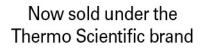
Monoclonal Antibody Characterization

Achieving Higher Throughput and Productivity







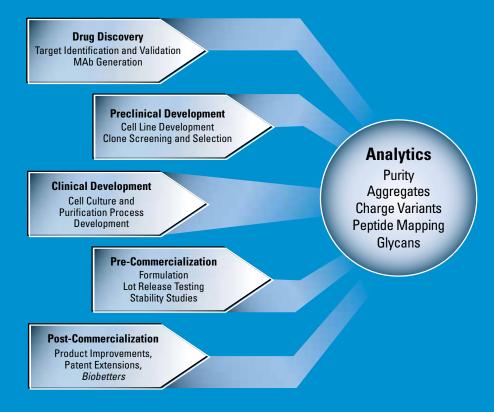


Dionex Solutions to Accelerate Monoclonal Antibody R&D and Characterization

The throughput and productivity challenge

- Increasing number of MAb candidates entering the clinical pipeline
- Advances in automation in upstream processes like cell culture and purification process development, and formulation screening
- Strict Quality-by-Design (QbD) guideline requiring enhanced antibody characterization

All result in a large increase in sample requests for characterization.



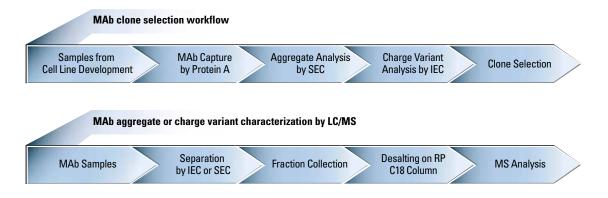
The analytical solutions

Dionex provides best-in-class analytical solutions for MAb therapeutics, improving throughput and productivity, thus reducing time to market.

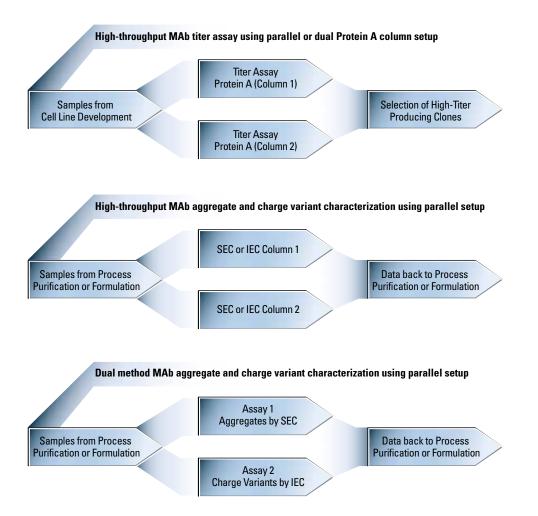
- Best-in-class columns for high-resolution and high-throughput characterization using ion-exchange chromatography (IEC) and size-exclusion chromatography (SEC)
- MAb characterization platforms to increase sample throughput and streamline multistep workflows
- Powerful and flexible Chromeleon[®] Chromatography Data System software
- Innovative UltiMate[®] 3000 RSLC system for UHPLC solutions for fast MAb characterization and peptide mapping
- High performance MAb glycan analysis

Whatever Your Workflow, You Can Achieve Higher Throughput and Productivity

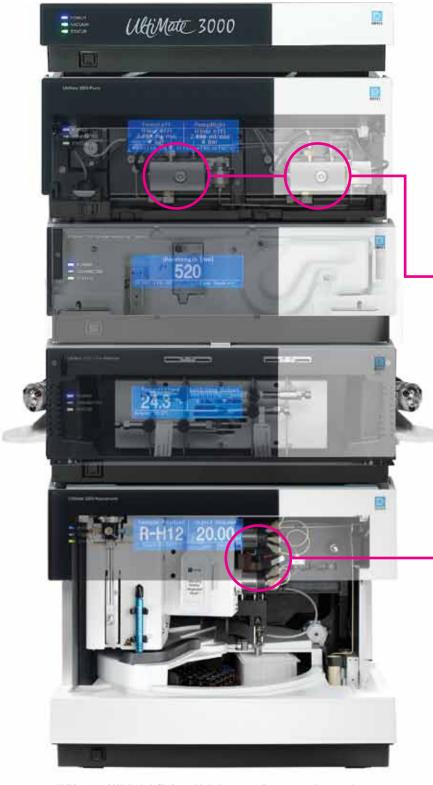
Multistep MAb characterization workflows



High-throughput MAb characterization workflows



The Dionex MAb Characterization Platform-



UltiMate 3000 MAb Analysis Platform with dual ternary gradient pumps and autosampler with integrated fraction collector.

UltiMate 3000 MAb fully biocompatible platform

- Titanium pumps; PEEK[™] fluidics, valves and injection needle
- Full compatibility with all biological buffers
- Prevents iron poisoning of columns
- Maintains protein modifications
- Less system corrosion and maintenance

Dual gradient pump design: 2-in-1 system

- Multistep Automation: allows automation of two or more methods like Protein A capture, SEC aggregate, and IEC charge variant analysis
- Tandem Analyses: shortens run times by utilizing the power of off-line column regeneration
- Parallel LC: double throughput for productivity, cost, and space saving

Autosampler with fraction collection and re-injection

- Dual valve design allows injection, fraction collection, and re-injection
- Optimized for automated workflows like:
 - Automated multistep workflow automation
 - Protein purification
 - Sample fractionation and desalting prior to MS detection
 - Sample derivatization like digestion or neutralization between multistep separations

For Your Throughput and Productivity Needs

Chromeleon Chromatography Data System (CDS) software

Features of Chromeleon CDS software:

- Automated peak integration
- · Automated control of fraction collection into autosampler
- · Automated method template and sequence generation for multi-dimensional workflows
- Automatic rejection of samples based on set criteria e.g., automated rejection of clones with low antibody titer
- Dilutions or variable injection volumes for the next steps
- One-click report generation
- Validation report templates and sequences for increased automation of method validation

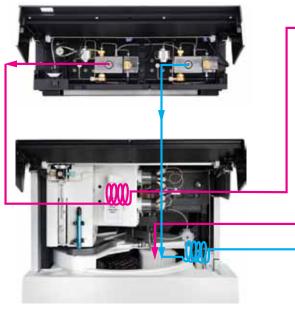


Fully integrated solutions for R&D, analytical method development and QA/QC

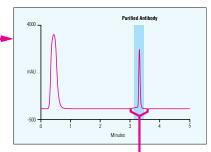
Boost Productivity with Automated Multistep MAb Capture and Characterization

Automate your multistep MAb capture and characterization reduce hands-on time and increase productivity

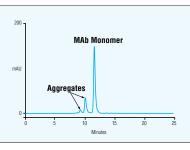
Dual Ternary Gradient Pump. 2-in-1 Pump Design.



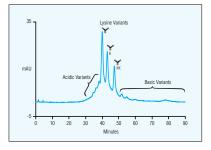
Step 1: Protein A capture of lgG (MAb) and automated fraction collection.



Step 2: SEC aggregate analysis.

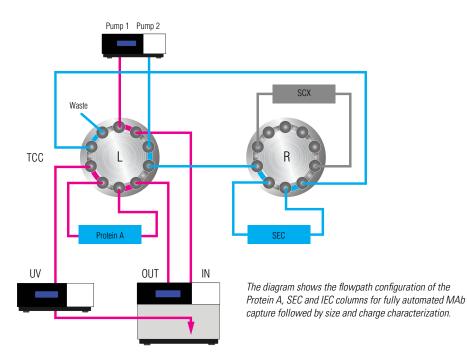


Step 3: IEC charge variant analysis.



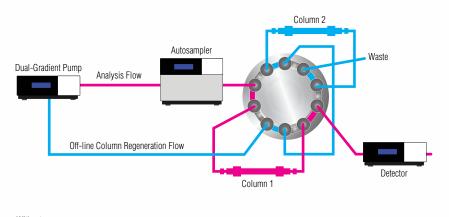
Autosampler with Injection, Fraction Collection, and Re-Injection. Two sampling technologies in one.

The Multistep MAb Analysis Platform allows the purification and analysis of hundreds of MAb samples. Cell culture fluid samples containing antibodies are injected onto a Protein A column for antibody recovery, then the purified antibody samples are automatically injected, first onto the SEC column, and then onto the IEC column—fully automated.



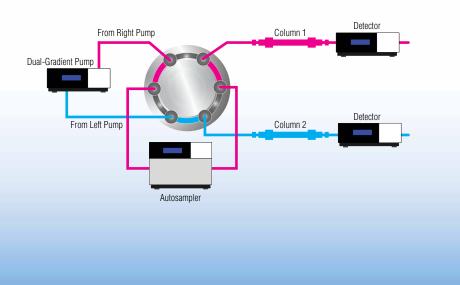
Accelerate Product Development by Increasing Sample Throughput

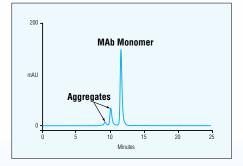
Analyze over a 1000 MAb samples a day by utilizing the power of off-line column regeneration

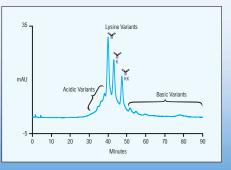




With the same samples perform multiple separation steps including SEC and IEC in a parallel configuration







Monoclonal Antibody Characterization Using

Charge Characterization with IEC

- Glycosylation
- Sialylation
- C-Terminal Lysine
- Deamidation
- Glutamine cyclization
- Maleuric acid adduct
- Oxidation
- Cysteinylation
- Disulfide related
- Succinimide
- Isomerization

Hydrophobicity Characterization with HIC

- Isomerization
- Succinimide
- Oxidation
- Amidation
- Aggregation
- Clipping

Column

 $\label{eq:proPac} \begin{array}{l} \mbox{ProPac}^{\mbox{\tiny (B)}} \mbox{ WCX-10} \mbox{ (} 4 \times 250 \mbox{ mm} \mbox{)} \\ \mbox{ProPac} \mbox{ WCX-10HT} \mbox{ (} 4 \times 50 \mbox{ mm} \mbox{)} \end{array}$

NEW!

 $\label{eq:masses} \begin{array}{l} \mathsf{MAbPac}^{\texttt{M}} \ \mathsf{SCX-10} \ (4 \times 250 \ \mathsf{mm}) \\ \mathsf{MAbPac} \ \mathsf{SCX-10} \ (4 \times 150 \ \mathsf{mm}) \\ \mathsf{MAbPac} \ \mathsf{SCX-10HT} \ (4 \times 50 \ \mathsf{mm}) \\ \mathsf{MAbPac} \ \mathsf{SCX-10} \ \mathsf{for} \ \mathsf{LC/MS} \ (2 \times 250 \ \mathsf{mm}) \end{array}$

Column

ProPac HIC (4.6×250 mm) ProPac HIC (4.6×100 mm) ProPac HIC-10 (2.1×100 mm) ProPac HIC-10 (7.8×75 mm)

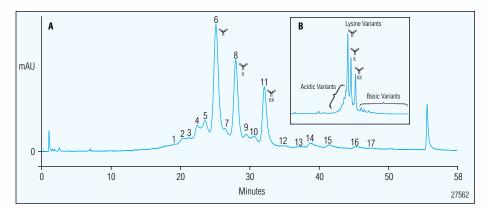
Size Characterization with SEC

- Monomers, aggregates, and fragments
- Under non-denaturing conditions using both high- and low-salt mobile phases and volatile eluents for LC/MS

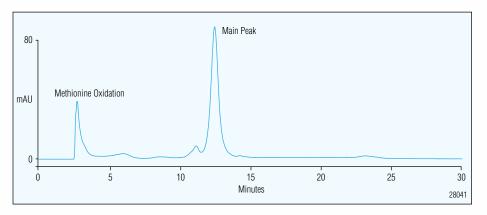
Column

MAbPac SEC-1, 5 μ m, 300 Å (4.0 × 300 mm) MAbPac SEC-1, 5 μ m, 300 Å (4.0 × 150 mm)

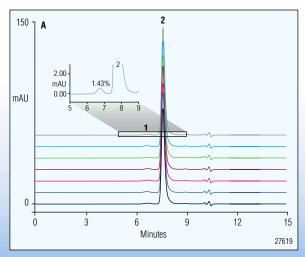
High-Resolution Columns



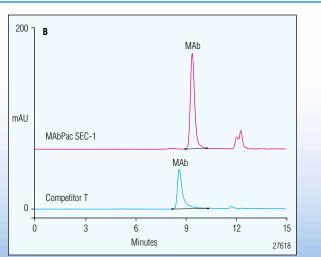
A. MAbPac SCX-10 column is the next generation IEC column for MAb charge characterization from Dionex. B. ProPac WCX-10 column (inset) is the industry gold standard for charge variant characterization.



ProPac HIC-10 column - Separation of populations of MAb variants using hydrophobic interaction chromatography (e.g., methionine oxidation monitoring).



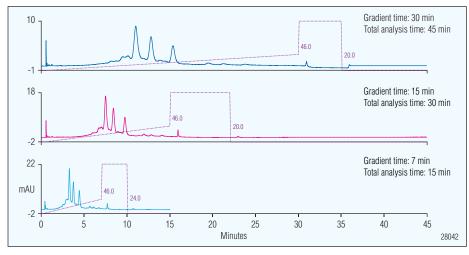




B. MAb analysis in volatile buffer for LC/MS—MAb Pac SEC-1 vs the competitor's column.

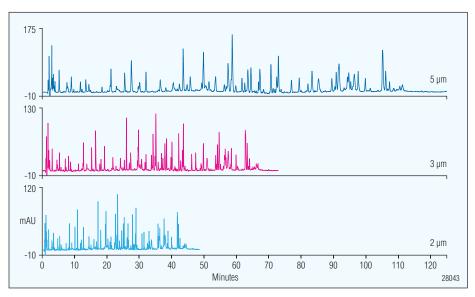
Fast MAb Characterization and Peptide Mapping

Method acceleration using the new MAbPac SCX-10 column, 4 × 150 mm



Elution profiles of a MAb sample from a 4×150 mm MAbPac SCX-10 column with different gradient elution times. The gradient time and total analysis time is indicated in the chromatogram.

Fast MAb peptide mapping—application of UHPLC and reduced particle size



Performing peptide mapping with smaller particle columns allows the user to achieve identical resolution in less time. Here a method transfer was performed using 2.1 \times 100 mm columns packed with 5, 3, and 2 µm particles respectively on the UltiMate 3000 Rapid Separation LC (RSLC).

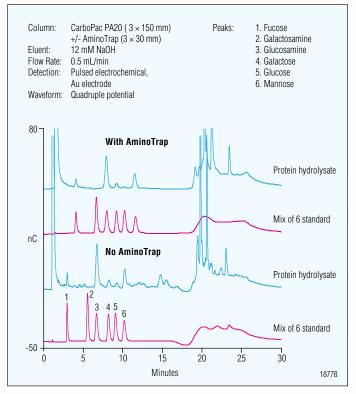
The Preferred and Proven Platform for MAb Glycan Characterization

MAb glycosylation-why measure it?

- Increasing relevance—biosimilars
- Glycosylation can affect:
 - Biological activity
 - Pharmacokinetics and clearance in vivo
 - Stability
 - Immunogenicity
- · Analysis of glycosylation is important to:
 - Meet regulatory requirement
 - Ensure product consistency
 - Develop new generation drugs with modified glycosylation

Dionex solution

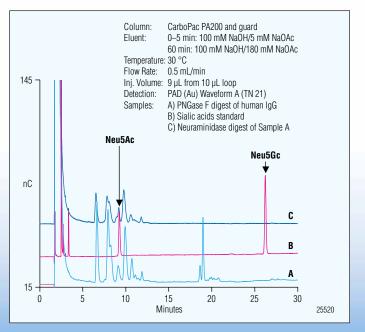
- High Performance Anion-Exchange with Pulsed Amperometric Detection (HPAE-PAD)
 - The workhorse in carbohydrate analysis
 - High Sensitivity-0.1 to 1 pmole detection limits
 - Label-Free—No derivatization necessary



Profiling MAb hydrolysate on the CarboPac[®] PA20 column, with and without an AminoTrap[™] precolumn.



ICS-5000—*New capillary system for carbohydrate analysis.*



Ionitoring release of sialic acids from human IgG N-linked oligosaccharides by HPAE-PAD.

Selected Peer Reviewed Publications

- Lyubarskaya, Y.; Houde, D.; Woodard, J. Murphy, D.; Mhatre, R. Analysis of Recombinant Monoclonal Antibody Isoforms by Electrospray Ionization Mass Spectrometry as a Strategy for Streamlining Characterization of Recombinant Monoclonal Antibody Charge Heterogeneity. *Anal. Biochem.* 2006, 348, 24–39.
- Vlasak, J.; Ionescu, R. Heterogeneity of Monoclonal Antibodies Revealed by Charge-Sensitive Methods. *Curr. Pharm. Biotechnol.* 2008, *9*, 468–481.
- Valliere-Douglass, J.; Wallace, A.; Balland, A. Separation of Populations of Antibody Variants by Fine Tuning of Hydrophobic Interaction Chromatography Operating Conditions. *J. Chromatogr.*, A 2008, 1214, 81–89.
- 4. Decrop, W.; Gendeh, G.; Swart, R. Development of an Automated Method for Monoclonal Antibodies Purification and Analysis. *Chromatography Today*, Jun 2009, 8–10.
- Farnan, D.; Moreno, G.T. Multiproduct High-Resolution Monoclonal Antibody Charge Variant Separations by pH Gradient Ion-Exchange Chromatography. *Anal. Chem.* 2009, *81*(21), 8846–8857.
- Farnan, D.; Moreno, D.T.; Stults, J.; Becker, A.; Tremintin, G.; van Gils, M. Interlaced Size Exclusion Liquid Chromatography of Monoclonal Antibodies. *J. Chromatogr., A* 2009, *1216*(51), 8904–9.
- Rea, J.C.; Moreno, G.T.; Lou, Y.; Parikh, R.; Farnan, D. High-Throughput Multi-Product Liquid Chromatography for Characterization of Monoclonal Antibodies. *BioPharm International* 2010, 23, 44–51.
- 8. Grey, C.; Edebrink, P.; Krook, M.; Jacobsson, S.P. Development of a High Performance Anion Exchange Chromatography Analysis for Mapping of Oligosaccharides. *J. Chromatogr., B Anal. Technol. Biomed. Life Sci.* **2009**, *877*, 1827–1832.



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