

BioLC columns

Connected chromatography solutions

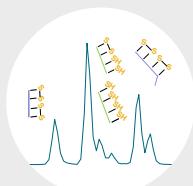
BioLC columns and accessories

thermo scientific

Introduction

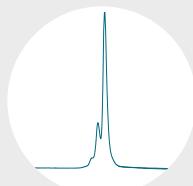
Your complete tool kit

Thermo Fisher Scientific has innovative Thermo Scientific™ BioLC columns for each step of your therapeutic protein characterization, no matter how challenging your separation. Here is just one example, a fully characterized model sample of pertuzumab. Discover our full range in this catalogue.



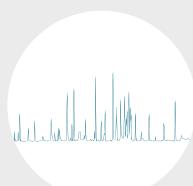
Intact or subunit analysis

Thermo Scientific™ MAbPac™ RP columns are ideal for intact and subunit analysis by MS or UV detection. The polymeric packing material offers column longevity, high resolution and the wide pores to allow for low carryover profiling of your sample.



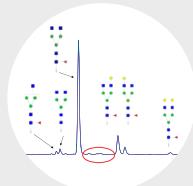
Oxidation monitoring

Deduce protein folding errors or charge-neutral amino acid modifications with the Thermo Scientific™ MAbPac™ HIC-20 hydrophobic interaction column. Our range of innovative HIC chemistries deliver native separations not seen on other columns.



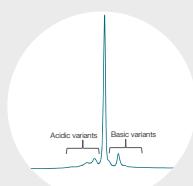
Peptide mapping

Experience reproducible peptide mapping and quantitation. The combination of rapid digestion from the Thermo Scientific™ SMART Digest kit and separation with the high resolution Thermo Scientific™ Hypersil™ GOLD column delivers outstanding, reproducible and efficient peptide mapping separations.



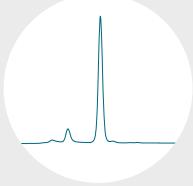
Released glycan analysis

Fully characterize your released N-glycans with the Thermo Scientific™ Accucore™ 150 Amide-HILIC column. This solid core column offers high resolution, durability, and the ability to run separations at lower temperatures to reveal the complete glycan profile.



Charge variant analysis

For charge variant analysis by LC-UV or LC-MS/MS Thermo Scientific™ ProPac™ 3R SCX and Thermo Scientific™ ProPac™ 3R SAX columns deliver outstanding resolution on a highly robust, reproducible and high-resolution platform. Combine ProPac 3R SCX columns with our proprietary Thermo Scientific™ CX-1 gradient buffers formulations to enable fast, robust and reproducible pH gradients that are simple to optimize and easily automated - without the need for time-consuming mobile phase adjustments.

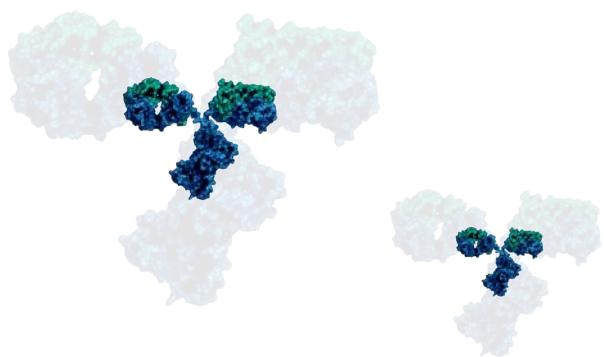


Aggregate analysis

Thermo Scientific™ MAbPac™ SEC-1 offers excellent size exclusion separation even under challenging conditions for aggregate analysis. Compatible with mass spectrometry for native LC-MS/MS workflows.

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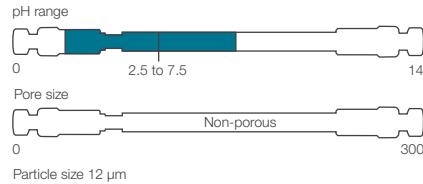
BioLC column selection quick guide

| Target applications | Column type | Mode of analysis | Recommended column | Particle size (µm) | Pore size (Å) | pH range | Maximum backpressure (psi) | Solvent compatibility |
|------------------------------------|---|-------------------------|--------------------------|--------------------|-----------------------|----------------------------------|---|--|
| Affinity | Affinity columns | Affinity | MAbPac Protein A | 12 | Non-porous | 2.5-7.5 | 1,000 | — |
| Intact analysis by HIC | Silica-based hydrophobic interaction chromatography columns | Hydrophobic interaction | MAbPac HIC-10 | 5 | 1,000 | 2-8 | 4.6 × 100 mm = 6,000 | Compatible with organic solvents and aqueous mobile phases |
| | | | MAbPac HIC-20 | 5 | 1,000 | 2-9 | 4.6 × 250 mm = 8,000 | |
| | | | MAbPac HIC Butyl | 5 | Non-porous | 2-12 | 4,000 | Compatible with up to 50% organic solvents |
| Released glycan analysis | Silica based, mixed-mode columns | Mixed-mode | GlycanPac AXH-1 | 1.9 | 175 | 2-8 | 10,000 | 0 – 90% aqueous buffer; 10 – 100% acetonitrile or alcohols |
| | | | | 3 | 120 | 2-8 | 6,000 | |
| | | | GlycanPac AXR-1 | 1.9 | 175 | 2-8 | 10,000 | Compatible with 0 – 100% aqueous and common HPLC solvents (except acetone) |
| Aggregate fragment analysis | Silica-based size exclusion chromatography phases | Size exclusion | Accucore 150 Amide HILIC | 2.6 | 150 | 2-8 | 14,500 | — |
| | Polymeric reversed-phase columns | Reversed-phase | MAbPac SEC-1 | 5 | 300 | 2.5-7.5 | 1,000 for 300 mm 600 for 150 mm | 100% organic solvents |
| | Polymeric reversed-phase columns | Reversed-phase | MAbPac RP | 4 | 1,500 | 2.1, 3.0 mm (0-14) 1 mm (1-7) | 4,000 | Up to 100% ACN, IPA, MeOH |
| Intact and subunit analysis | Polymeric reversed-phase columns | Reversed-phase | ProSwift RP-1S | | | | 2,800 | — |
| | | | ProSwift RP-2H | | | | 2,800 | |
| | | | ProSwift RP-3U | Monolith | Monolith | 1-14 | 2,800 | Most common organic solvents |
| Charge variant analysis | Monolithic ion-exchange columns | Ion-exchange | ProSwift RP-4H | | | | 1 × 50 mm = 2,000 2 × 250 mm = 3,000 | |
| | | | ProPac 3R SCX | | | | 4,500 | — |
| | | | ProPac 3R SAX | | | | 4,500 | — |
| Peptide mapping | Silica based, reversed-phase columns | Reversed-phase | MAbPac SCX-10RS | 5 | | | 7,000 | — |
| | | | MAbPac SCX-10 | 5, 10 | Non-porous | 2-12* | 3,000 for 10 µm 5,000 for 5 µm | *Please consult column manual |
| | | | ProPac SAX-10 | 10 | | | 3,000 | *Please consult column manual |
| Nucleic acids and oligonucleotides | Polymeric ion-exchange columns | Ion-exchange | ProPac Elite WCX | 5 | | | 4,500 | *Please consult column manual |
| | | | Hypersil GOLD C18 | 1.9 | 175 | 1-11 | 18,130 | — |
| | | | | 3 | 175 | | 5,800 | — |
| | Polymeric reversed-phase | Reversed-phase | Acclaim 120 C18 | 2.2 | 120 | | — | |
| | | | | 3 | 120 | 2-8 | Various | — |
| | | | | 5 | 120 | | — | |
| | Polymeric ion-exchange columns | Ion-exchange | DNAPac PA200 | 8 | Non-porous | 2.5-12.5* | 4,000 | *Please consult column manual |
| | | | DNAPac PA200RS | 4 | Non-porous | 2.5-12.5* | 10,000 | — |
| | | | DNASwift SAX-1S | Monolith | Monolith | 3-14* | 1,500 | *Please consult column manual |
| | Polymeric reversed-phase | Reversed-phase | DNAPac RP | 4 | Proprietary wide pore | 0-14 | 4,000 | — |

Affinity columns

Providing fast, accurate titer analysis of monoclonal antibodies in harvest cell cultures, the nonporous, polymeric Thermo Scientific™ MAbPac™ Protein A HPLC Column delivers reproducible, highly efficient separations.

MAbPac Protein A column

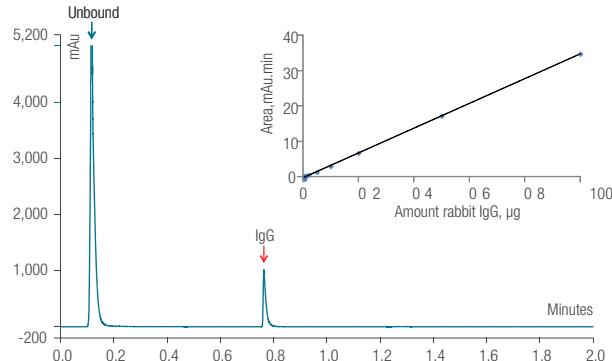


Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | MAbPac Protein A: A novel affinity Protein A column |
| | Learn more | thermofisher.com/biolc |

Harvest cell culture titer analysis

| MAbPac Protein A, 12 μm , 35 x 4.0 mm | |
|--|--|
| Flow rate | 2 mL/min |
| Mobile phase A | 50 mM sodium phosphate, 150 mM NaCl, 5% acetonitrile, pH 7.5 |
| Mobile phase B | 50 mM sodium phosphate, 150 mM NaCl, 5% acetonitrile, pH 2.5 |
| Gradient | 0% B for 0.2 mins, 100% B for 0.60 mins, 0% B for 1.20 mins |
| Temperature | 30 °C |
| Injection volume | 10 μL |
| Detection | UV at 280 nm |
| Sample | mAb B, 5 mg/mL harvest cell culture |



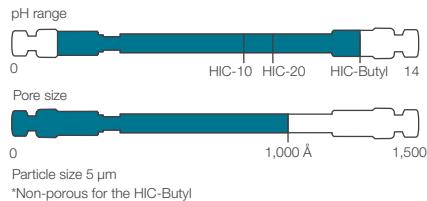
MAbPac Protein A column

| Particle size (μm) | Format | Length (mm) | 4.0 mm ID |
|---------------------------------|-------------|-------------|------------------------|
| 12 | HPLC column | 35 | 082539 |

Intact analysis by HIC

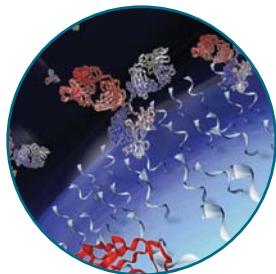
Orthogonal to IEX and SEC, Hydrophobic Interaction Chromatography (HIC) offers selectivity to resolve charge neutral protein oxidations and protein misfolds. Our proprietary 1000 Å silica Thermo Scientific™ MAbPac™ HIC-10 and Thermo Scientific™ MAbPac™ HIC-20 provide unique separation profiles offering high resolution for protein samples. For more hydrophobic samples, select the Thermo Scientific™ MAbPac™ HIC-Butyl column.

MAbPac HIC-10, HIC-20, HIC-Butyl columns

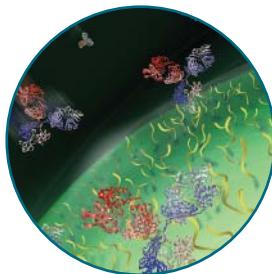


Additional reading

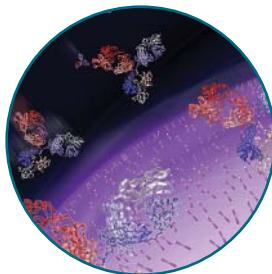
| Links | Type | Description |
|-------|------------------|--|
| | Application note | MAbPac HIC-10 High resolution separation of a fusion protein on MAbPac HIC-10 column |
| | Application note | HIC as a complementary, confirmatory tool to SEC for the analysis of mAb aggregates |
| | Application note | MAbPac HIC-20 High resolution separation of mAb fragments on MAbPac HIC-20 column |
| | Application note | High resolution separation of monoclonal antibody (mAb) oxidation variants |
| | Application note | High resolution separation of cysteine-conjugated antibody drug mimics |
| | Learn more | thermofisher.com/biolc |



MAbPac HIC-10



MAbPac HIC-20



MAbPac HIC-Butyl

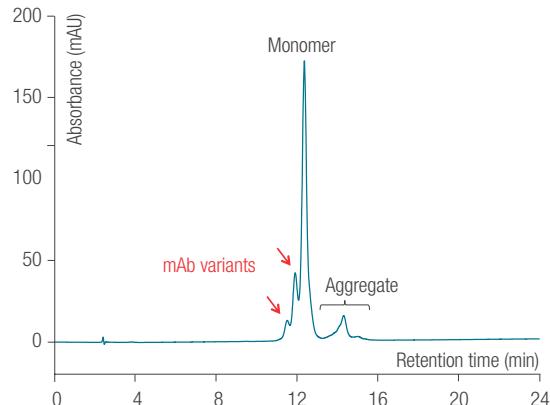
Intact analysis by HIC

continued

Separation of mAb aggregates

MAbPac HIC-10, 5 µm, 100 x 4.6 mm

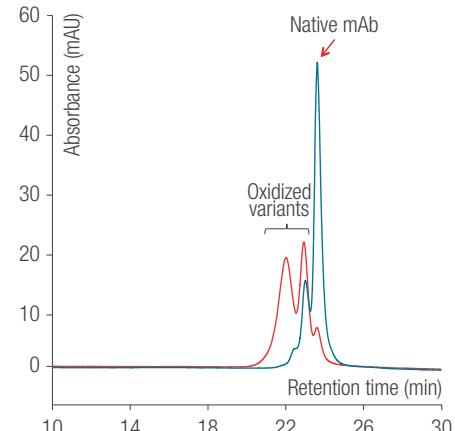
| | | |
|------------------|---|-------|
| Flow rate | 0.5 mL/min | |
| Mobile phase A | 2 mM ammonium sulfate, 100 mM sodium phosphate, pH 7.0 | |
| Mobile phase B | 100 mM sodium phosphate, pH 7.0 | |
| Temperature | 20 °C | |
| Injection volume | 15 µL | |
| Detection | UV at 280 nm | |
| Sample | Monoclonal antibody (4 mg/mL) | |
| | Time (min) | %A %B |
| Gradient | -5.0 | 60 40 |
| | 0.0 | 60 40 |
| | 1.0 | 60 40 |
| | 29.0 | 0 0 |
| | 34.0 | 0 0 |



Separation of mAb fragments

MAbPac HIC-20, 5 µm, 250 x 4.6 mm

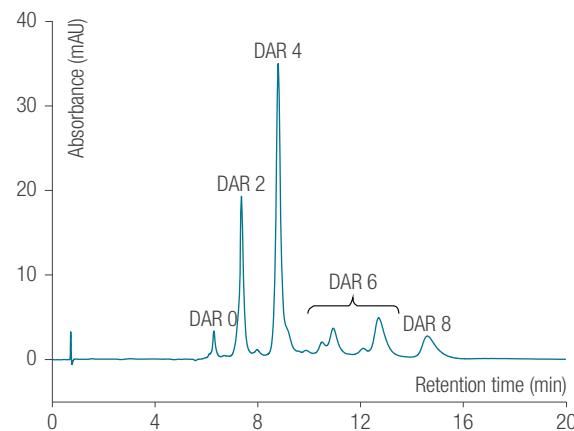
| | | |
|------------------|---|-------|
| Flow rate | 0.5 mL/min | |
| Mobile phase A | 2 mM ammonium sulfate, 100 mM sodium phosphate, pH 7.0 | |
| Mobile phase B | 100 mM sodium phosphate, pH 7.0 | |
| Temperature | 30 °C | |
| Injection volume | Untreated mAb: 20 µL (1.25 mg/mL) Oxidized mAb: 20 µL (1.25 mg/mL) | |
| Detection | UV at 280 nm | |
| Sample | Untreated mAb H ₂ O ₂ oxidized mAb | |
| | Time (min) | A% %B |
| Gradient | -6.0 | 50 50 |
| | 0.0 | 50 50 |
| | 2.0 | 50 50 |
| | 30.0 | 0 100 |
| | 35.0 | 0 100 |



Separation of Antibody Drug Conjugates (ADCs)

MAbPac HIC-Butyl, 5 µm, 100 x 4.6 mm

| | | |
|------------------|---|-------|
| Flow rate | 1.0 mL/min | |
| Mobile phase A | 1.5 mM ammonium sulfate, 50 mM sodium phosphate, pH 7.0/ isopropanol (95:5 v/v) | |
| Mobile phase B | 50 mM sodium phosphate, pH 7.0/isopropanol (80:20 v/v) | |
| Temperature | 25 °C | |
| Injection volume | 5 µL | |
| Detection | UV at 280 nm | |
| Sample | Cys-conjugated ADC mimic (5 mg/mL) | |
| | Time (min) | %A %B |
| Gradient | -5.0 | 100 0 |
| | 0.0 | 100 0 |
| | 1.0 | 100 0 |
| | 15.0 | 0 100 |
| | 20.0 | 0 100 |



Intact analysis by HIC continued

MAbPac HIC selection guide

| Column | MAbPac HIC-10 | MAbPac HIC-20 | MAbPac HIC-Butyl |
|--------------------------------------|---------------|---------------|------------------|
| Intact mAbs/proteins | ++++ | +++ | ++ |
| mAb aggregates | ++++ | +++ | ++ |
| mAb fragments (F_{ab} and F_c) | +++ | ++++ | +++ |
| Oxidized mAbs | +++ | ++++ | +++ |
| Antibody Drug Conjugates (ADCs) | +++ | +++ | ++++ |
| Bispecific mAbs | +++ | ++++ | ++ |

Greater number of ++++ denotes greater suitability



MAbPac HIC family columns

| Description | Particle size (μm) | Format | Length (mm) | 4.6 mm ID |
|------------------------|---------------------------------|-------------------------|-------------|------------------------|
| MAbPac HIC-10 | 5 | Guard cartridges (2/pk) | 10 | 088482 |
| | | HPLC column | 100 | 088480 |
| | | | 250 | 088481 |
| MAbPac HIC-20 | 5 | Guard cartridges (2/pk) | 10 | 088555 |
| | | HPLC column | 100 | 088553 |
| | | | 250 | 088554 |
| MAbPac HIC-Butyl | 5 | Guard cartridges (2/pk) | 10 | 088559 |
| | | HPLC column | 100 | 088558 |
| Guard cartridge holder | — | — | — | 069580 |



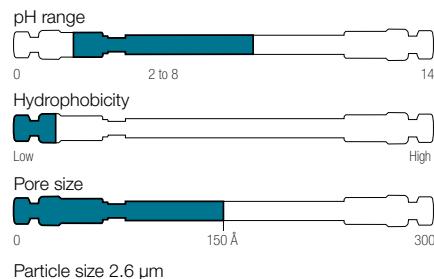
Video:

Introduction to hydrophobic interaction chromatography

Released glycan analysis

For monoclonal antibodies, or protein samples with a lot of neutral glycans, the Thermo Scientific™ Accucore™ 150-Amide HILIC offers outstanding separation on a solid core particle. The low backpressure of this particle allows users to experiment with optimum temperature of their separation, to maximize the elucidation of their released glycan profile. For proteins with charged glycans, we offer two mixed mode column chemistries combining anion exchange with HILIC or RP separations. Thermo Scientific™ GlycanPac™ AXH-1 separates the glycan profile by charge, size, and hydrophilicity. Thermo Scientific™ GlycanPac™ AXR-1 separates the profile by charge, size, and branch isomers.

Accucore 150-Amide-HILIC column

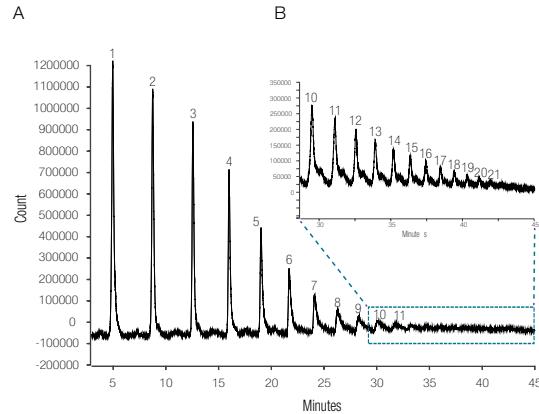


Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | Analysis of human IgG glycans on a solid core amide HILIC stationary phase |
| | Learn more | thermofisher.com/biolc |

2-AB labeled dextran ladder

| Accucore 150-Amide-HILIC, 2.6 μm, 100 x 2.1 mm | |
|--|---|
| Flow rate | 500 μL/min |
| Mobile phase A | Acetonitrile |
| Mobile phase B | 50 mM ammonium formate, pH 4.5 |
| Temperature | 60 °C |
| Injection volume | 2 μL to 5 μL |
| Backpressure at starting conditions | 110 bar |
| Injection wash solvent | 80:20 (v/v) acetonitrile:water |
| Detector | Fluorescence, 330 nm excitation wavelength; 420 nm emission wavelength; acquisition start after 3 min from gradient start |
| Run time | 50 min |
| Gradient | 20–50% B in 40.0 minutes; 50% B for 5.0 minutes 50–20% B in 0.5 minutes; 50% B for 4.5 minutes |



(A) 2 μL injection of sample, where 11 glycans were separated

(B) 5 μL injection of sample, zoomed-in to the later part of the gradient rise.
A further 10 glycans were detected

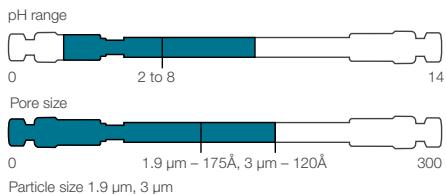


Accucore 150-Amide-HILIC columns

| Particle size (μm) | Format | Length (mm) | 2.1 mm ID | 3.0 mm ID | 4.6 mm ID |
|--------------------|------------------------|-------------|------------------------------|------------------------------|------------------------------|
| 2.6 | HPLC column | 10 | 16726-012105 | — | — |
| | | 50 | 16726-052130 | 16726-053030 | — |
| | | 100 | 16726-102130 | 16726-103030 | 16726-104630 |
| | | 150 | 16726-152130 | 16726-153030 | 16726-154630 |
| | | 250 | 16726-252130 | — | — |
| — | Guard cartridge holder | — | 852-00 | 852-00 | 850-00 |

Released glycan analysis continued

GlycanPac AXH-1 column

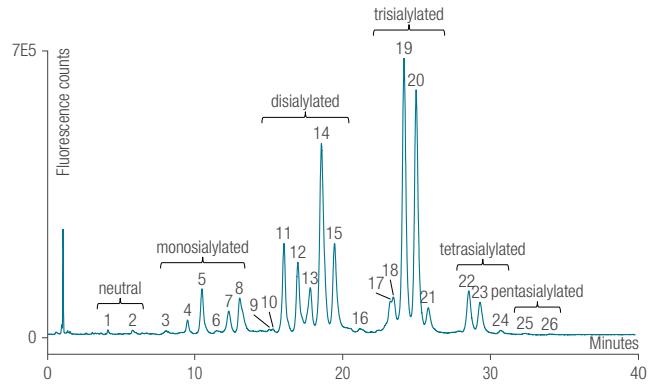


Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | Separation of 2AB-labeled N-linked glycans from bovine fetuin |
| | Application note | Separation of 2AA-labeled N-linked glycans from human IgG |
| | Application note | Separation of 2AA-labeled N-linked glycans from glycoproteins |
| | Learn more | thermofisher.com/biolc |

Separation of 2AB labeled N-glycans from bovine fetuin by charge, size and polarity

| GlycanPac AXH-1, 1.9 μm, 150 x 2.1 mm | | | | |
|---------------------------------------|---|----|----|----|
| Flow rate | 0.4 mL/min | | | |
| Mobile phase A | Acetonitrile (100%) | | | |
| Mobile phase B | Water | | | |
| Mobile phase C | Ammonium formate (100 mM, pH = 4.4) | | | |
| Temperature | 30 °C | | | |
| Injection volume | 5 μL | | | |
| Detection | Fluorescence, 320/420 nm | | | |
| Sample | 2AB labeled N-glycan from bovine fetuin | | | |
| Curve | 5 | | | |
| | Time (min) | %A | %B | %C |
| Gradient | -10.0 | 78 | 20 | 2 |
| | 0.0 | 78 | 20 | 2 |
| | 30.0 | 70 | 20 | 10 |
| | 35.0 | 60 | 20 | 20 |
| | 40.0 | 50 | 20 | 30 |

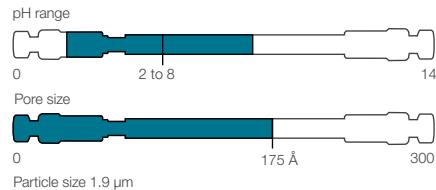


GlycanPac AXH-1 columns

| Particle size (μm) | Format | Length (mm) | 2.1 mm ID | 3.0 mm ID | 4.6 mm ID |
|--------------------|-------------------------|-------------|------------------------|------------------------|------------------------|
| 1.9 | UHPLC column | 100 | 082473 | — | — |
| | | 150 | 082472 | — | — |
| | | 250 | 082521 | — | — |
| 3 | Guard cartridges (2/pk) | 10 | 082476 | 082475 | 082474 |
| | HPLC column | 150 | 082470 | 082469 | 082468 |
| — | Guard cartridge holder | — | 069580 | 069580 | 069580 |

Released glycan analysis continued

GlycanPac AXR-1 column



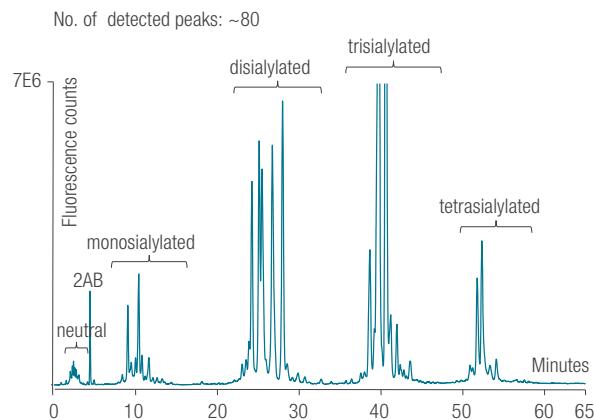
Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | Separation of 2AB labeled N-glycans from bovine fetuin |
| | Application note | Structural analysis of native N-glycans released from proteins |
| | Learn more | thermofisher.com/biolc |

Separation of 2AB labeled N-glycans from bovine fetuin

GlycanPac AXR-1, 1.9 µm, 150 x 2.1 mm

| Flow rate | 0.4 mL/min | | | | | | | | | | | | | | | | | | | | | | | | |
|----------------|--|------------|----|----|----|-------|---|----|---|-----|---|----|---|-----|---|----|----|------|---|----|----|------|----|----|----|
| Mobile phase A | Acetonitrile | | | | | | | | | | | | | | | | | | | | | | | | |
| Mobile phase B | Water | | | | | | | | | | | | | | | | | | | | | | | | |
| Mobile phase C | Ammonium formate (100 mM, pH = 4.4) | | | | | | | | | | | | | | | | | | | | | | | | |
| Temperature | 40 °C | | | | | | | | | | | | | | | | | | | | | | | | |
| Sample load | 100 pmoles | | | | | | | | | | | | | | | | | | | | | | | | |
| Detection | Fluorescence, 320/420 nm | | | | | | | | | | | | | | | | | | | | | | | | |
| Sample | 2AB labeled N-glycan from bovine fetuin | | | | | | | | | | | | | | | | | | | | | | | | |
| Curve | 5 | | | | | | | | | | | | | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time (min)</th> <th>%A</th> <th>%B</th> <th>%C</th> </tr> </thead> <tbody> <tr> <td>-10.0</td> <td>0</td> <td>95</td> <td>5</td> </tr> <tr> <td>0.0</td> <td>0</td> <td>95</td> <td>5</td> </tr> <tr> <td>1.0</td> <td>0</td> <td>95</td> <td>15</td> </tr> <tr> <td>30.0</td> <td>1</td> <td>74</td> <td>25</td> </tr> <tr> <td>65.0</td> <td>20</td> <td>50</td> <td>30</td> </tr> </tbody> </table> | Time (min) | %A | %B | %C | -10.0 | 0 | 95 | 5 | 0.0 | 0 | 95 | 5 | 1.0 | 0 | 95 | 15 | 30.0 | 1 | 74 | 25 | 65.0 | 20 | 50 | 30 |
| Time (min) | %A | %B | %C | | | | | | | | | | | | | | | | | | | | | | |
| -10.0 | 0 | 95 | 5 | | | | | | | | | | | | | | | | | | | | | | |
| 0.0 | 0 | 95 | 5 | | | | | | | | | | | | | | | | | | | | | | |
| 1.0 | 0 | 95 | 15 | | | | | | | | | | | | | | | | | | | | | | |
| 30.0 | 1 | 74 | 25 | | | | | | | | | | | | | | | | | | | | | | |
| 65.0 | 20 | 50 | 30 | | | | | | | | | | | | | | | | | | | | | | |



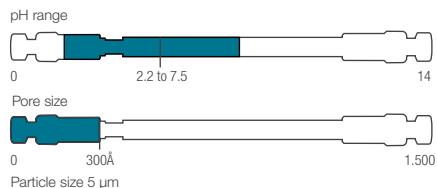
GlycanPac AXR-1 columns

| Particle size (µm) | Format | Length (mm) | 2.1 mm ID | 3.0 mm ID | 4.6 mm ID |
|--------------------|------------------------|-------------|------------------------|------------------------|------------------------|
| 1.9 | UHPLC column | 150 | 088136 | — | — |
| | | 250 | 088135 | — | — |
| — | Guard cartridge holder | — | 069580 | 069580 | 069580 |

Aggregate fragment analysis

For mAb samples, our 300 Å silica Thermo Scientific™ MAbPac™ SEC-1 provides separation of aggregates and fragments samples to characterize your analyte by LC-UV or LC-MS.

MAbPac SEC-1 column

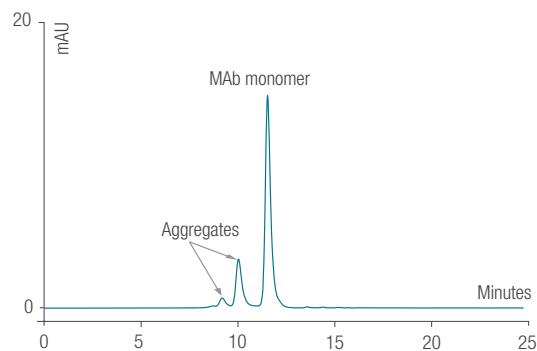


Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | Lifetime stability of size-exclusion chromatography columns for protein aggregate analysis |
| | Application note | Analysis of monoclonal antibodies and their fragments |
| | Learn more | thermofisher.com/biolc |

Monoclonal antibody aggregate separation

| MAbPac SEC-1, 5 μm, 300 x 4.0 mm (PEEK) | |
|---|---|
| Flow rate | 0.20 mL/min |
| Mobile phase | 0.3 mM NaCl in 50 mM phosphate buffer pH 6.8 |
| Gradient | 0% B for 0.2 mins, 100% B for 0.60 mins, 0% B for 1.20 mins |
| Temperature | 30 °C |
| Injection volume | 2 μL |
| Detection | 280 nM |
| Sample | mAb (10 mg/mL) |



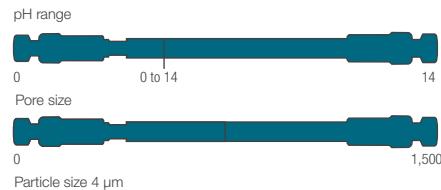
MAbPac SEC-1 columns

| Particle size (μm) | Format | Length (mm) | 2.1 mm ID | 4.0 mm ID | 7.8 mm ID |
|--------------------|--------------|-------------|------------------------|------------------------|------------------------|
| 5 | Guard column | 50 | — | 074697 | — |
| — | HPLC column | 150 | 088790 | 075592 | — |
| — | | 300 | 088789 | 074696 | 088460 |

Intact and subunit analysis (RP)

The wide pore (1500 Å) polymeric Thermo Scientific™ MAbPac™ RP columns offers high resolution separation and minimal carryover for monoclonal antibody samples. Excellent lifetime and ability to separate intact and protein subunits, compatible with LC-UV and LC-MS/MS applications. The monolithic Thermo Scientific™ ProSwift™ RP columns offer unique selectivity, high throughput separations for a wide range of protein sizes. These columns provide high loadability and operate under very low backpressure.

MAbPac RP column



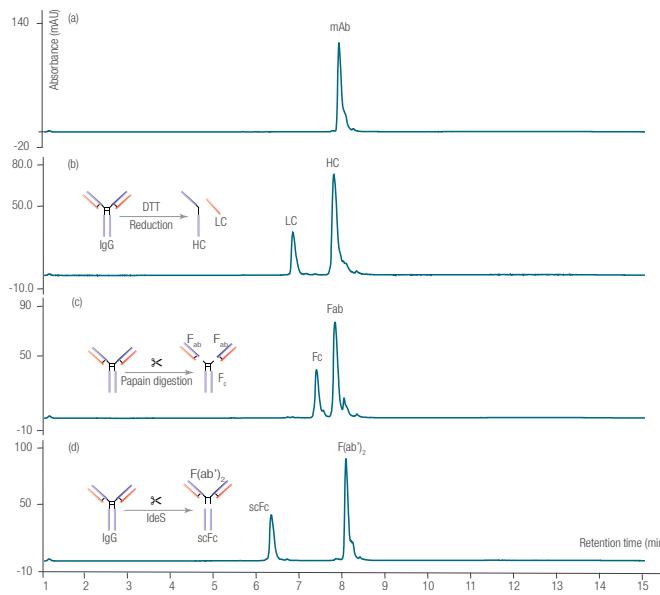
Additional reading

| Links | Type | Description |
|-------|------------------|---|
| | Application note | Confident monoclonal antibody sequence verification by complementary LC-MS techniques |
| | Application note | Fast analysis of therapeutic monoclonal antibody fragments |
| | Learn more | thermofisher.com/biolc |

mAb and mAb fragments analysis

MAbPac RP, 4 µm, 50 x 3.0 mm

| Flow rate | 0.5 mL/min | | | | | | | | | | | | | | | | | | | | | |
|------------------|--|------------|----|----|-----|----|----|-----|----|----|------|----|----|------|----|----|------|----|----|------|----|----|
| Mobile phase A | H ₂ O/FA/TFA (99.88 : 0.1 : 0.02 v/v/v) | | | | | | | | | | | | | | | | | | | | | |
| Mobile phase B | ACN/H ₂ O/FA/TFA 90 : 9.88 : 0.1 : 0.02 v/v/v/v | | | | | | | | | | | | | | | | | | | | | |
| Temperature | 80 °C | | | | | | | | | | | | | | | | | | | | | |
| Injection volume | 5 µL | | | | | | | | | | | | | | | | | | | | | |
| Detection | UV at 280 nm | | | | | | | | | | | | | | | | | | | | | |
| Sample | (a) trastuzumab (5 mg/mL) (b) trastuzumab + DTT (4 mg/mL) (c) trastuzumab + Papain (2 mg/mL) (d) trastuzumab + IdeS (2 mg/mL) | | | | | | | | | | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time (min)</th> <th>%A</th> <th>%B</th> </tr> </thead> <tbody> <tr> <td>0.0</td> <td>80</td> <td>20</td> </tr> <tr> <td>1.0</td> <td>80</td> <td>20</td> </tr> <tr> <td>11.0</td> <td>55</td> <td>45</td> </tr> <tr> <td>12.0</td> <td>55</td> <td>45</td> </tr> <tr> <td>14.0</td> <td>80</td> <td>20</td> </tr> <tr> <td>16.0</td> <td>80</td> <td>20</td> </tr> </tbody> </table> | Time (min) | %A | %B | 0.0 | 80 | 20 | 1.0 | 80 | 20 | 11.0 | 55 | 45 | 12.0 | 55 | 45 | 14.0 | 80 | 20 | 16.0 | 80 | 20 |
| Time (min) | %A | %B | | | | | | | | | | | | | | | | | | | | |
| 0.0 | 80 | 20 | | | | | | | | | | | | | | | | | | | | |
| 1.0 | 80 | 20 | | | | | | | | | | | | | | | | | | | | |
| 11.0 | 55 | 45 | | | | | | | | | | | | | | | | | | | | |
| 12.0 | 55 | 45 | | | | | | | | | | | | | | | | | | | | |
| 14.0 | 80 | 20 | | | | | | | | | | | | | | | | | | | | |
| 16.0 | 80 | 20 | | | | | | | | | | | | | | | | | | | | |



Intact and subunit analysis (RP) continued



MAbPac RP columns

| Particle size (μm) | Format | Length (mm) | 2.1 mm ID | 3.0 mm ID |
|---------------------------------|-------------------------|-------------|------------------------|------------------------|
| 4 | Guard cartridges (2/pk) | 10 | 088649 | 088646 |
| | | 50 | 088648 | 088645 |
| | HPLC column | 100 | 088647 | 088644 |
| | | 150 | 303270 | 303269 |
| — | Guard cartridge holder | — | 069580 | 069580 |

MAbPac RP 1 mm columns

| Particle size (μm) | Length (mm) | 1 mm ID |
|---------------------------------|-------------|------------------------|
| 4 | 50 | 303182 |
| | 100 | 303183 |
| | 150 | 303184 |



Webinars

Analytical and life science webinars live and on-demand



NIBRT collaboration information

A collaboration built for Biopharma between the National Institute for Bioprocessing Research and Training (NIBRT) and Thermo Fisher Scientific
thermofisher.com/nibrt

Intact and subunit analysis (RP) continued

ProSwift RP column

Additional reading

| Links | Description |
|--|-------------|
|  Learn more thermofisher.com/biolc | |

ProSwift column

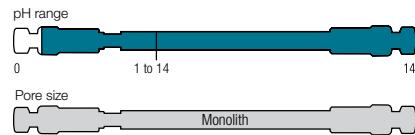
ProSwift RP-2H, 50 x 4.6 mm

| | |
|------------------|---|
| Flow rate | 1, 2, 4, or 8 mL/min |
| Mobile phase A | H ₂ O/ACN (95:5; V/V) + 0.1% TFA |
| Mobile phase B | H ₂ O/ACN (5:95; V/V) + 0.1% TFA |
| Injection volume | 2 µL |
| Detection | UV at 214 nm |
| Sample | Mixture of five proteins |
| Gradient | 1 mL/min: 1-75% B in 12 min 2 mL/min: 1-75% B in 6 min 4 mL/min: 1-75% B in 3 min 8 mL/min: 1-75% B in 1.5 min |
| Analytes | 1. Ribonuclease A 1.5 mg/mL 2. Cytochrome C 0.5 mg/mL 3. BSA 1.5 mg/mL 4. Carbonic anhydrase 0.9 mg/mL 5. Ovalbumin 1.5 mg/mL |

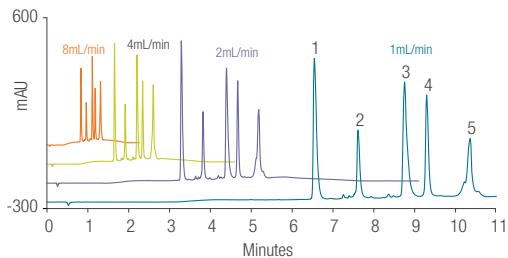


ProSwift RP columns

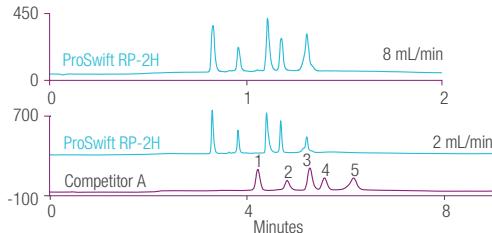
| Functional group | Length (mm) | 1.0 mm ID | 4.6 mm ID |
|------------------|-------------|------------------------|------------------------|
| RP-1S | 50 | — | 064297 |
| RP-2H | 50 | — | 064296 |
| RP-3U | 50 | — | 064298 |
| RP-4H | 50 | 069477 | — |
| | 250 | 066640 | — |



Proteins



Competitive comparison



Charge variant analysis

For charge variant analysis by LC-UV or LC-MS/MS Thermo Scientific™ ProPac™ 3R SAX and Thermo Scientific™ ProPac™ 3R SCX columns deliver outstanding resolution on a highly robust, reproducible and high-resolution platform. Combine ProPac 3R SCX columns with our proprietary CX-1 buffers formulations to enable fast, robust and reproducible pH gradients that are simple to optimize and easily automated - without the need for time-consuming mobile phase adjustments.

Protein isoelectric point (pI)

<7

>7

ProPac 3R SAX column

ProPac 3R SCX column

MAbPac SCX-10 column

ProPac WCX-10 column

ProPac Elite WCX column

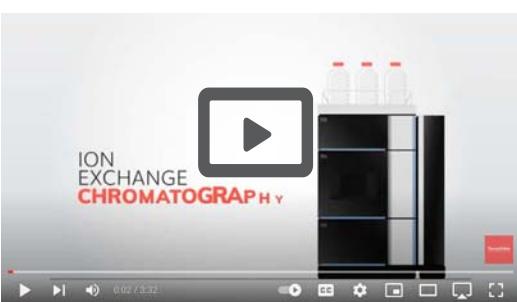
- Works well with salt and pH gradient buffers
- Best choice for proteins with acidic pI
- Analyze full/empty AAV capsid ratios

- Highest resolution with excellent reproducibility
- Works well with CX-1 buffers

- Alternative selectivity to WCX, scalable from short methods over comprehensive analysis to semi-prep formats
- Works well with CX-1 buffers

- Industry GOLD standard – widely used and published

- Improved resolution, speed and reproducibility over ProPac WCX-10 column
- Works well with CX-1 buffers



Video:

Tips to improve your charge variant analysis by ion exchange



Charge variant analysis continued

ProPac 3R SCX column

Additional reading

| Links | Type | Description |
|-------|------------------|---|
| | Application note | Salt gradient analysis of monoclonal antibodies using a 3 µm monodisperse SCX column |
| | Application note | Method development for pH gradient analysis of monoclonal antibodies using SCX column |
| | Learn more | thermofisher.com/propac |

Salt gradient analysis of infliximab

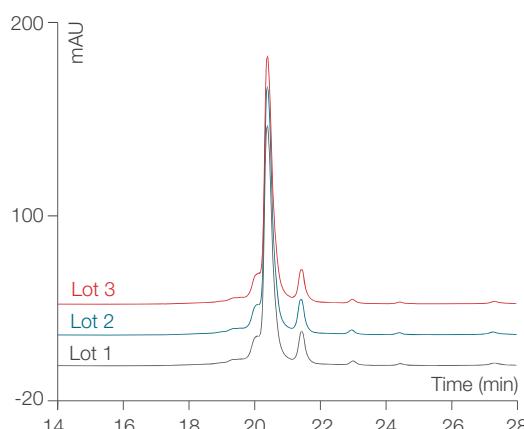
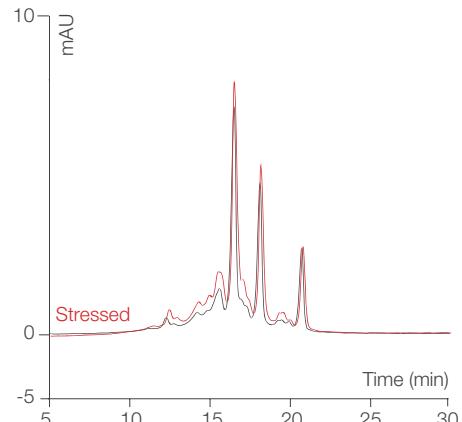
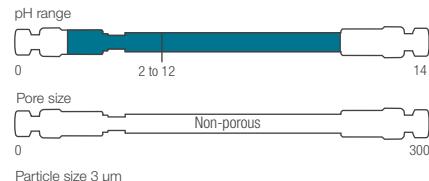
ProPac 3R SCX column, 3 µm

| | | |
|--------------|---|----|
| Format | 4 x 100 mm | |
| Mobile phase | A: 20 mM MES, pH 6.5 B: 20 mM MES, pH 6.5 + 0.5 M NaCl | |
| Flow rate | 0.3 mL/min | |
| Injection | 2 µL | |
| Temp | 30 °C | |
| Detection | UV, 280 nm | |
| Sample | Infliximab – 5 mg/mL | |
| Gradient | %A | %B |
| Time (min) | 0.0 | 93 |
| | 30.0 | 78 |
| | 30.1 | 20 |
| | 33.0 | 20 |
| | 33.1 | 93 |
| | 40.0 | 93 |
| | | 7 |

Lot-to-lot reproducibility of NISTmAb salt gradient separation

ProPac 3R SCX column, 3 µm

| | | |
|--------------|---|----|
| Format | 4 x 100 mm | |
| Mobile phase | A: 20 mM MES, pH 6.5 B: 20 mM MES, pH 6.5 + 0.5 M NaCl | |
| Flow rate | 0.3 mL/min | |
| Injection | 2 µL | |
| Temp | 30 °C | |
| Detection | UV, 280 nm | |
| Sample | NISTmAb – 10 mg/mL | |
| Gradient | %A | %B |
| Time (min) | 0.0 | 95 |
| | 30.0 | 75 |
| | 30.1 | 20 |
| | 33.0 | 20 |
| | 33.1 | 95 |
| | 40.0 | 95 |
| | | 10 |



ProPac 3R SCX 3 µm columns

| Particle size (µm) | Length (mm) | 2.0 mm ID | 4.0 mm ID |
|--------------------|-------------|------------------------------|------------------------------|
| 3 | 50 | 43103-052068 | 43103-054068 |
| | 100 | 43103-102068 | 43103-104068 |

Charge variant analysis

continued

ProPac 3R SAX column

Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | Salt gradient analysis of Protein G using a 3 µm monodisperse SAX column |
| | Application note | Salt gradient separation and analysis of adeno-associated virus samples using SAX column |
| | Learn more | thermofisher.com/propac |

Salt gradient analysis of protein G

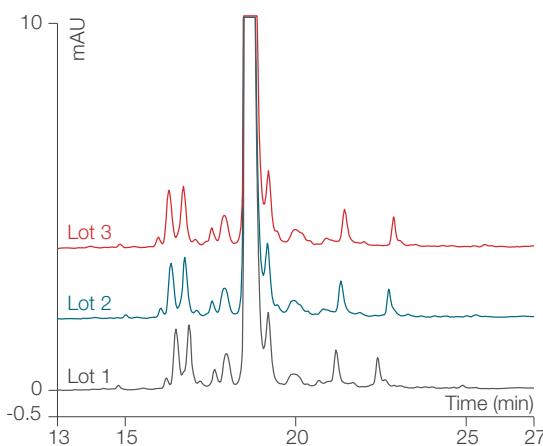
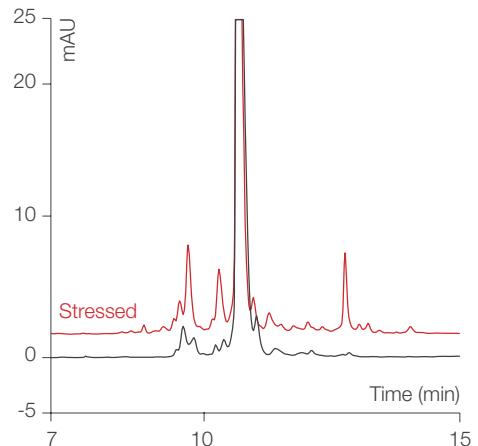
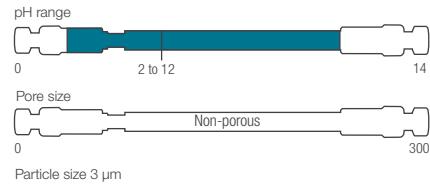
ProPac 3R SAX column, 3 µm

| Format | 4 x 100 mm | |
|--------------|--|-----|
| Mobile phase | A: 20 mM Tris, pH 8.0 B: 20 mM Tris + 500 mM NaCl, pH 8.0 | |
| Flow rate | 0.5 mL/min | |
| Injection | 1 µL | |
| Temp | 30 °C | |
| Detection | UV, 280 nm | |
| Sample | Protein G – 5 mg/mL | |
| Gradient | %A | %B |
| 0.0 | 88 | 12 |
| 1.0 | 88 | 12 |
| 16.0 | 58 | 42 |
| 16.1 | 0 | 100 |
| 18.0 | 0 | 100 |
| 18.1 | 88 | 12 |
| 30.0 | 88 | 12 |

Lot-to-lot reproducibility of protein G salt gradient separation

ProPac 3R SAX column, 3 µm

| Format | 4 x 100 mm | |
|--------------|--|-----|
| Mobile phase | A: 20 mM Tris, pH 8.0 B: 20 mM Tris + 500 mM NaCl, pH 8.0 | |
| Flow rate | 0.5 mL/min | |
| Injection | 1 µL | |
| Temp | 30 °C | |
| Detection | UV, 280 nm | |
| Sample | Protein G – 5 mg/mL | |
| Gradient | %A | %B |
| 0.0 | 88 | 12 |
| 1.0 | 88 | 12 |
| 31.0 | 58 | 42 |
| 31.1 | 0 | 100 |
| 33.0 | 0 | 100 |
| 33.1 | 88 | 12 |
| 45.0 | 88 | 12 |

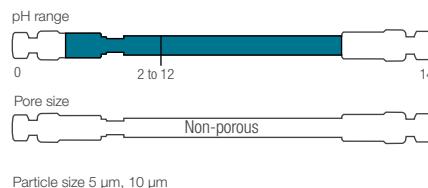


ProPac 3R SAX 3 µm columns

| Particle size (µm) | Length (mm) | 2.0 mm ID | 4.0 mm ID |
|--------------------|-------------|------------------------------|------------------------------|
| 3 | 50 | 43203-052068 | 43203-054068 |
| | 100 | 43203-102068 | 43203-104068 |

Charge variant analysis continued

MAbPac SCX-10 column

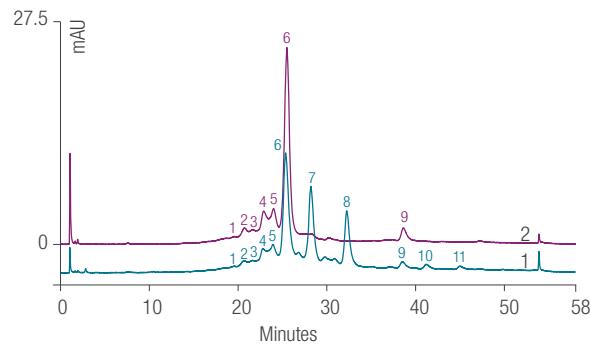


Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Application note | A global pH-gradient based charge variant analysis |
| | Application note | High throughput, high resolution monoclonal antibody analysis |
| | Learn more | thermofisher.com/biolc |

Baseline resolution of C-terminal lysine variants of a monoclonal antibody

| MAbPac SCX-10, 5 μm, 250 x 4.0 mm | |
|-----------------------------------|--|
| Flow rate | 1 mL/min |
| Mobile phase A | 20 mM MES (pH 5.6) + 60 mM NaCl |
| Mobile phase B | 20 mM MES (pH 5.6) + 300 mM NaCl |
| Gradient | 15–36% B in 50 min |
| Temperature | 30 °C |
| Injection volume | 5 μL |
| Detection | UV at 280 nm |
| Sample | 1. mAb B, 900 μg in 100 μL (no carboxypeptidase) 2. mAb B, 900 μg in 100 μL + carboxypeptidase, 50 μg, incubation at 37 °C for 3 h |
| Both chromatograms | Peaks 1–5: acidic variants |
| Sample 1 | Peaks 6–8: C-Terminal lysine truncation variants of main peak. Peaks 9–11: C-Terminal lysine truncation variants of minor variant peak |
| Sample 2 | Peak 6 results from peaks 6, 7, and 8 after CBP treatment. Peak 9 results from peaks 9, 10, and 11 after CBP treatment |

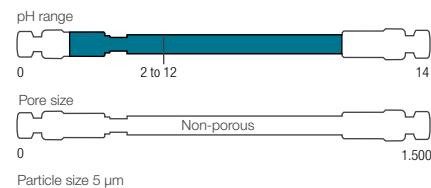


MAbPac SCX-10 columns

| Particle size (μm) | Format | Length (mm) | 2.0 mm ID | 4.0 mm ID | 9.0 mm ID |
|--------------------|--------------|-------------|------------------------|------------------------|------------------------|
| 5 | HPLC column | 50 | — | 078656 | — |
| | | 150 | — | 085198 | — |
| | | 250 | — | 078655 | — |
| 10 | Guard column | 50 | 075749 | 074631 | — |
| | | 50 | — | 075603 | — |
| | HPLC column | 150 | — | 075602 | — |
| | | 250 | 075604 | 074625 | 088784 |

Charge variant analysis continued

MAbPac SCX-10 RS column



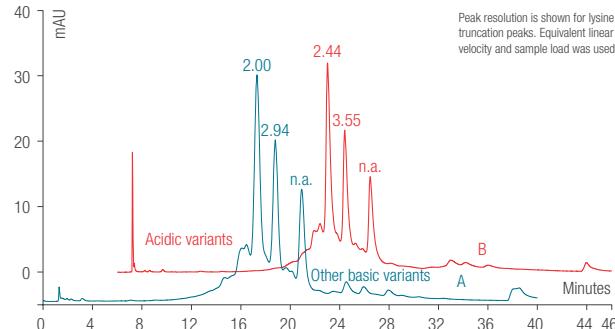
Additional reading

| Links | Description |
|------------------|---|
| | Learn more thermofisher.com/biolc |

Lysine variants

MAbPac SCX, 5 μm, 250 x 4.6 mm

| | |
|--------------------|------------------------------|
| Flow rate | 1.5 mL/min |
| Mobile phase A | 20 mM MES pH 5.6 + 60 mM |
| Mobile phase B | 20 mM MES pH 5.6 + 3 mM NaCl |
| Injection volume | 15 μL |
| Detection | UV at 280 nm |
| Sample | mAb 5 mg/mL |
| Both chromatograms | Peaks 1–5: acidic variants |
| Chromatogram A | Gradient: 33–53% B in 30 min |
| Chromatogram B | Gradient: 33–53% B in 20 min |

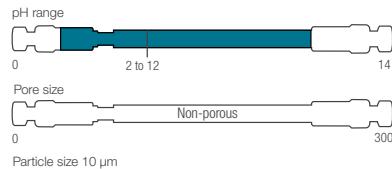


MAbPac SCX-10 RS columns

| Particle size (μm) | Format | Length (mm) | 2.1 mm ID | 4.6 mm ID |
|--------------------|--------------|-------------|------------------------|------------------------|
| 5 | UHPLC column | 50 | 082675 | 082674 |
| | | 150 | 088242 | 085209 |
| | | 250 | 082515 | 082673 |

Charge variant analysis continued

ProPac SAX-10 column



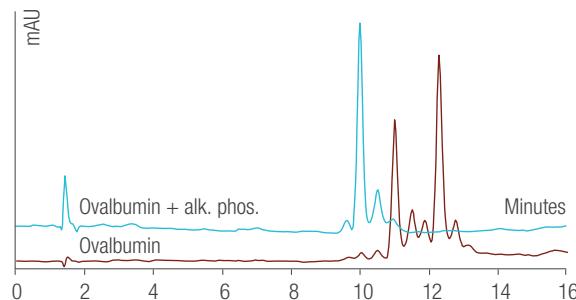
Additional reading

| Links | Description |
|---|-------------|
| Learn more thermofisher.com/biolc | |

Resolution of phosphorylation variants of ovalbumin

ProPac SAX-10, 10 μm, 250 x 4.0 mm

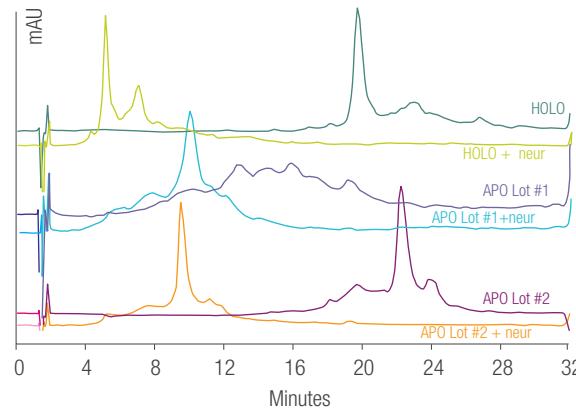
| Flow rate | 1.0 mL/min | | | | | | | | | | | | |
|------------------|--|------------|----|----|----|-----|----|---|----|------|------|------|----|
| Mobile phase A | Water | | | | | | | | | | | | |
| Mobile phase B | 2.0 mM NaCl | | | | | | | | | | | | |
| Mobile phase C | 0.1 mM Tris/HCl (pH 8.5) | | | | | | | | | | | | |
| Injection volume | 1.0 μL | | | | | | | | | | | | |
| Detection | UV at 214 nm | | | | | | | | | | | | |
| Sample | Ovalbumin before and after alkaline phosphatase treatment | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time (min)</th> <th>%A</th> <th>%B</th> <th>%C</th> </tr> </thead> <tbody> <tr> <td>0.0</td> <td>80</td> <td>0</td> <td>20</td> </tr> <tr> <td>15.0</td> <td>67.5</td> <td>12.5</td> <td>20</td> </tr> </tbody> </table> | Time (min) | %A | %B | %C | 0.0 | 80 | 0 | 20 | 15.0 | 67.5 | 12.5 | 20 |
| Time (min) | %A | %B | %C | | | | | | | | | | |
| 0.0 | 80 | 0 | 20 | | | | | | | | | | |
| 15.0 | 67.5 | 12.5 | 20 | | | | | | | | | | |



Effect of sialylation on transferrin chromatography

ProPac SAX-10, 10 μm, 250 x 4.0 mm

| Flow rate | 1.0 mL/min | | | | | | | | | | | | |
|------------------|---|------------|----|----|----|-----|----|---|----|------|----|---|----|
| Mobile phase A | Water | | | | | | | | | | | | |
| Mobile phase B | 2.0 mM NaCl | | | | | | | | | | | | |
| Mobile phase C | 0.2 mM Tris/HCl (pH 9) | | | | | | | | | | | | |
| Injection volume | 50.0 μL | | | | | | | | | | | | |
| Detection | UV at 214 nm | | | | | | | | | | | | |
| Sample | HOLO (iron rich) and APO (iron poor) human transferrin samples before and after neuraminidase treatment. Digestions were carried out overnight at 37 °C in sodium acetate buffer at pH 5. | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time (min)</th> <th>%A</th> <th>%B</th> <th>%C</th> </tr> </thead> <tbody> <tr> <td>0.0</td> <td>87</td> <td>3</td> <td>10</td> </tr> <tr> <td>30.0</td> <td>83</td> <td>7</td> <td>10</td> </tr> </tbody> </table> | Time (min) | %A | %B | %C | 0.0 | 87 | 3 | 10 | 30.0 | 83 | 7 | 10 |
| Time (min) | %A | %B | %C | | | | | | | | | | |
| 0.0 | 87 | 3 | 10 | | | | | | | | | | |
| 30.0 | 83 | 7 | 10 | | | | | | | | | | |

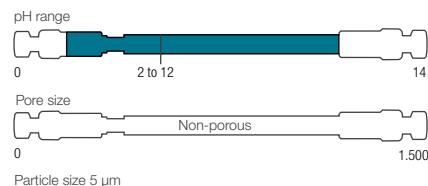


ProPac SAX-10 columns

| Particle size (μm) | Format | Length (mm) | 2.0 mm ID | 4.0 mm ID | 9.0 mm ID | 22.0 mm ID | 4 x 50 mm ID |
|--------------------|--------------|-------------|------------------------|------------------------|------------------------|------------------------|------------------------|
| 10 | Guard column | 50 | 063454 | 054998 | — | — | — |
| | HPLC column | 250 | 063448 | 054997 | 063703 | 088770 | 078990 |

Charge variant analysis continued

ProPac Elite WCX column

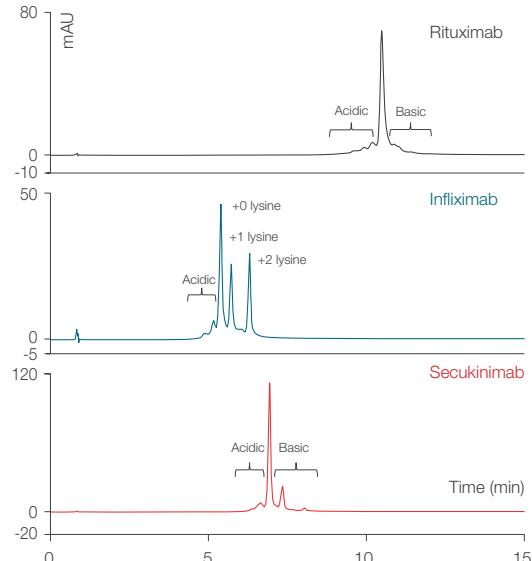


Additional reading

| Links | Type | Description |
|-------|------------------|---|
| | Application note | Confident monoclonal antibody sequence verification by complementary LC-MS techniques |
| | Application note | Fast analysis of therapeutic monoclonal antibody fragments |
| | Learn more | thermofisher.com/biolc |

ProPac Elite WCX, 5 μ m, 150 x 4.0 mm

| | |
|------------------|---|
| Flow rate | 1.0 mL/min |
| Mobile phase A | 1x CX-1 pH Gradient buffer A |
| Mobile phase B | 1x CX-1 pH Gradient buffer B |
| Temperature | 30 °C |
| Injection volume | 2 μ L |
| Detection | UV at 280 nm |
| Sample | Top: rituximab, 5 mg/mL Middle: infliximab, 5 mg/mL Bottom: secukinimab, 5 mg/mL |
| Gradient | Time (min) %A %B 0.0 80 20 15.0 20 80 15.1 0 100 17.0 0 100 17.1 80 20 25.0 80 20 |



ProPac Elite WCX columns

| Particle size (μ m) | Format | Length (mm) | 2.0 mm ID | 4.0 mm ID |
|--------------------------|-------------|-------------|------------------------|------------------------|
| 5 | HPLC column | 50 | 303028 | 302973 |
| | | 100 | 303027 | 302972 |
| | | 250 | 303026 | 303025 |

ProPac Elite WCX kits

| Particle size (μ m) | Set contents | Length (mm) | 4.0 mm ID |
|--------------------------|-----------------------|-------------|------------------------|
| 5 | 3 columns from 1 lot | 150 | 302976 |
| | 3 columns from 3 lots | 150 | 302977 |
| | 3 columns from 1 lot | 250 | 303061 |
| | 3 columns from 3 lots | 250 | 303062 |

Charge variant analysis continued

pH gradient buffers

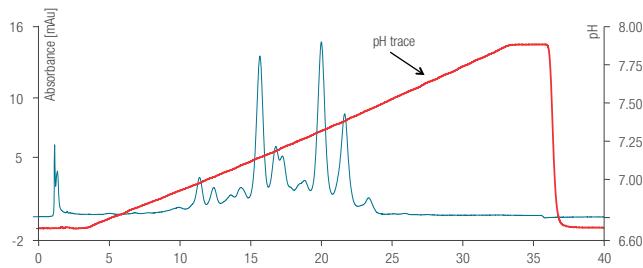
Ready-to-use buffers for simple method development during charge variant characterization

Thermo Scientific™ pH gradient platform accelerates method development and facilitates method transfer to QA/QC for a wide range of protein and mAb charge variants through a generic LC-based approach to charge variant characterization.

- Patented buffer formulations enable fast, robust and reproducible pH gradients that are simple to optimize and easily automated
- Ready to use with existing LC columns and systems, without the need for time consuming mobile phase adjustments
- Applicable to the majority of mAbs



Optimization of mAb charge variant separation using a linear pH gradient: 25% B (pH 6.75) to 50% B (pH 7.9)



pH gradient buffers

| Description | Buffer bottle size | | | |
|-------------------------------------|------------------------|------------------------|------------------------|------------------------|
| Buffer | 125 mL | 250 mL | 500 mL | 1000 mL |
| CX-1 pH gradient buffer A (pH 5.6) | 083273 | 085346 | 302779 | 303274 |
| CX-1 pH gradient buffer B (pH 10.2) | 083275 | 085348 | 302780 | 303275 |



NIBRT collaboration information

A collaboration built for Biopharma between the National Institute for Bioprocessing Research and Training (NIBRT) and Thermo Fisher Scientific
thermofisher.com/nibrt

Peptide mapping and MAM

Thermo Scientific™ Hypersil GOLD™ VANQUISH™ C18 UHPLC columns are an excellent column choice for a broad range of peptides, offering high resolution for all critical quality attributes, without extremely long retention for more hydrophobic peptides. For faster separation of peptide samples select the Thermo Scientific™ Accucore™ C18 VANQUISH™ column. The column offers sub-2 µm particles providing ultra-short diffusion paths that result in extremely efficient separations.

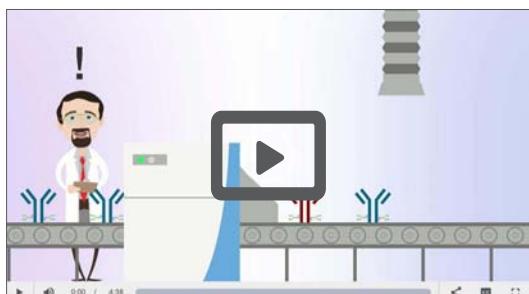
Additional reading

| Links | Type | Description |
|---|---|--|
|  | Landing page | Multi-Attribute Method (MAM): Straight through to breakthrough |
|  | Learning centre | Biopharmaceutical Multi-Attribute Method (MAM) |
|  | Learn more thermofisher.com/biolc | |



Video:

End-to-end MAM solution to move biopharma forward



Video:

Learn how innovation and monitoring strategies can reduce the number of tests and enhance the methodology of validating impurity

Peptide mapping and MAM

continued

Hypersil GOLD VANQUISH column

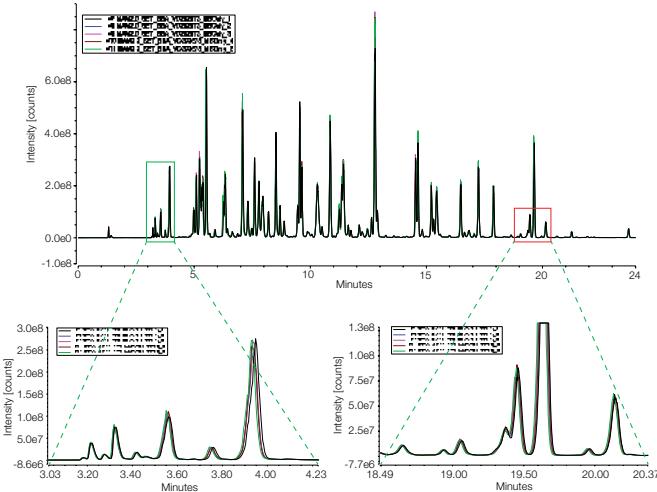
Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Flyer | VANQUISH UHPLC columns. Delivering powerful separations |
| | Application note | An integrated LC-MS system performance evaluation test |
| | Learn more | thermofisher.com/biolc |

Overlay of 5 TIC traces from the SET injection sequence

Hypersil GOLD VANQUISH C18 UHPLC column, 150 × 2.1 mm, 1.9 µm

| | |
|------------------|--|
| Flow rate | 0.25 mL/min |
| Mobile phase A | H ₂ O + 0.1% FA |
| Mobile phase B | ACN + 0.1% FA |
| Injection volume | 5 µL |
| Detection | Mass spectrometer – Full scan |
| Sample | Pierce BSA protein digest standard, MS grade, UD294474 (P/N 88341) |
| Chromatogram B | Gradient: 33-53% B in 20 min |



Hypersil GOLD Vanquish columns

| Particle size (µm) | Length (mm) | 2.1 mm ID |
|--------------------|-------------|--------------------------------|
| 1.9 | 50 | 25002-052130-V |
| | 100 | 25002-102130-V |
| | 150 | 25002-152130-V |

Peptide mapping and MAM

continued

Accucore VANQUISH C18+ column

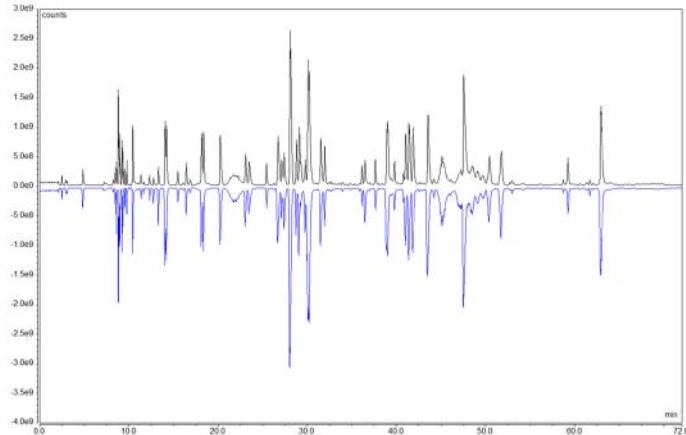
Additional reading

| Links | Type | Description |
|---|------------------|---|
|  | Application note | Comparative analysis of innovator and biosimilar monoclonal antibodies using MAM method |
|  | Technical guide | Powerful separations are our core performance |
|  | Poster | Application of a MS in QC method for characterization and attribute monitoring |
|  | Learn more | thermofisher.com/biolc |

Mirrored base peak chromatograms of rituximab innovator (black) and its biosimilar product (blue)

Accucore Vanquish C18+ UHPLC column, 1.5 µm, 2.1 x 150 mm

| | |
|------------------|----------------------------|
| Flow rate | 0.25 mL/min |
| Mobile phase A | H ₂ O + 0.1% FA |
| Mobile phase B | ACN + 0.1% FA |
| Injection volume | 8 µL |
| Detection | Mass spectrometer |
| Sample | Rituximab innovator |
| Temperature | 50 °C |



Accucore Vanquish C18+ columns

| Particle size (µm) | Length (mm) | 2.1 mm ID |
|--------------------|-------------|------------------------------|
| 1.5 µm | 50 mm | 27101-052130 |
| | 100 mm | 27101-102130 |
| | 150 mm | 27101-152130 |

Nucleic acids/oligonucleotides

Thermo Scientific™ DNAPac™ RP column offers ion-pair reversed phase separations of nucleic acid mixtures. Samples from siRNA to mRNA easily resolve on this polymer chemistry. Compatible with LC-UV and LC-MS/MS methodologies this column delivers outstanding separations.

Thermo Scientific™ DNAPac™ PA200 and Thermo Scientific™ DNAPac™ PA200RS columns are strong anion exchange columns for n-1 separation of oligo samples. Compatible with LC-UV, these columns offer orthogonal separation to reversed phase columns, separating the oligonucleotide sample by size and charge.

Thermo Scientific™ DNASwift™ column is a monolithic column designed for users who would like to do SAX purification of oligonucleotide samples using their analytical HPLC. These monolithic columns offer high loadability, with slightly less resolution than our analytical columns.



Webinars

Analytical and life science webinars live and on-demand



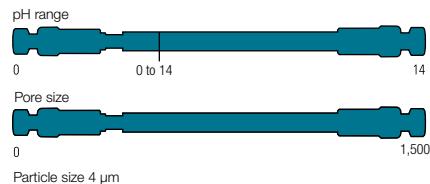
NIBRT collaboration information

A collaboration built for Biopharma between the National Institute for Bioprocessing Research and Training (NIBRT) and Thermo Fisher Scientific

thermofisher.com/nibrt

Nucleic acids/oligonucleotides continued

DNAPac RP column

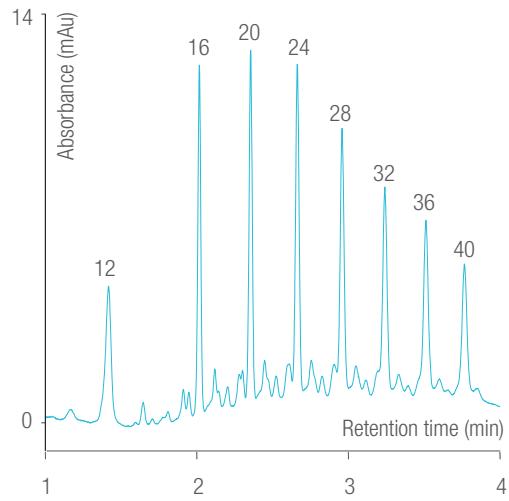


Additional reading

| Links | Description |
|--|-------------|
| Learn more thermofisher.com/biolc | |

Fast analysis of mixed base DNA

| DNAPac RP, 4 µm, 50 x 2.1 mm | | | |
|------------------------------|--|----|----|
| Flow rate | 0.8 mL/min | | |
| Mobile phase A | 25 mM HAA, pH 8.5 | | |
| Mobile phase B | 25 mM HAA, pH 8.5/acetonitrile (50:50 v/v) | | |
| Temperature | 65 °C | | |
| Injection volume | 4 µL | | |
| Detection | UV at 260 nm | | |
| Sample | 8-Combo DNA | | |
| Gradient curve | 3 | | |
| Peak label | Length of DNA | | |
| | Time (min) | %A | %B |
| Gradient | -0.1 | 67 | 33 |
| | 0.0 | 67 | 33 |
| | 3.0 | 41 | 59 |
| | 3.1 | 5 | 95 |
| | 4.9 | 5 | 95 |
| | 5.0 | 67 | 33 |
| | 8.0 | 67 | 33 |

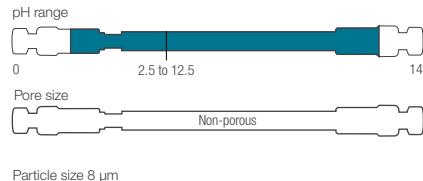


DNAPac RP columns

| Particle size (µm) | Format | Length (mm) | 2.1 mm ID | 3.0 mm ID |
|--------------------|-------------------------|-------------|------------------------|------------------------|
| 4 | Guard cartridges (2/pk) | 10 | 088925 | 088921 |
| | | 50 | 088924 | 088920 |
| | HPLC column | 100 | 088923 | 088919 |
| | | 250 | 303324 | — |
| — | Guard cartridge holder | — | 069580 | 069580 |

Nucleic acids/oligonucleotides continued

DNAPac PA200 column



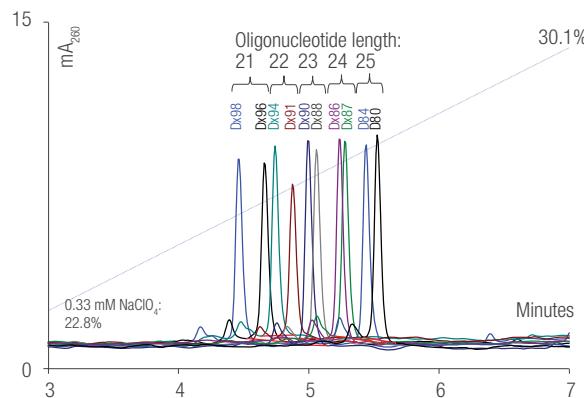
Additional reading

| Links | Description |
|--|-------------|
| Learn more thermofisher.com/biolc | |

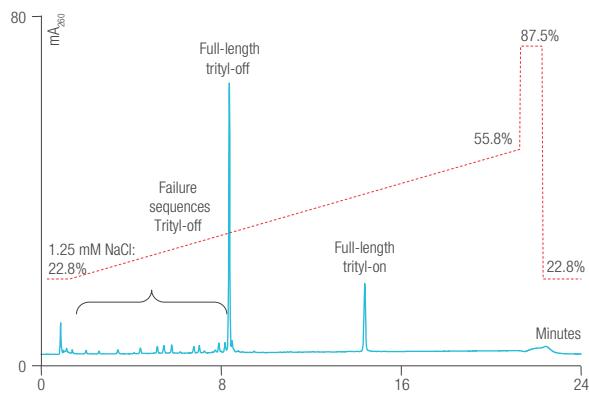
DNAPac PA200, 8 μ m, 250 x 4.0 mm

| | |
|--------------|--|
| Flow rate | 1.2 mL/min |
| Mobile phase | NaClO ₄ , pH 6.5 with 20% ACN |
| Detection | UV at 260 nm |
| Flow rate | 1.2 mL/min |

Separation of oligonucleotides by length



Target, failure and trityl-on oligonucleotides



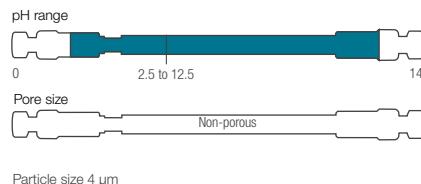
DNAPac PA200 columns

| Particle size (μ m) | Format | Length (mm) | 2.0 mm ID | 4.0 mm ID | 9.0 mm ID | 22.0 mm ID |
|--------------------------|--------------|-------------|------------------------|------------------------|------------------------|------------------------|
| 8 | Guard column | 50 | 063423 | 062998 | 063419 | 088780 |
| | HPLC column | 250 | 063425 | 063000 | 063421 | 088781 |

Nucleic acids/oligonucleotides

continued

DNAPac PA200 RS column



Additional reading

| Links | Type | Description |
|-------|------------------|--|
| | Brochure | Superior oligonucleotide analysis |
| | Application note | High resolution separation of oligonucleotides |
| | Application note | Ultra-high-resolution separation of oligonucleotides by UHPLC |
| | Application note | Separation of mixed-base oligonucleotides |
| | Learn more | thermofisher.com/biolc |

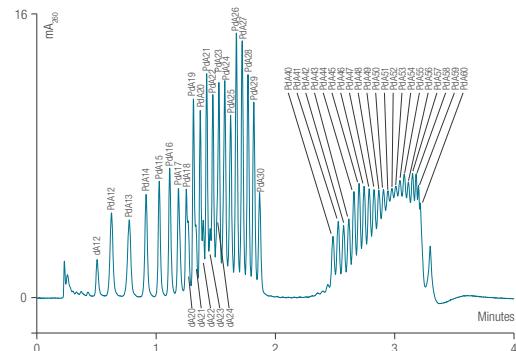
Partial resolution of 46 oligonucleotides

DNAPac PA200 RS, 4 μm, 50 x 4.6 mm

| | |
|------------------|--|
| Flow rate | 1.30 mL/min |
| Mobile phase A | 20 mM Tris pH 8 |
| Mobile phase B | A + 1.25 mM NaCl |
| Temperature | 30 °C |
| Injection volume | 2.5 μL |
| Gradient | 28–43% B in 4 CV* (2.56 min) curve 3** |
| Sample | PdA12–30, 40–60 |

*CV = column volumes

**Curve 3 indicates continuously changing gradient, asymptotically approaching a maximum salt concentration. Programmed in Thermo Scientific™ Chromeleon™ 6.8.

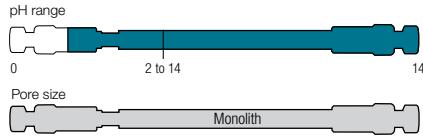


DNAPac PA200 RS columns

| Particle size (μm) | Format | Length (mm) | 4.6 mm ID |
|--------------------|--------------|-------------|------------------------|
| 4 | BioRS column | 50 | 082508 |
| | | 150 | 082509 |
| | | 250 | 082510 |

Nucleic acids/oligonucleotides continued

DNA Swift SAX-1S column



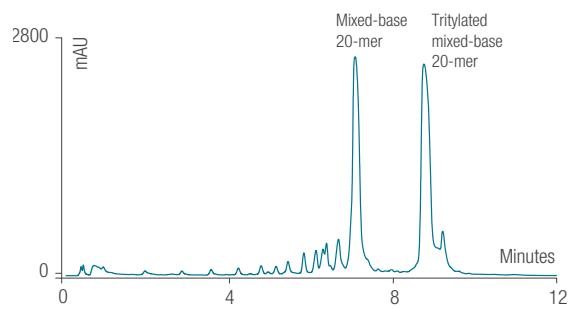
Additional reading

| Links | Description |
|---|-------------|
| Learn more thermofisher.com/biolc | |

Triylated oligonucleotide

DNA Swift SAX-1S, 150 x 5.0 mm

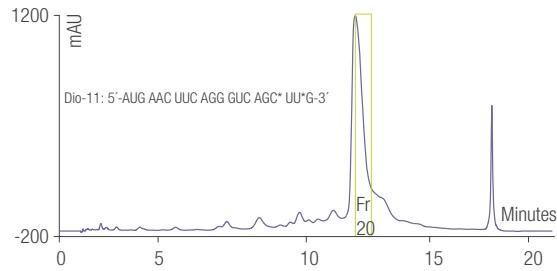
| | |
|------------------|-------------------------------|
| Flow rate | 1.5 mL/min |
| Mobile phase A | 15 mM Tris, pH 8 |
| Mobile phase B | 15 mM Tris, pH 8, 1.25 M NaCl |
| Temperature | 30 °C |
| Injection volume | 20 µL |
| Detection | UV at 260 nm |
| Gradient | 8–64% B in 10 min |



Purification of a 21-base RNA sample with aberrant 2'-5' linkages at the 1 and 3 positions from the 3' end

DNA Swift SAX-1S, 150 x 5.0 mm

| | |
|------------------|--------------------------------|
| Flow rate | 1.5 mL/min |
| Mobile phase A | 40 mM Tris, pH 7 |
| Mobile phase B | 40 mM Tris, pH 7 + 1.25 M NaCl |
| Temperature | 30 °C |
| Injection volume | 125 µg |
| Detection | UV at 260 nm |
| Gradient | 26–42% B in 10 column volumes |



DNA Swift SAX-1S column

Length (mm)

150

5.0 mm ID

[066766](#)