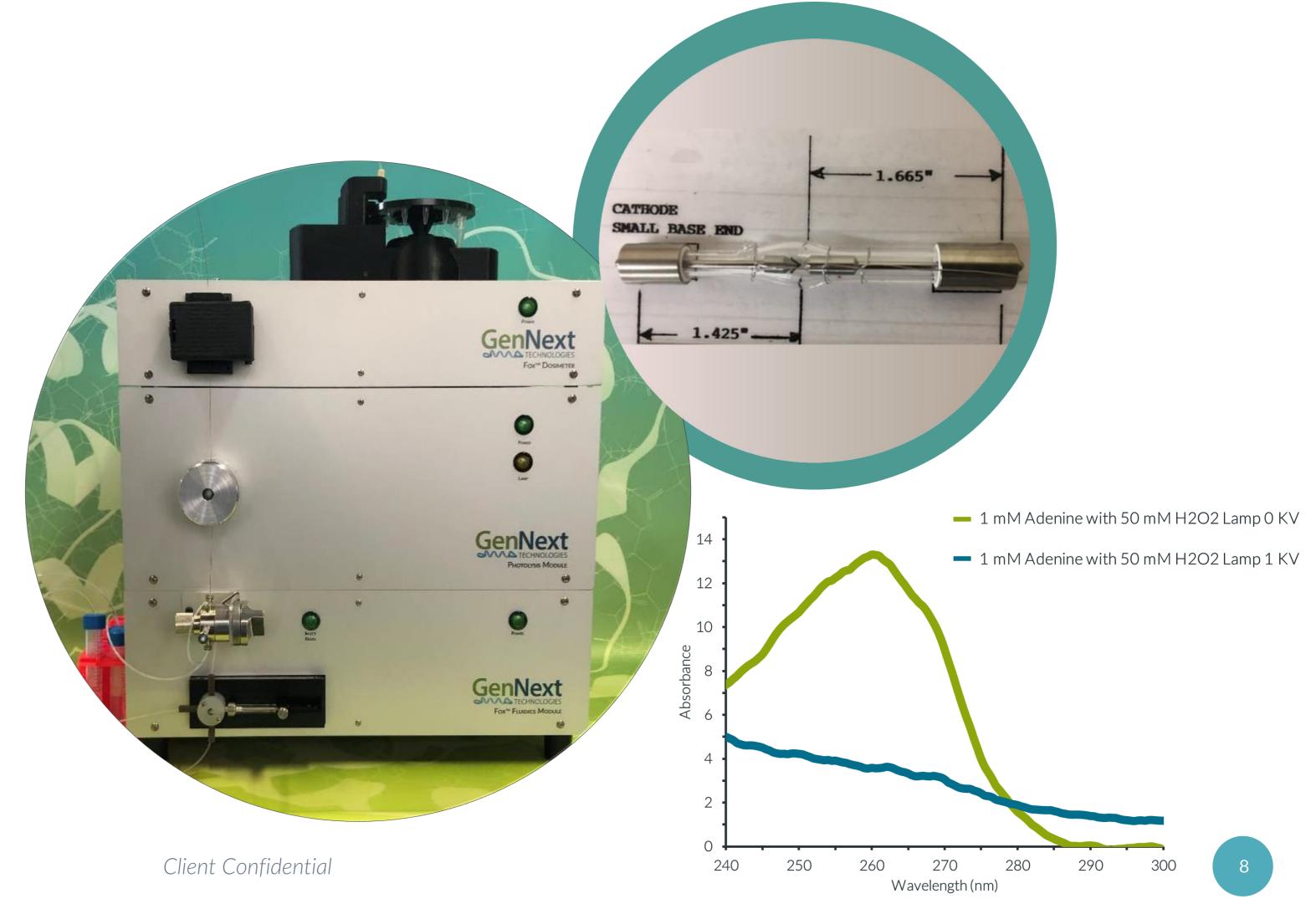




### Fox<sup>™</sup> Protein Footprinting System

Laser-free Delivery of Reproducible & Reliable Structural Biology Data

- Fox system replaces hazardous lasers with proprietary plasma lamp technology in a safe and easy-to-use benchtop package
- Dosimeter enables real-time adjustment of scavenging, providing confident and actionable results
- Automated product collector facilitates collection of properly labeled sample while disposing unwanted products and reagents to waste

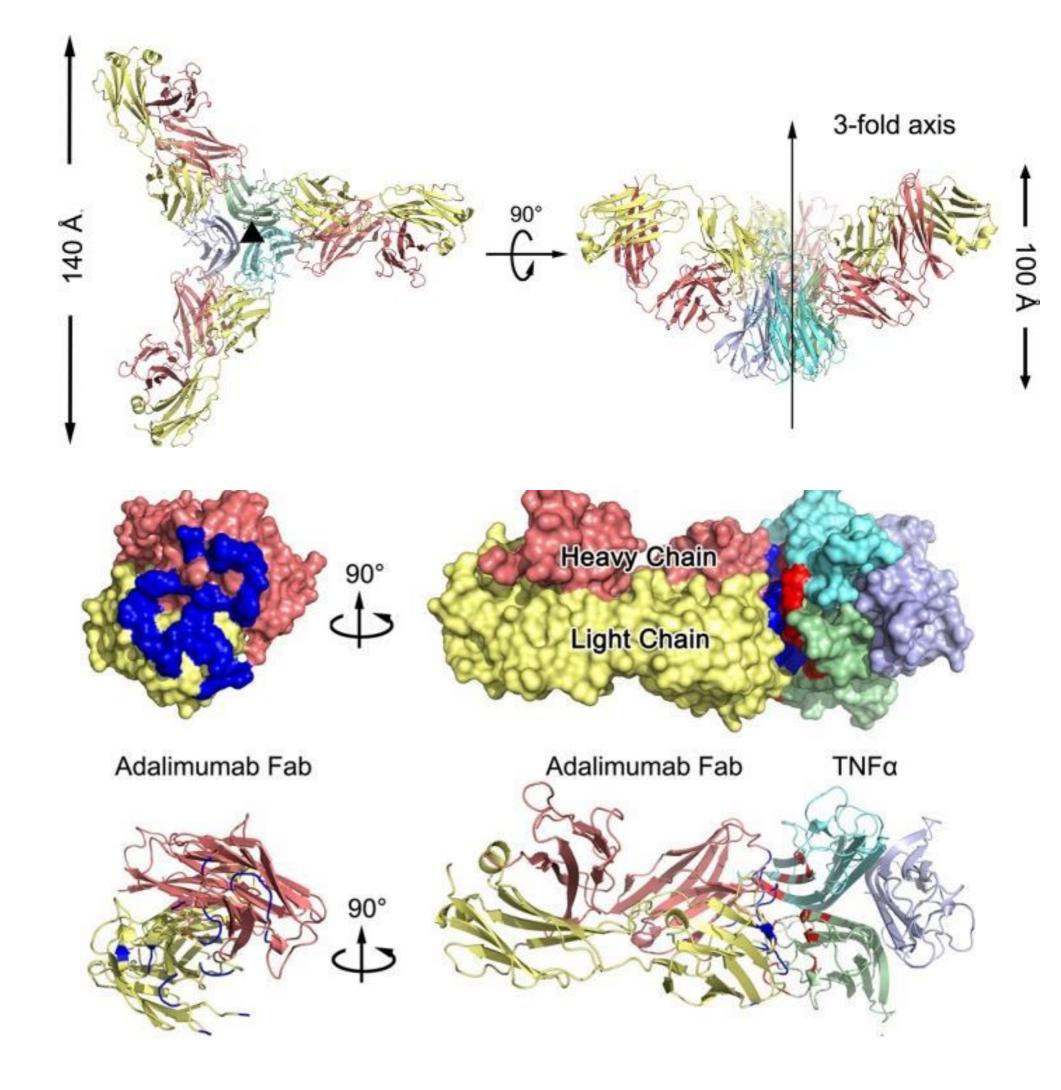




### **Epitope and Paratope** Mapping Fox<sup>™</sup> System Case Study: TNFα:Adalimumab

- Tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) is a pro-inflammatory cytokine
- Adalimumab (Humira) is a monoclonal antibody prescribed to treat inflammatory diseases
- TNFα:Adalimumab epitope has been well characterized with an available crystal structure
- Can Fox<sup>™</sup> Protein Footprinting accurately detect regions involved in the epitope and paratope?





J Biol Chem. 2013 Sep 20; 288(38): 27059-27067. PDB ID 3WD5



### Hydroxyl Radical Protein Footprinting Amino acids can stabilize proteins, minimizing intermolecular interactions

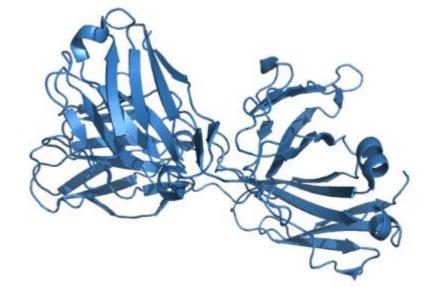
Condition 1: TNF $\alpha$  alone



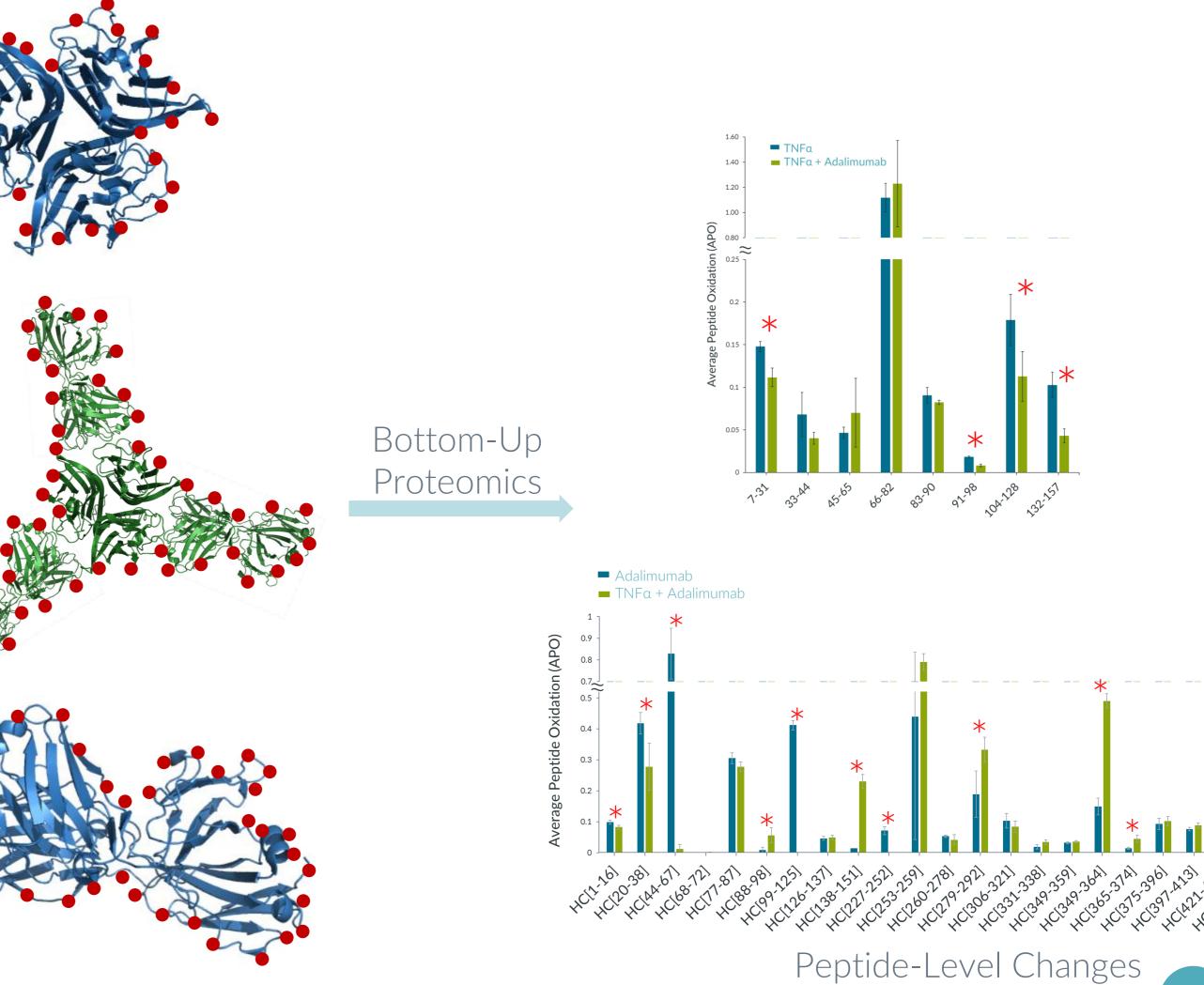
Condition 2: TNFa with Adalimumab



Condition 3: Adalimumab Alone

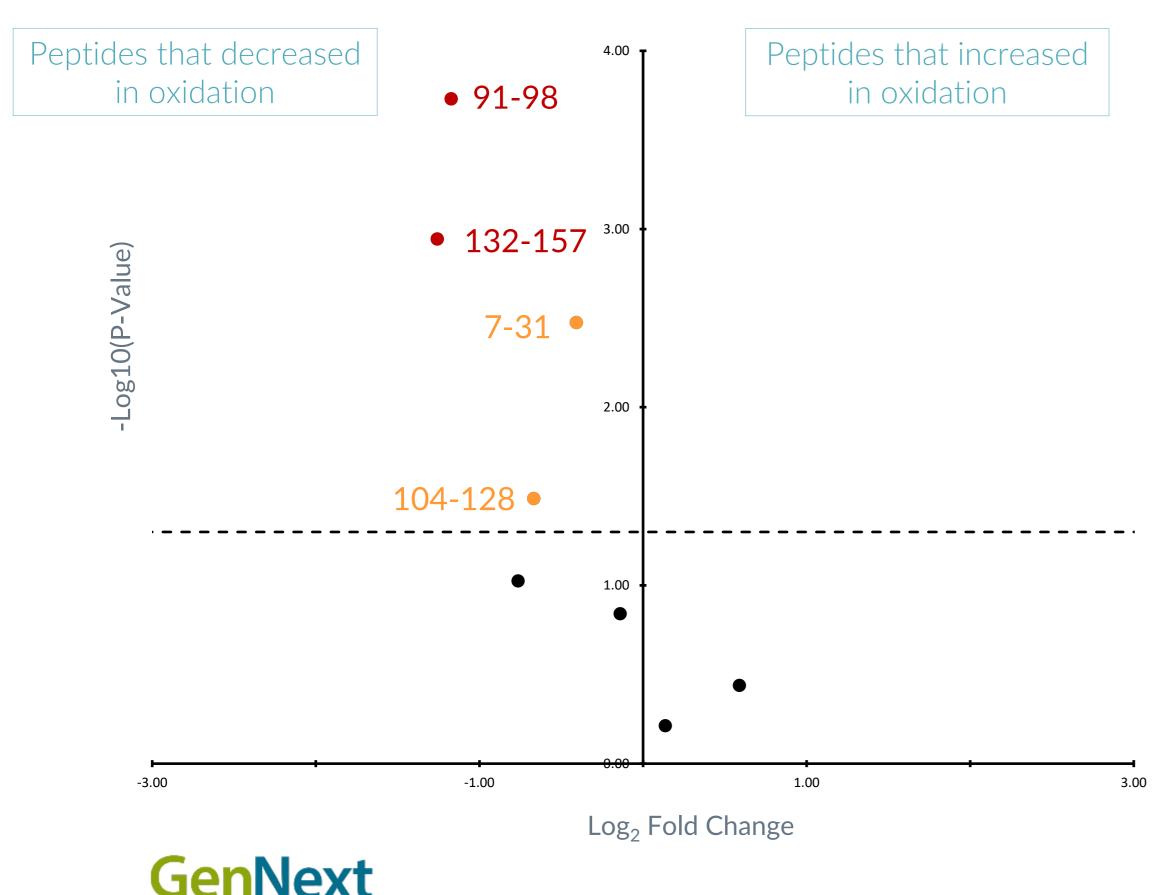


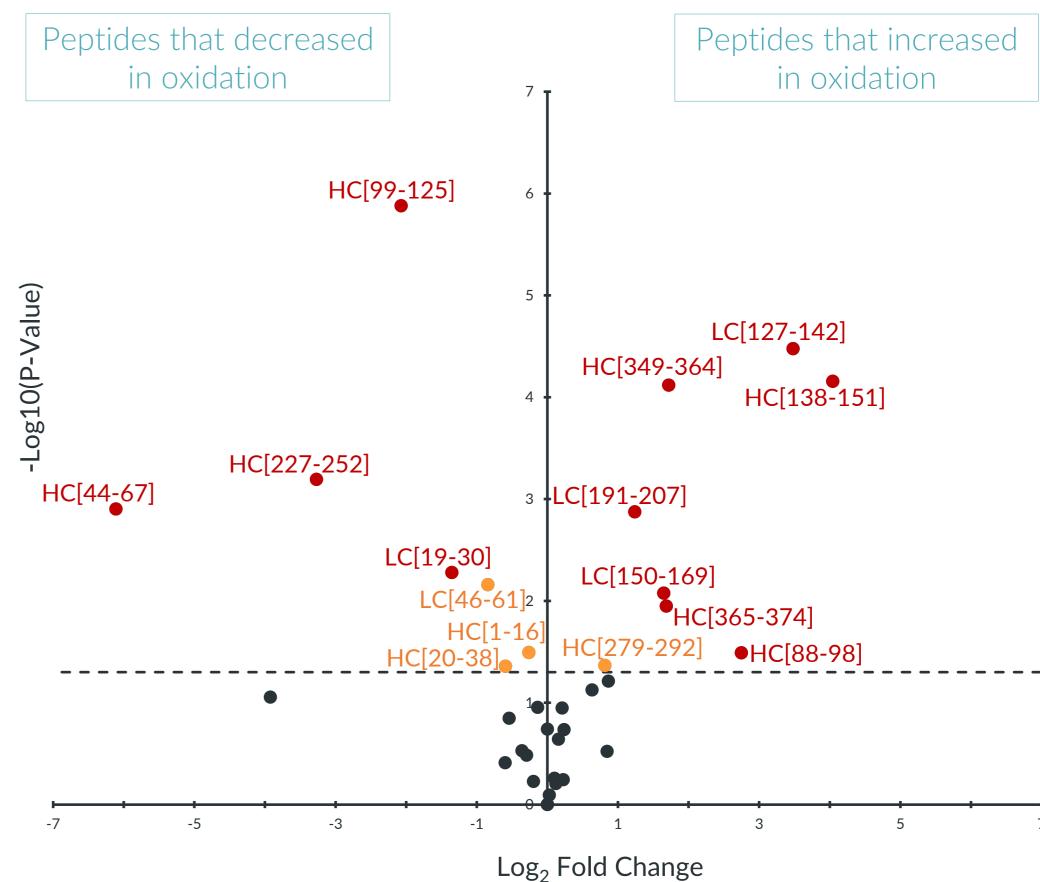






### **Epitope Mapping** Fox<sup>™</sup> Case Study: Histogram & Volcano Plot Pairwise Analysis

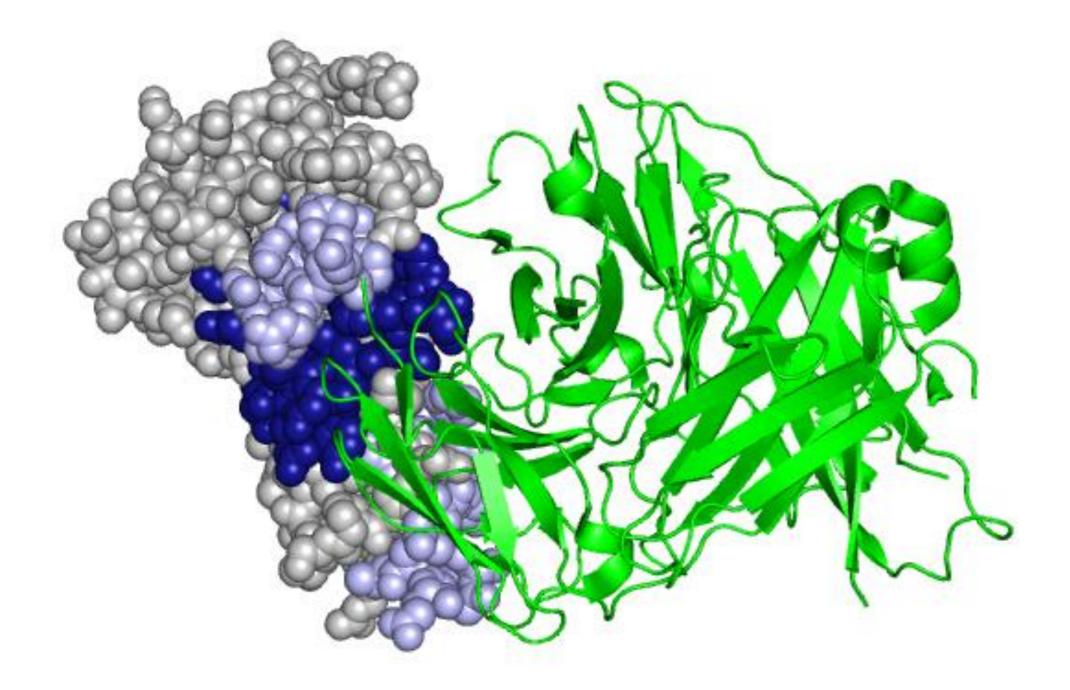






### TNFα – Adalimumab HRPF Characterized Epitope and Paratope

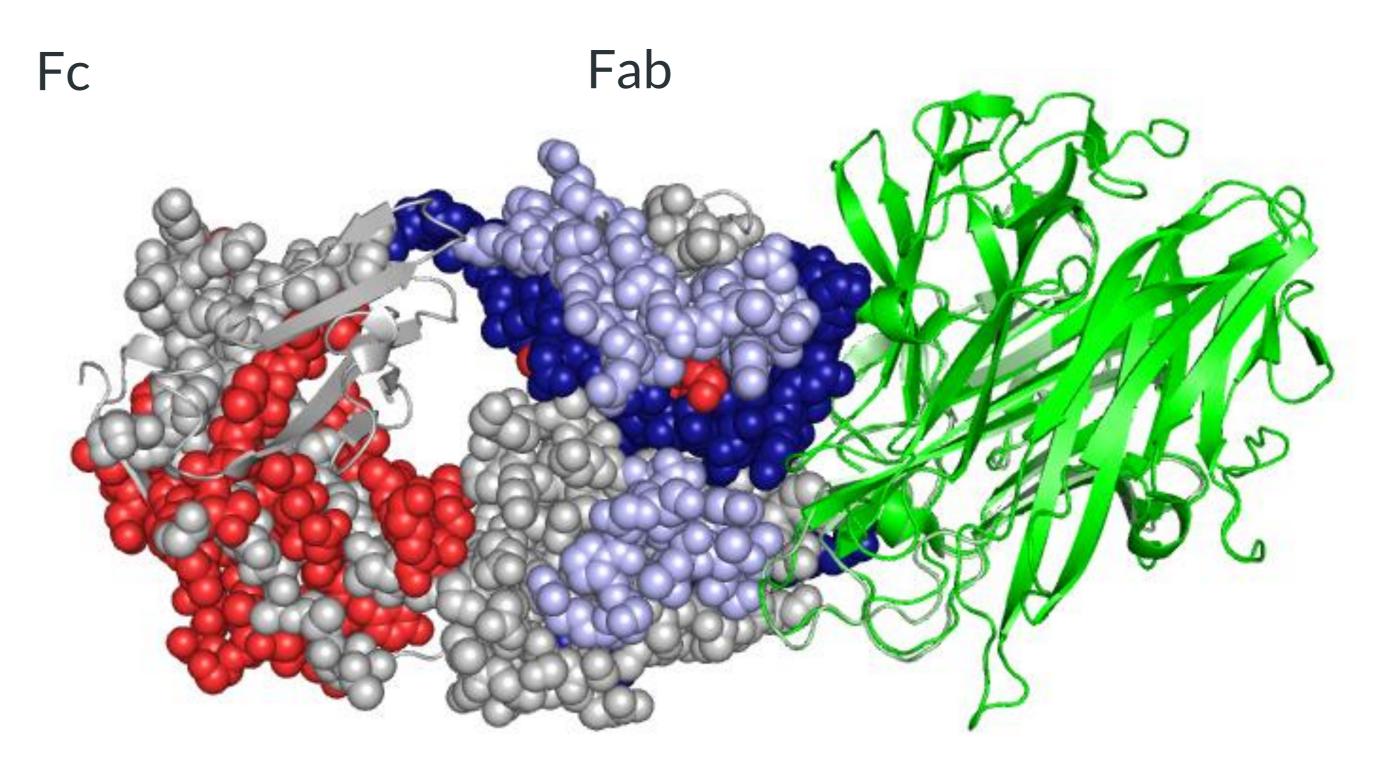
### TNFα – Adalimumab HRPF Characterized Epitope





Fox<sup>™</sup> System Case Study: Map Oxidation Changes on Crystal Structure

TNFα – Adalimumab HRPF Characterized Paratope





### Monoclonal Antibody Aggregation

#### Excipient effects on antibody aggregation

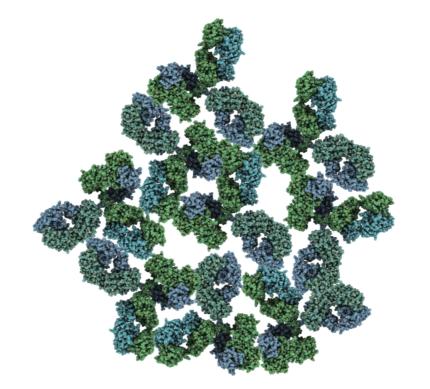
• Excipients, like amino acids, have been effective in stabilizing proteins that undergo intermolecular interactions at high concentrations

#### Fox<sup>™</sup> mAb aggregation domain analysis

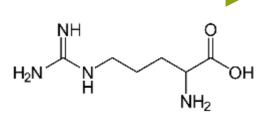
- Pairwise differential labeling experiment of aggregated and nascent Mab
- Aggregated surfaces are protected from labeling
- Labeling differences highlights interfacial domains for aggregation

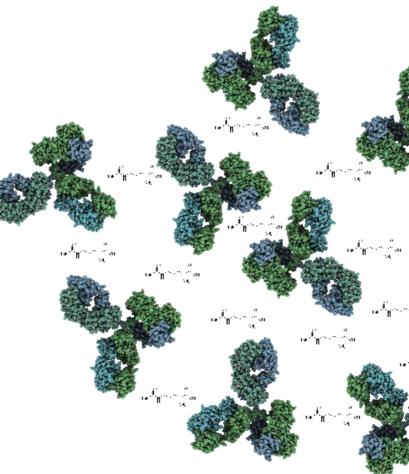


### Identification of Aggregate Domains







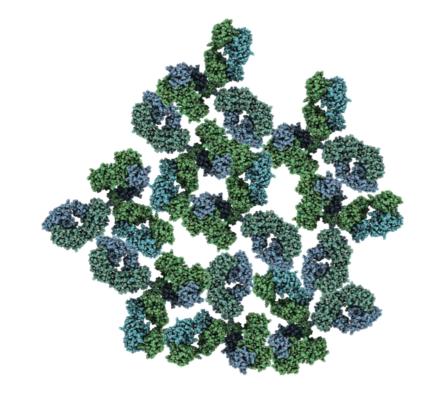






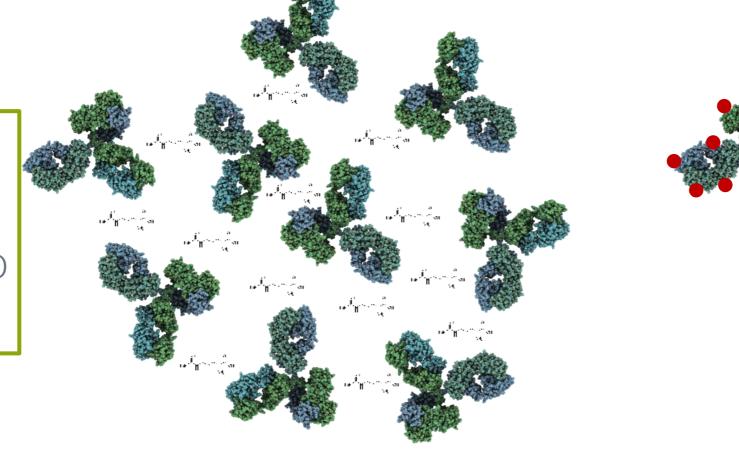
### **Hydroxyl Radical Protein Footprinting** Amino acids can stabilize proteins, minimizing intermolecular interactions

Condition 1: High concentration antibody without amino acid

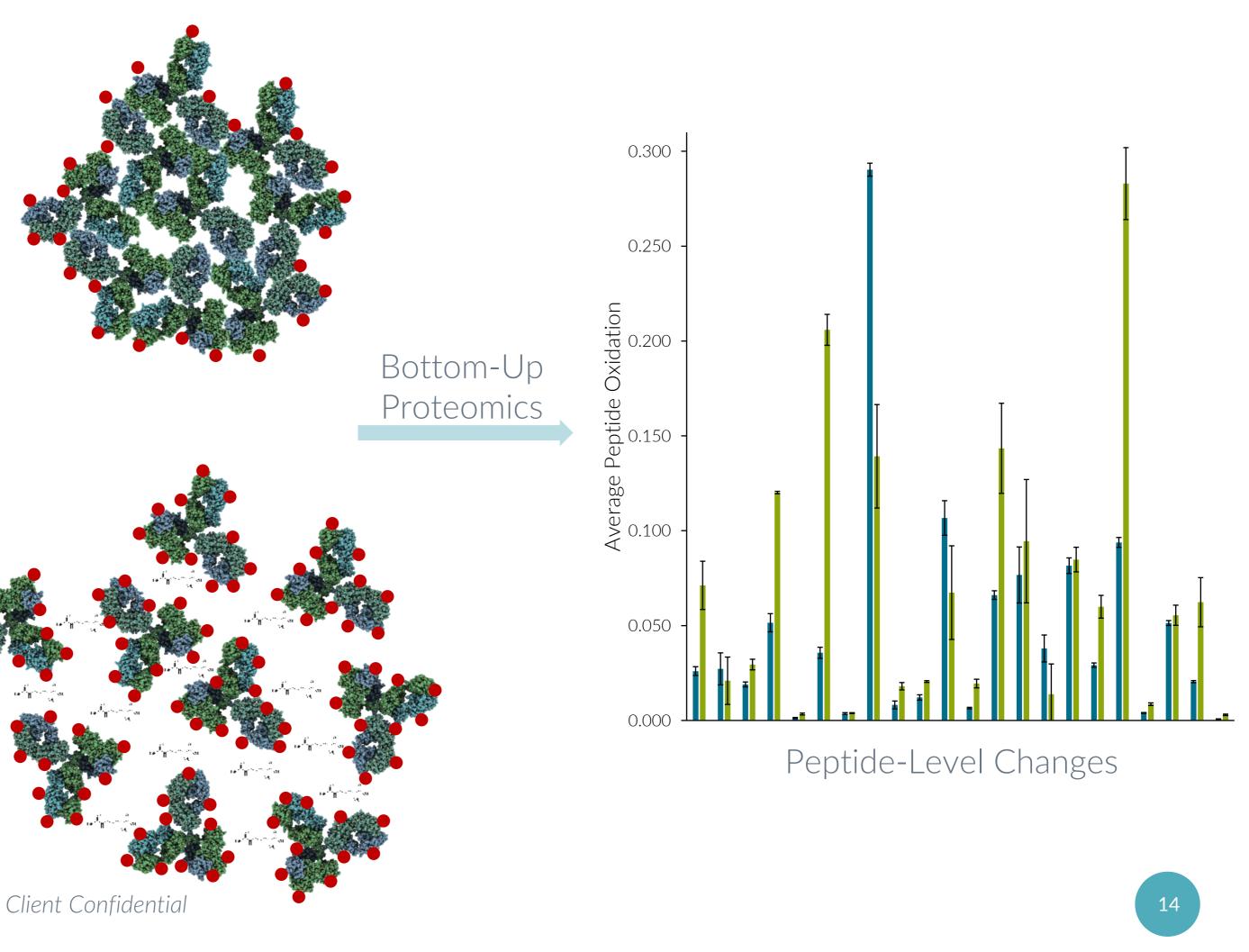


HRPF

Condition 2: High concentration antibody with amino acid



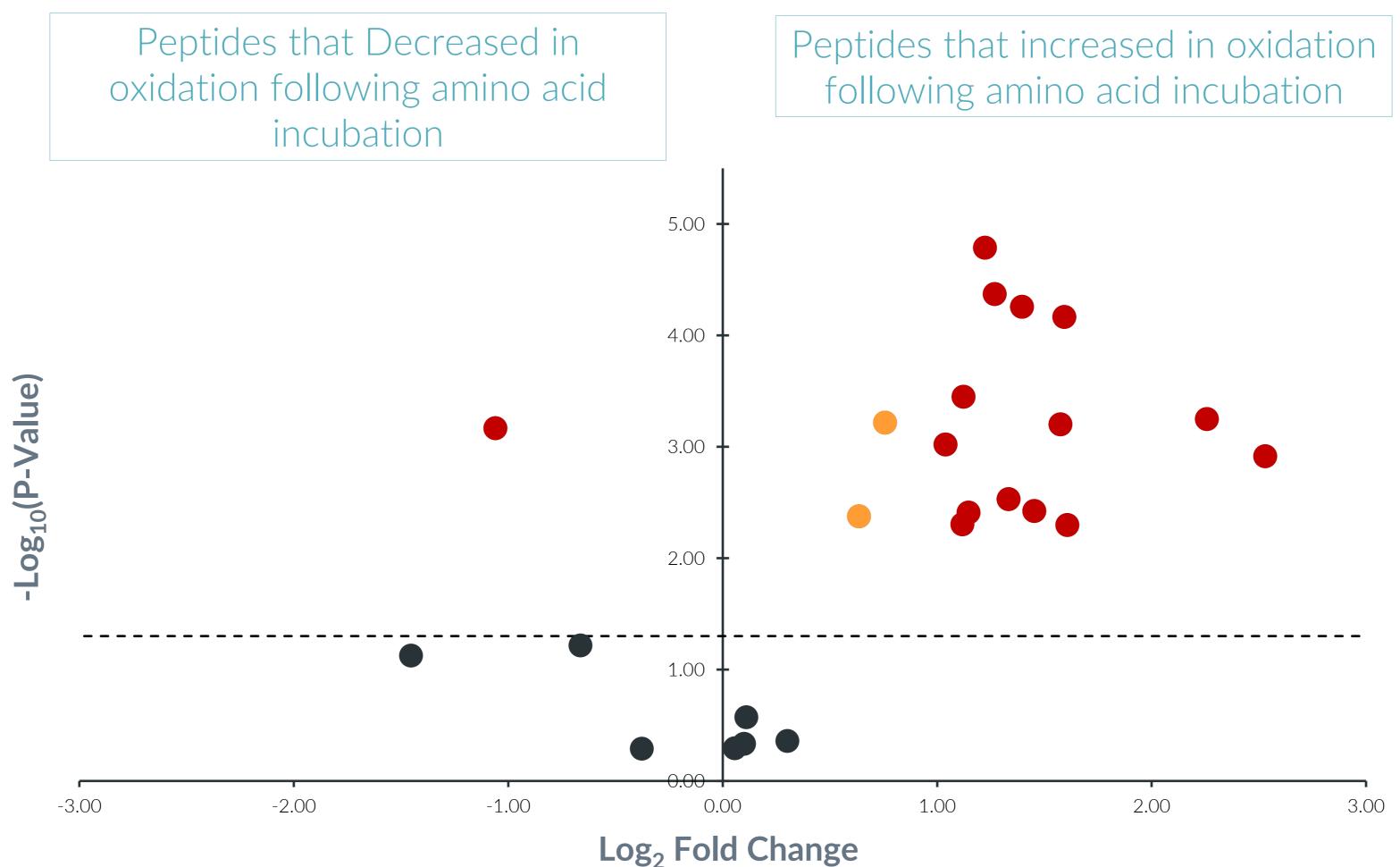




### Monoclonal Antibody Aggregation

Identifying peptides involved in intermolecular interactions

Peptides that Decreased in incubation







## **Protein Footprinting** Webinar Series

#### Introduction to Protein Footprinting

 Professors Mark Chance of Case Western Reserve University and Joshua Sharp of the University of Mississippi

#### **Protein Footprinting Applications in Structural Biology**

• Professor Michael Gross of the Washington University in St. Louis

#### Fast Photochemical Oxidation of Proteins (FPOP) HRPF

Professor Joshua Sharp of the University of Mississippi 







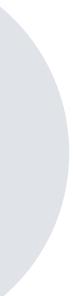
#### All Available **On-Demand at** www.gnxtech.com/ webinar-series

#### HRPF Data Processing and Higher Order Structural Analysis

Professor Joshua Sharp of the University of Mississippi 

#### In-cell and in vivo FPOP

• Professor Lisa Jones of the University of Maryland Baltimore







# Thank You for Your Participation!

To learn more, please visit our virtual booth.

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