



Detection of protein heterogeneity by HPLC

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Introduction

Protein heterogeneity is generated by...

- Post-translational modification
phosphorylation (dephosphorylation), glycosylation, lipidation,
methylation, acetylation, protein splicing, ...
- Decomposition
proteolysis, deamidation, oxydation, ...
- Others
chemical modification (e.g. PEGylation), denaturation,
aggregation, ...



Introduction

Therapeutic proteins

- Therapeutic antibodies and recombinant proteins are now widely used for therapeutic treatment.
- Heterogeneity evaluation is essential during development, stability testing, and in the quality control of the final product.
- Analysis of the aggregates and denaturated proteins is also important because they might increase the risk of anaphylaxis or immunoreaction.



Contents

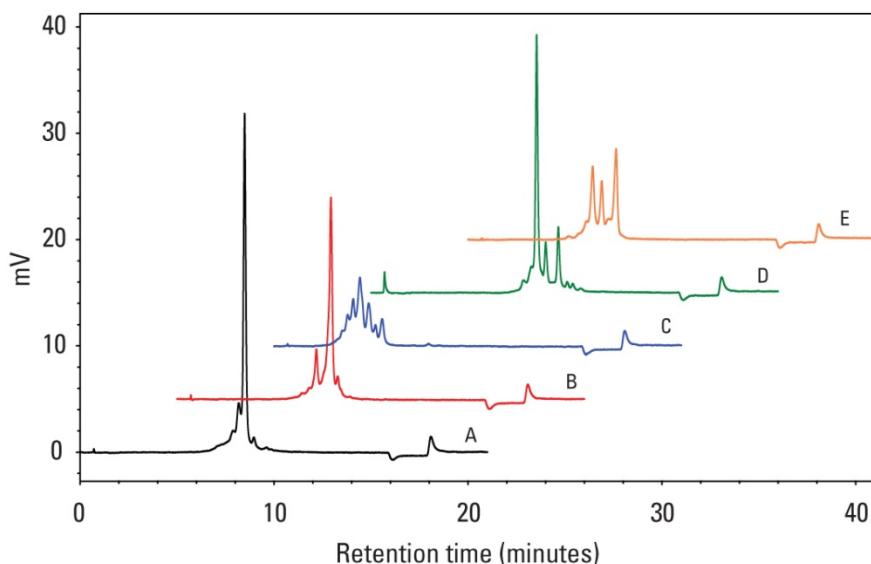
- 1. Ion Exchange Chromatography (IEC)**
- 2. Size Exclusion Chromatography (SEC)**
- 3. Hydrophobic Interaction Chromatography (HIC)**
- 4. Reversed Phase Chromatography (RPC)**



1. Ion Exchange Chromatography



Therapeutic antibody analysis (1) using a TSKgel CM-STAT column

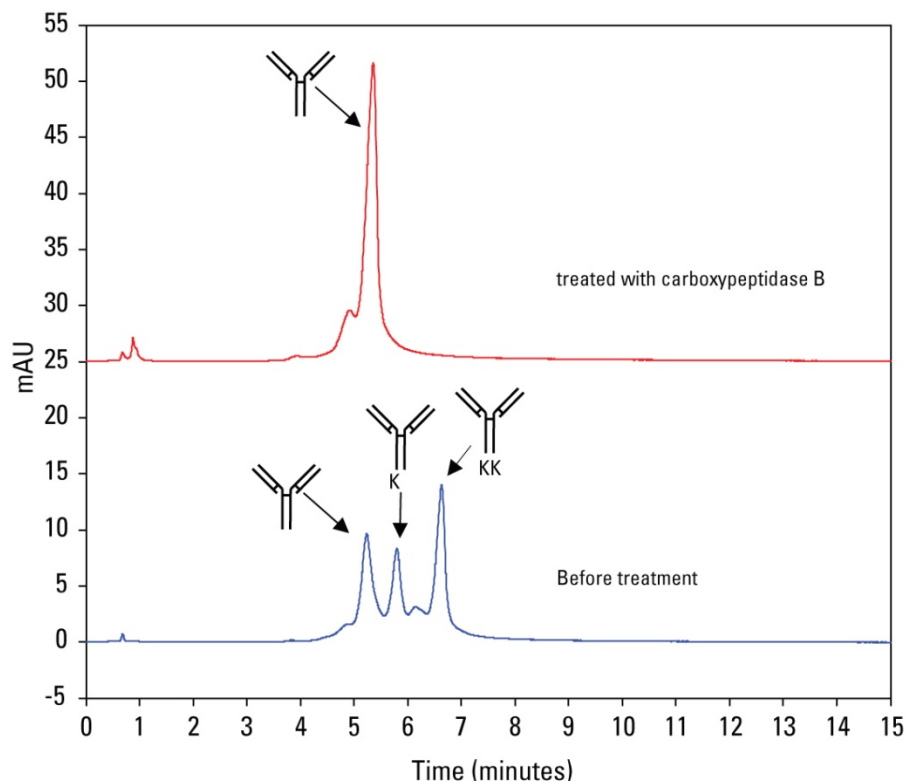


Column: TSKgel CM-STAT, 4.6 mm ID x 10 cm
Mobile phase A: 20 mmol/L MES, pH 6.0
Mobile phase B: 20 mmol/L MES + 0.5 mol/L NaCl, pH 6.0
Gradient: 10% B (0 min), 30% B (15 min), 100% B (15.1 min), 100% B (18 min), 10% B (18.1 min), 10% B (23 min)
Flow rate: 1.0 mL/min
Detection: UV@280 nm
Temperature: 25°C
Injection vol.: 20 µL
Concentration: 0.5 g/L
Sample: A: humanized, IgG₁
B: humanized, IgG₁
C: chimera, IgG₁
D: chimera, IgG₁
E: chimera, IgG₁
MES: 2-morpholinoethanesulfonic acid

High resolution analysis profiles of five antibodies were obtained on a TSKgel CM-STAT column. Each antibody shows multiple variants.



Therapeutic antibody analysis (2) using a TSKgel CM-STAT column

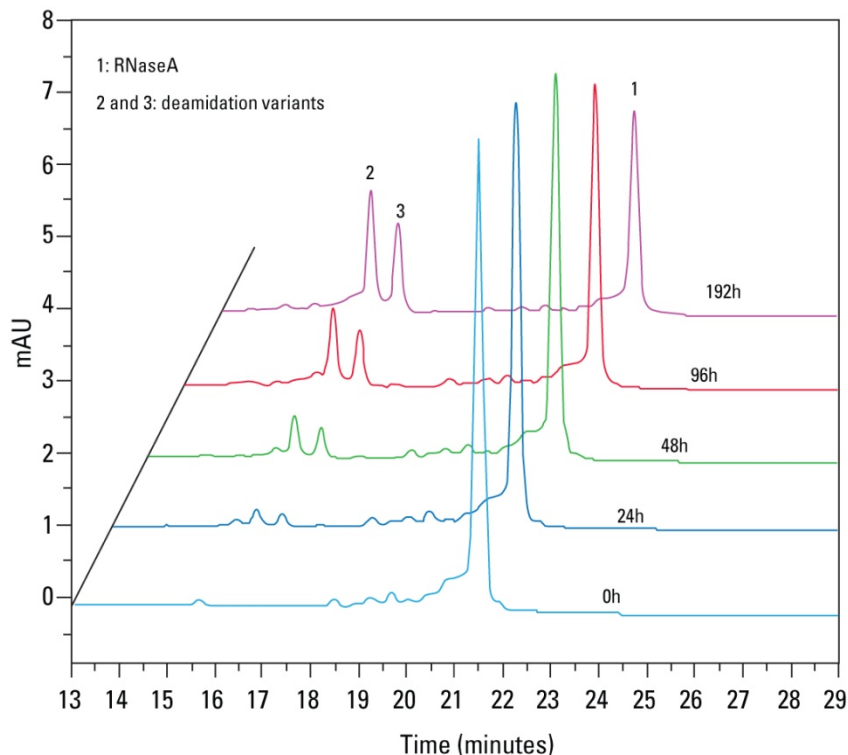


Column: TSKgel CM-STAT, 4.6 mm ID x 10 cm
Mobile phase A: 20 mmol/L MES, pH 6.0
Mobile phase B: 20 mmol/L MES + 0.5 mol/L NaCl, pH 6.0
Gradient: 10% B (0 min), 30% B (15 min)
100% B (15.1 min), 100% B (18 min),
10% B (18.1 min), 10% B (23 min)
Flow rate: 1.0 mL/min
Detection: UV@280 nm
Temperature: 25°C
Injection vol.: 20 µL
Concentration: 0.5 g/L
Sample: therapeutic antibody treated and
untreated with carboxypeptidase B
Procedure: To a 35 µL of therapeutic antibody (10 g/L),
1 µL of carboxypeptidase B (Sigma C9584,
140 U/mg protein, 5 g/L in PBS) was added
and incubated for 3 hours at 37°C. After
adding 664 µL of 20 mmol/L MES, pH 6.0,
to dilute the antibody concentration of 0.5 g/L,
20 µL of the diluted sample was injected.

The TSKgel CM-STAT column can detect even one amino residue difference.



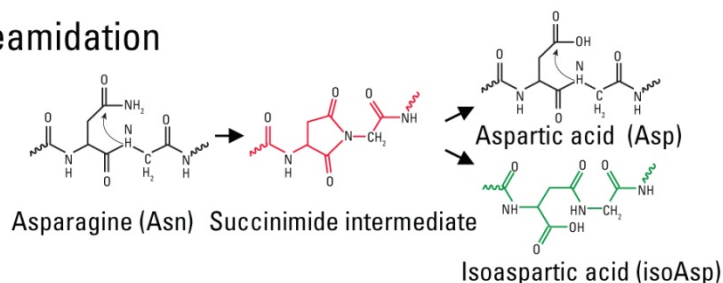
Deamidation of RNaseA using a TSKgel CM-STAT column



Column: TSKgel CM-STAT, 4.6 mm ID x 10 cm
 Mobile phase A: 20 mmol/L MES, pH 6.0
 Mobile phase B: 20 mmol/L MES + 1 mol/L NaCl, pH 6.0
 Gradient: 5% B (0 min), 25% B (30 min), 100% B (30 min), 100% B (34 min), 5% B (34 min), 5% B (40 min)
 Flow rate: 1.0 mL/min
 Detection: UV@280 nm
 Temperature: 25°C
 Injection vol.: 20 µL
 Concentration: 0.25 g/L

Procedure: RNaseA was dissolved at a concentration of 5 g/L in 1% ammonium carbonate buffer at pH 8.2 and incubated at 37°C for 0 to 192 hours. After incubation, diluted with 20 mmol/L MES, pH 6.0, at a concentration of 0.25 g/L and analyzed.

Deamidation



Note: Position 67 asparagine of the Ribonuclease A is well known of in the mild alkaline condition and generates 2 kinds of variants (Asp67 and isoAsp67).

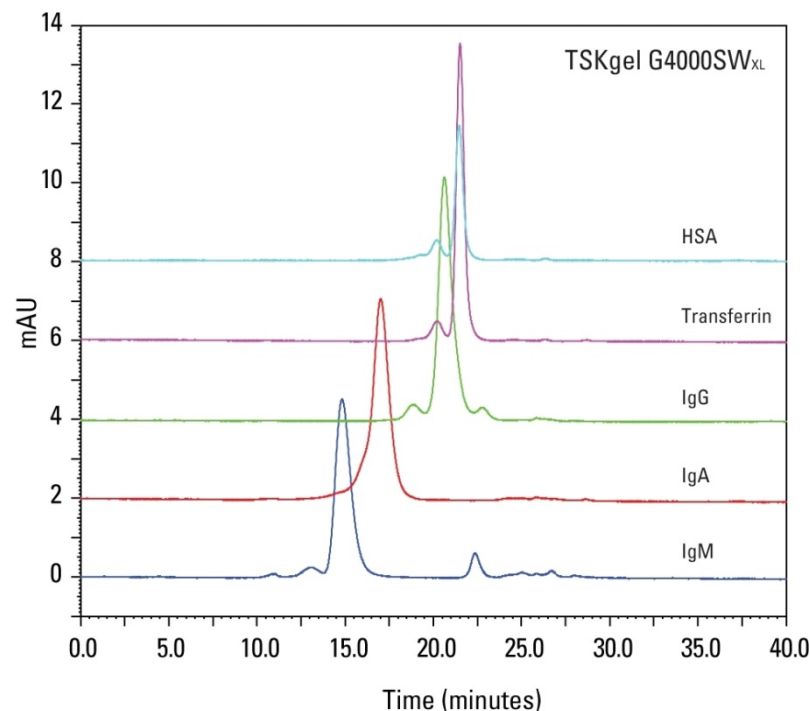
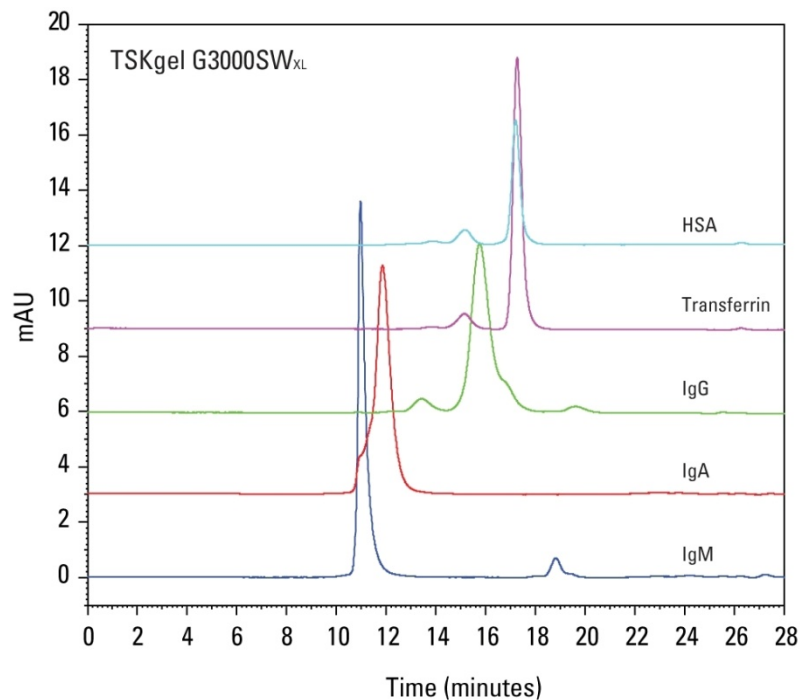
RNaseA and the two variants were completely separated using the TSKgel CM-STAT column.



2. Size Exclusion Chromatography



Separation of immunoglobulins using TSKgel G3000SW_{XL} and G4000SW_{XL} columns



Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm
TSKgel G4000SW_{XL}, 7.8 mm ID x 30 cm
Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl, pH 7.0
Flow rate: 0.5 mL/min
Detection: UV@280 nm
Temperature: 25°C
Injection vol.: 15 µL
Concentration: 0.17 g/L

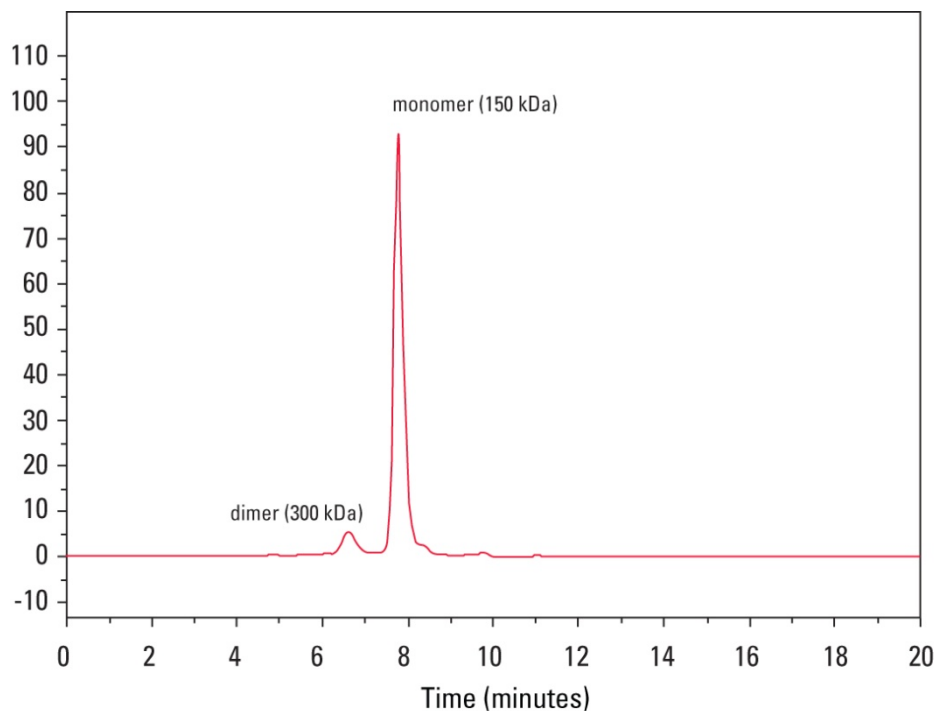
Samples:

1. human IgM (from myeloma, MW: 970 kDa)
2. human IgA (colostrum, MW: 390 kDa)
3. human IgG (MW: 150 kDa)
4. transferrin (MW: 80 kDa)
5. human serum albumin (MW: 66 kDa)

The separation of immunoglobulins and serum-related proteins was successfully analyzed using TSKgel SW_{XL} columns.



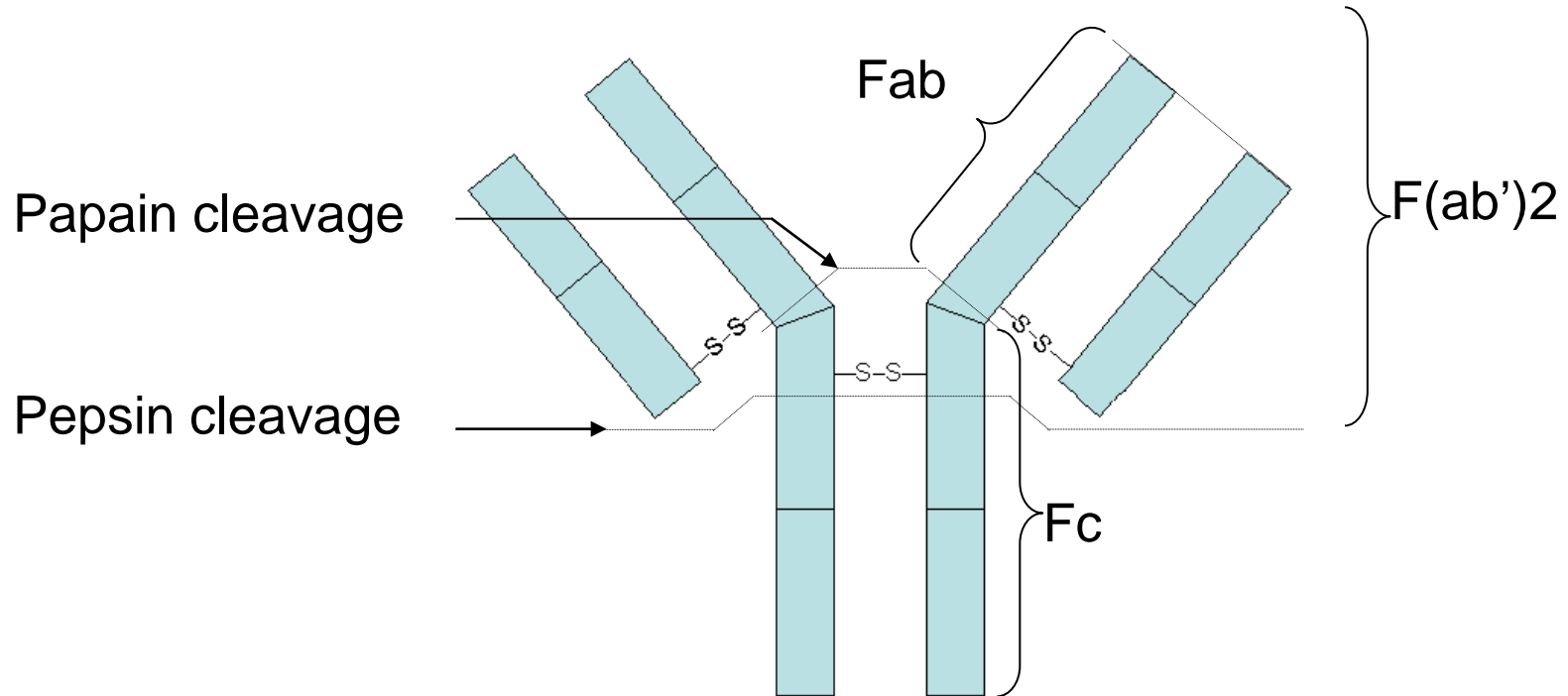
Aggregation analysis of an antibody using a TSKgel SuperSW3000 column



Column: TSKgel SuperSW3000, 4.6 mm ID x 30 cm
Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl, pH 7.0
Flow rate: 0.35 mL/min
Detection: UV@280 nm
Temperature: room temperature
Injection vol.: 5 μ L
Sample: human monoclonal antibody (4.6 g/L, IgG₁)

The monomer antibody (approximate molecular size of 150 kDa) and the dimer (molecular size of 300 kDa) are separated with the TSKgel SuperSW3000 column.

Structure of IgG₁

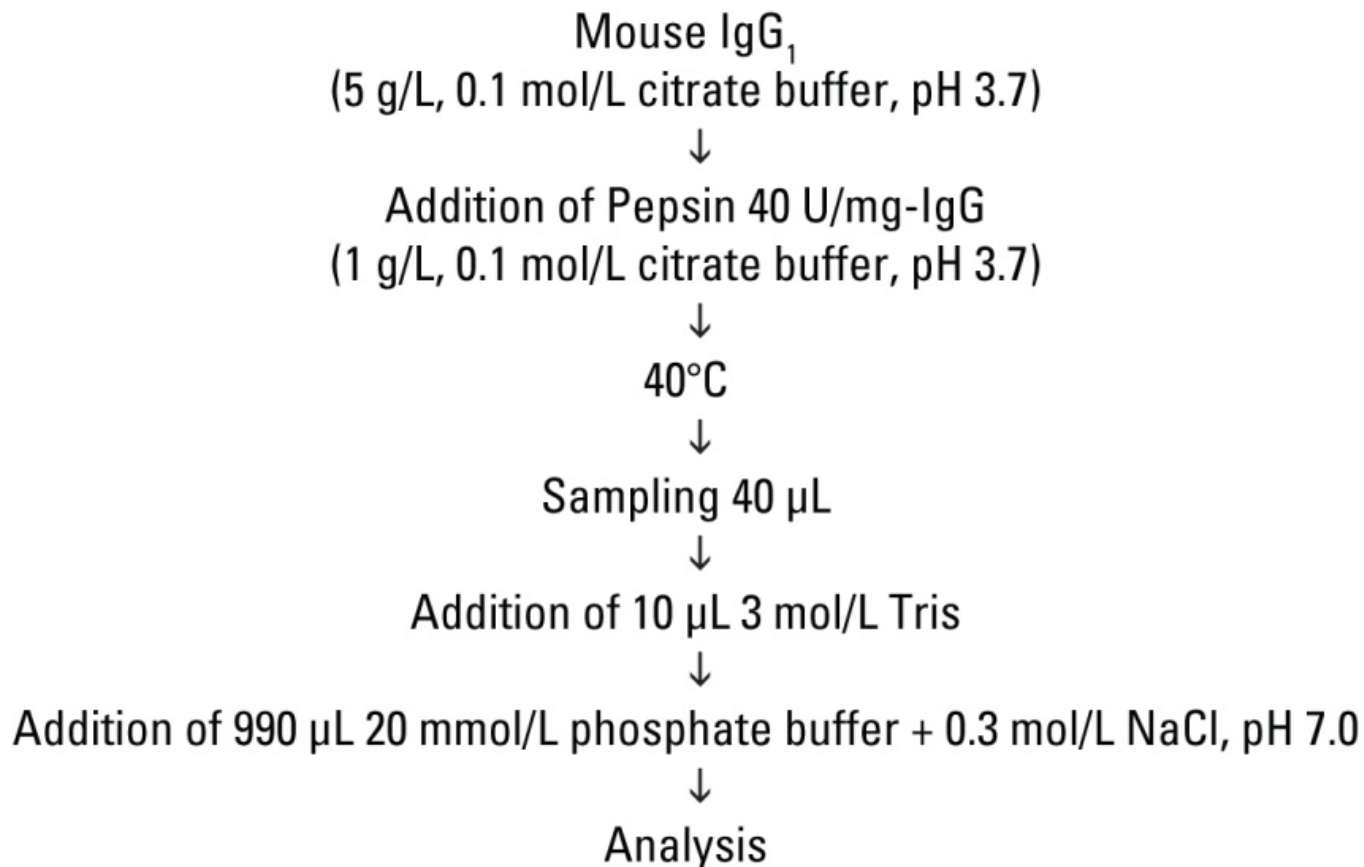


IgG is a relatively large molecule (approx. 150 kDa) and in order to improve the penetration to the tissue, fragmentation is carried out. Digestion with papain or pepsin is commonly applied to obtain antibody fragments without the loss of activity.

When papain is used for the antibody digestion, 2 Fab and 1 Fc are obtained from 1 antibody. When pepsin is used, a F(ab')₂ is obtained.



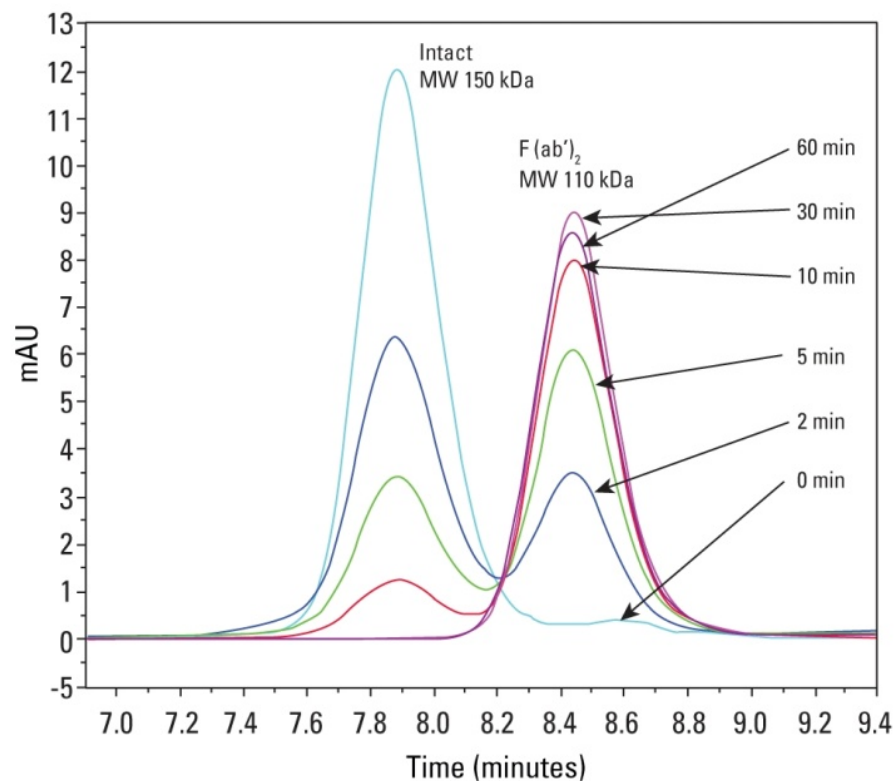
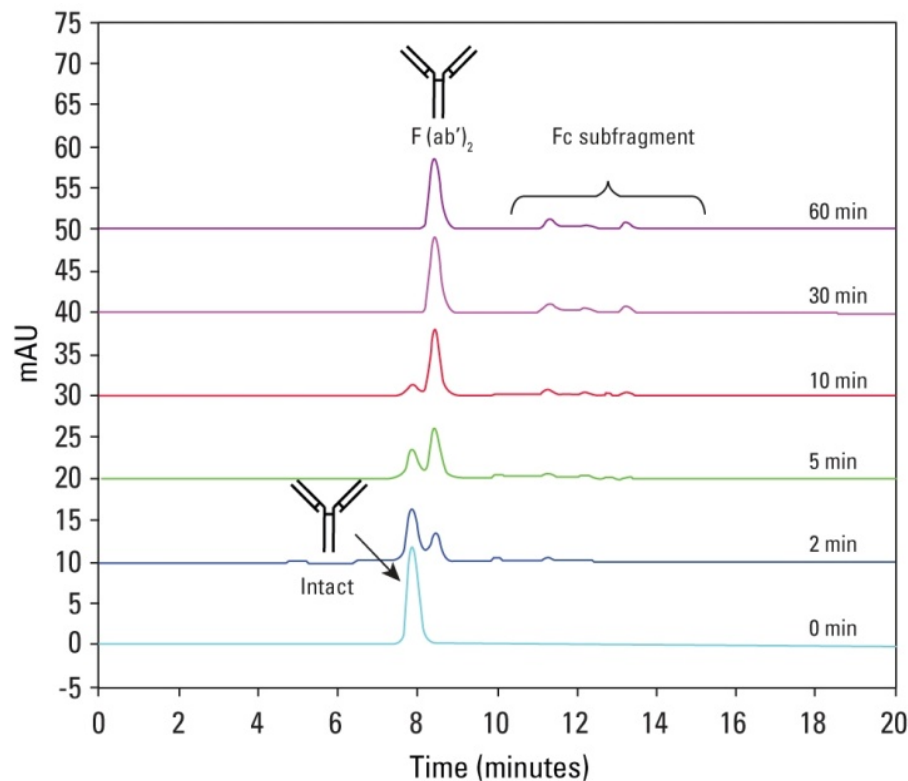
Pepsin digestion



Pepsin: from porcine gastric mucosa,
Sigma P7012, 2540 units/mg



Pepsin digestion using a TSKgel G3000SW_{XL} column

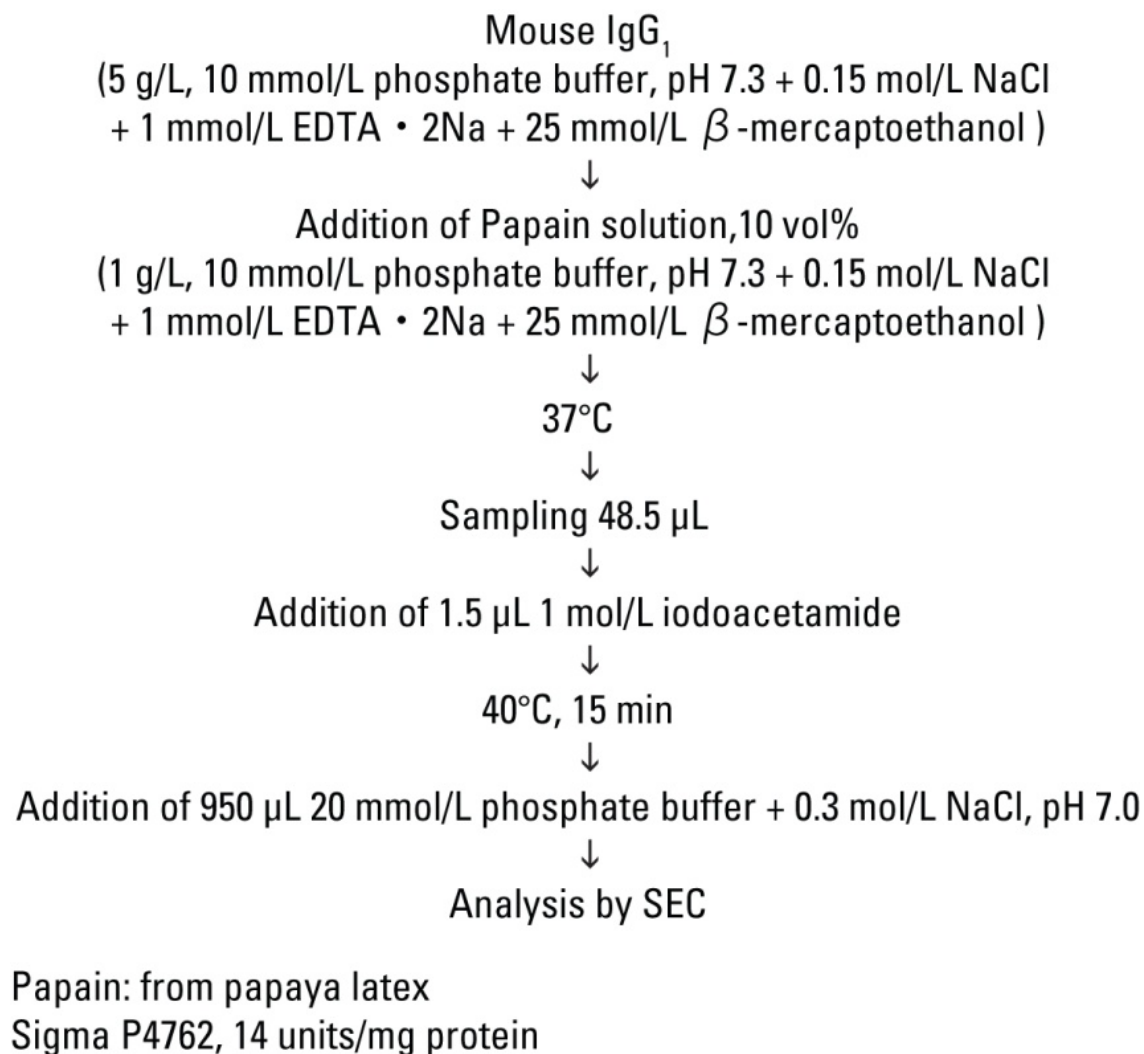


Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm
Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl, pH 7.0
Flow rate: 1.0 mL/min
Detection: UV@280 nm
Temperature: room temperature
Injection vol.: 10 μ L
Concentration: 0.19 g/L

As the reaction proceeds, the Fc fragment is further digested to a small subfragment so its peak is not seen, as is expected from its molecular size.

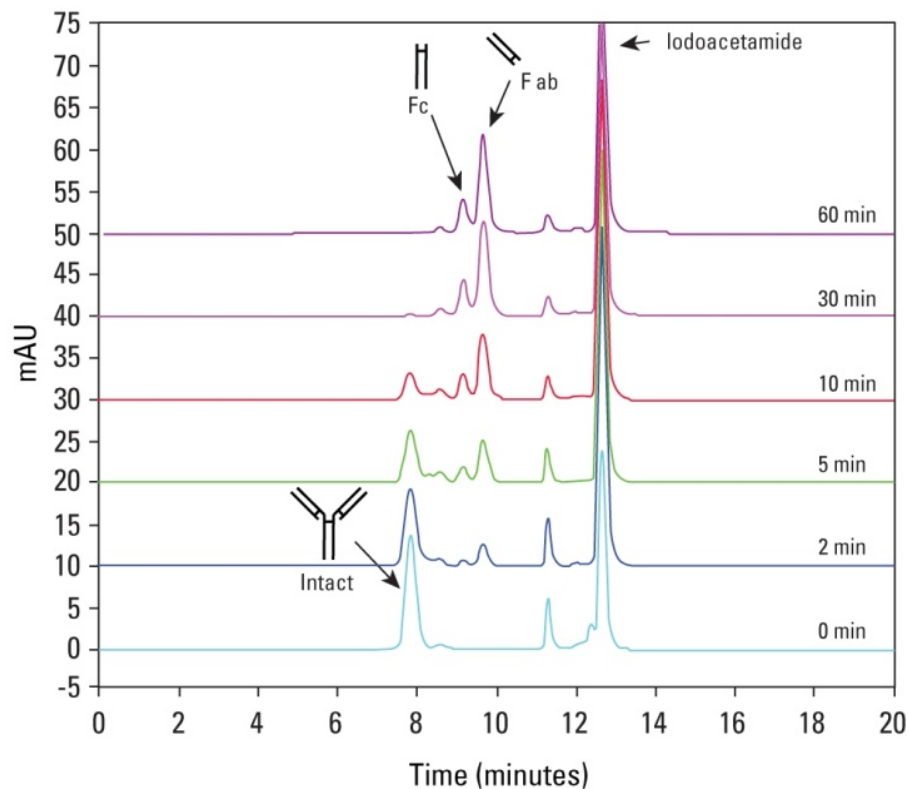


Papain digestion

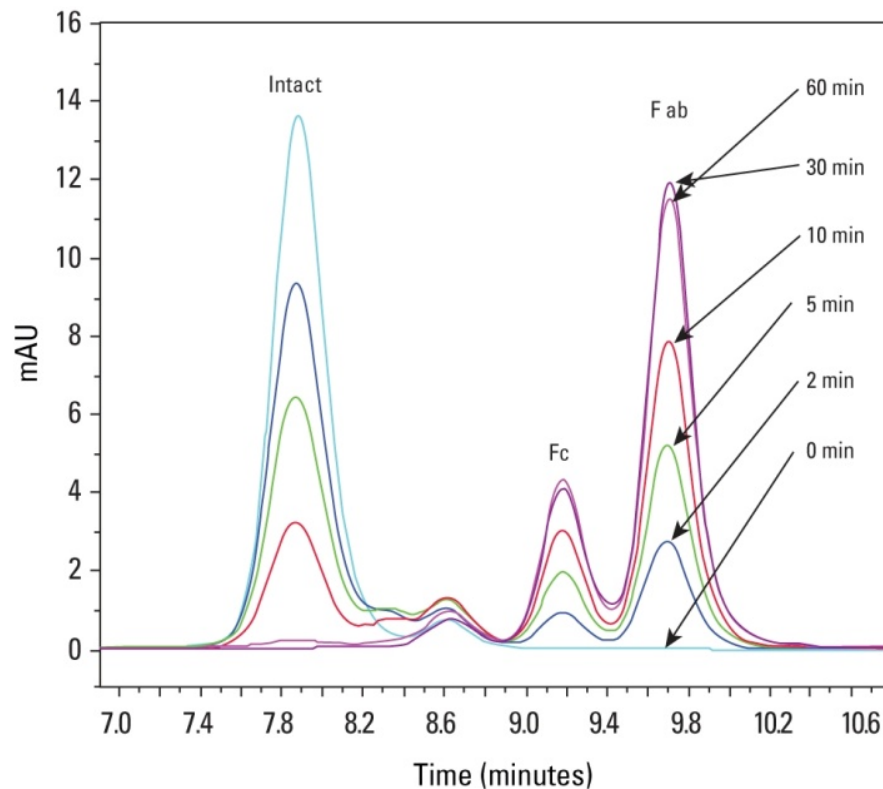




Papain digestion using a TSKgel G3000SW_{XL} column



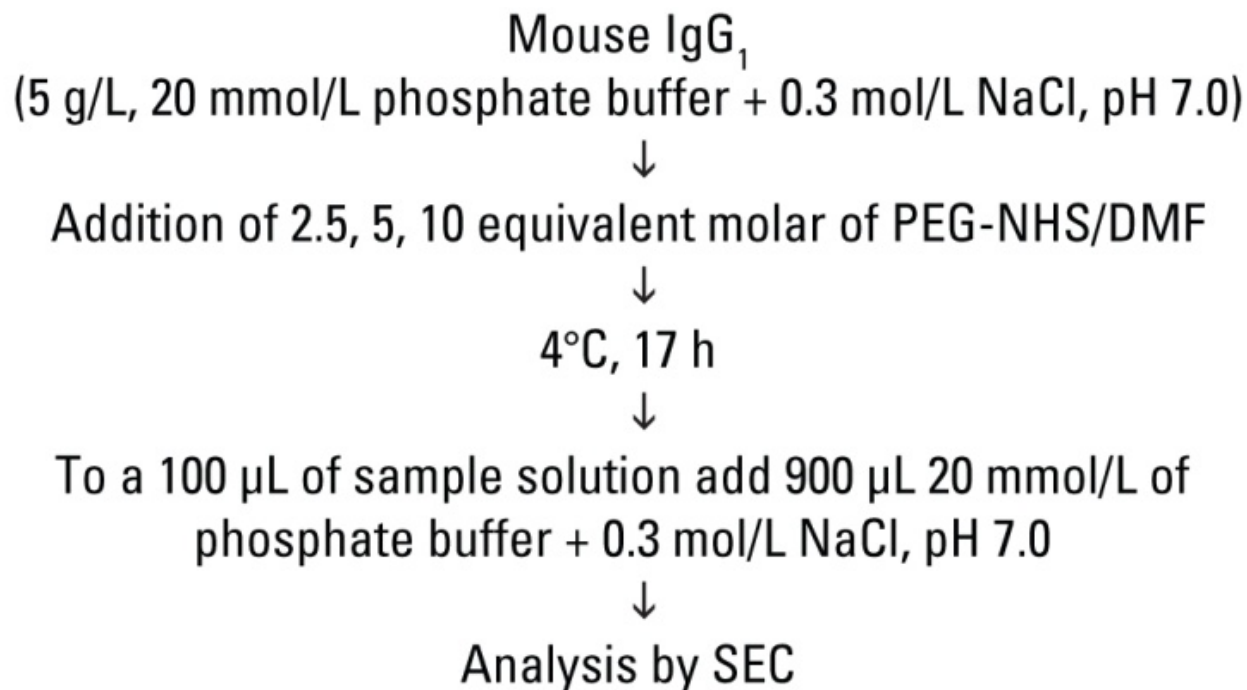
Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm
Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl, pH 7.0
Flow rate: 1.0 mL/min
Detection: UV@280 nm
Temperature: room temperature
Injection vol.: 10 μ L
Concentration: 0.24 g/L



As the reaction proceeds, the intact antibody reduces and Fab and Fc peaks become larger.



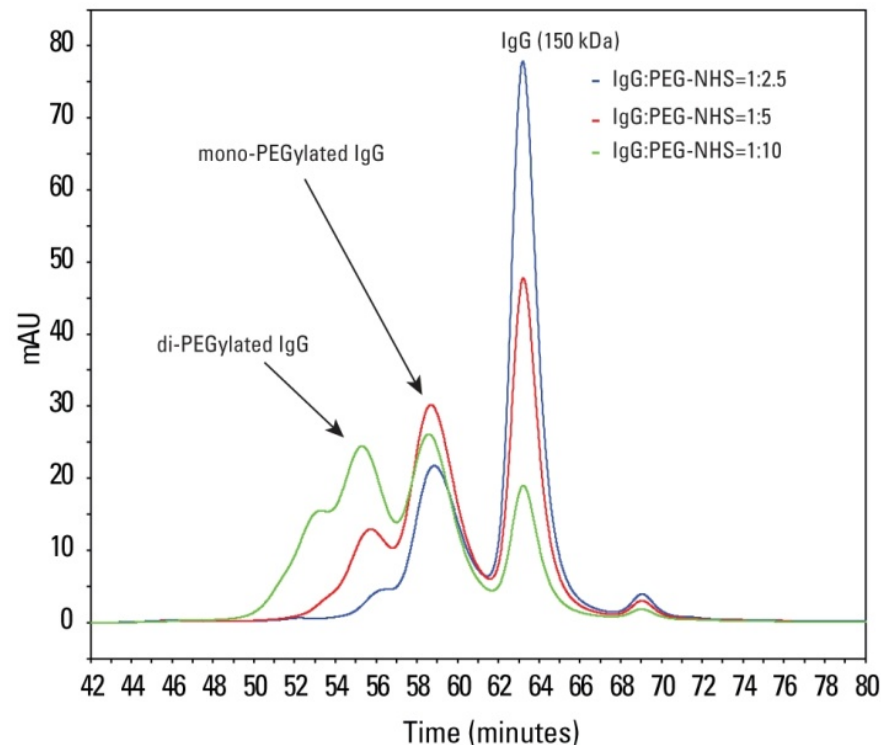
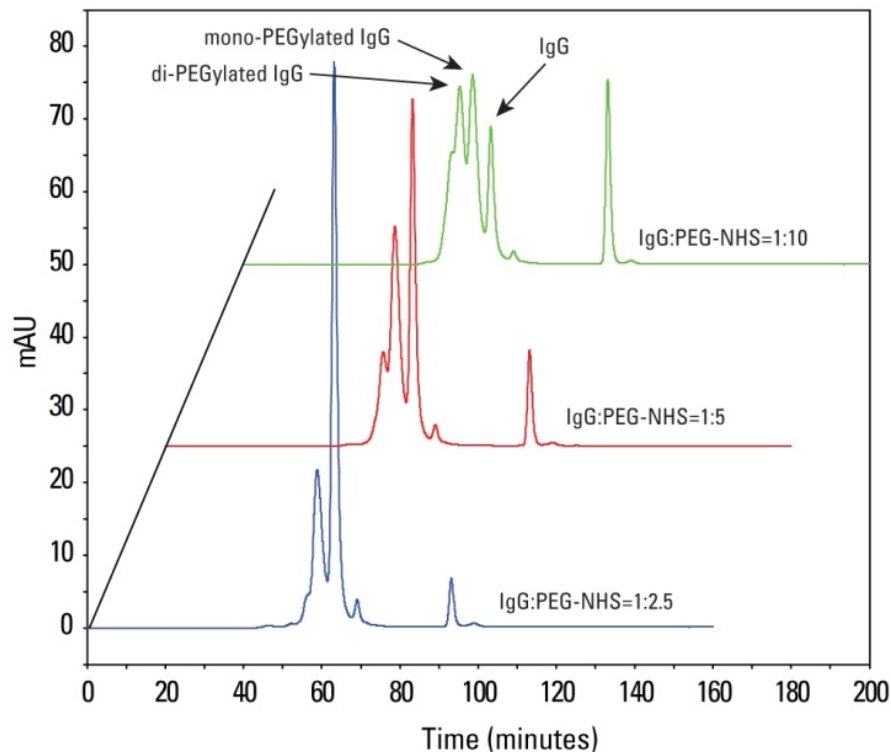
Separation of PEGylated protein (PEG:5000)



PEG-NHS: PEG, N-hydroxysuccinimidyl ester
SUNBRIGHT ME-050CS, MW: 5000, NOF CORPORATION, JAPAN



Separation of PEGylated protein (PEG:5000) using a TSKgel G3000SW_{XL} column



Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm,
2 columns
Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl,
pH 7.0
Flow rate: 0.25 mL/min
Detection: UV@280 nm
Temperature: 25°C
Injection vol.: 20 µL
Concentration: 0.5 g/L

From this analysis we can get information not only on the average number of PEG that were introduced to the antibody, but also the distribution of the PEG.

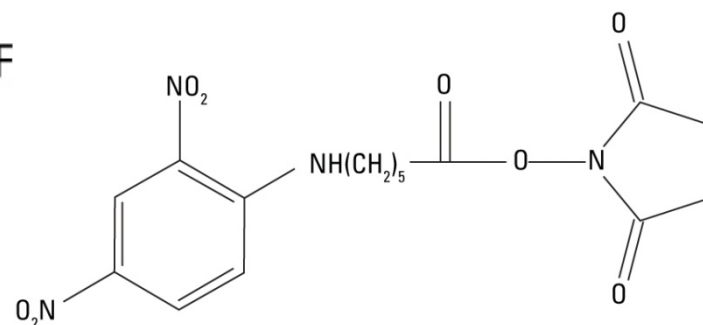


Addition of 1, 2, 4 equivalent molar of DNP-X-NHS/DMF

Purified by SEC to remove unreacted DNP

Addition of anti-DNP antibody (Fab)

Analysis by SEC



DNP-X-N-hydroxysuccinimidyl ester
MW: 394.34

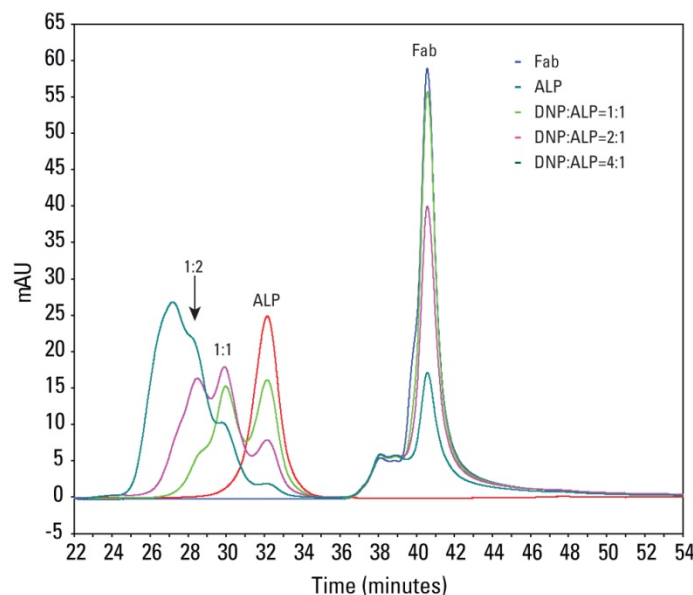
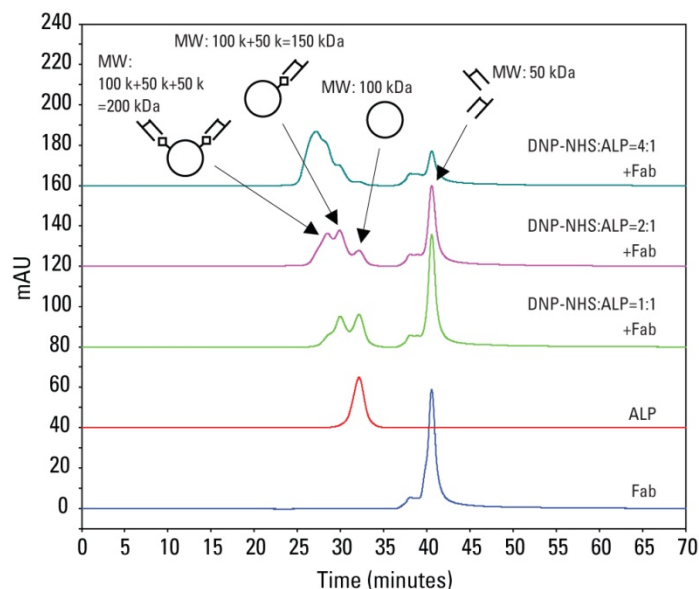
DNP-X-NHS: AnaSpec, Cat# 81228

Anti-DNP MAb: Cosmo Bio, Cat# LO-DNP-61

Antibody was papain digested and Fab was purified by SEC before use



Analysis of hapten-conjugated protein using a TSKgel G3000SW_{XL} column



Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm
 Mobile phase: 20 mmol/L phosphate buffer + 0.3 mol/L NaCl, pH 7.0
 Flow rate: 0.25 mL/min
 Detection: UV@280 nm
 Temperature: 25°C
 Injection vol.: 20 µL
 Concentration: DNP-ALP: 0.67 g/L, anti-DNP Fab: 1 g/L

Mixed ratio of DNP-NHS:ALP	Ratio (%)			
	DNP introduced (number)			
	0	1	2	≥3
1:1	61%	32%	7%	0%
2:1	30%	40%	22%	8%
4:1	7%	19%	30%	44%

Because the molecular weight of hapten is only about 300 Da compared to the 100 kDa weight of alkaline phosphatase, molecular size doesn't change enough to change the retention time of alkaline phosphatase when the hapten has conjugated to the protein.

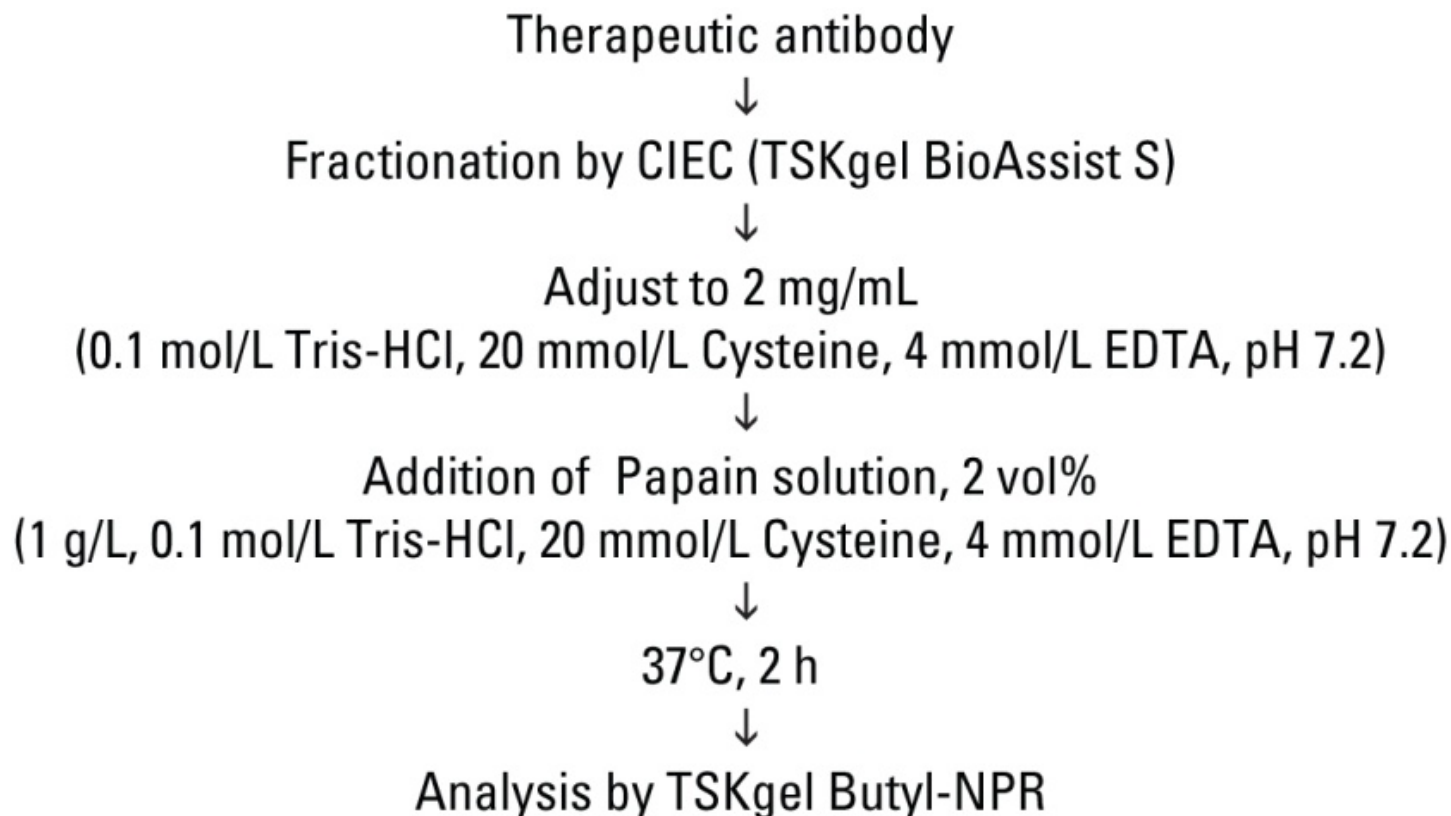
By adding anti-hapten Fab, in this case anti-DNP Fab, Fab binds to the hapten of the alkaline phosphatase surface. As a result, it shows a large molecular size increase, allowing hapten-conjugated and non-conjugated alkaline phosphatase to be separated by SEC.



3. Hydrophobic Interaction Chromatography

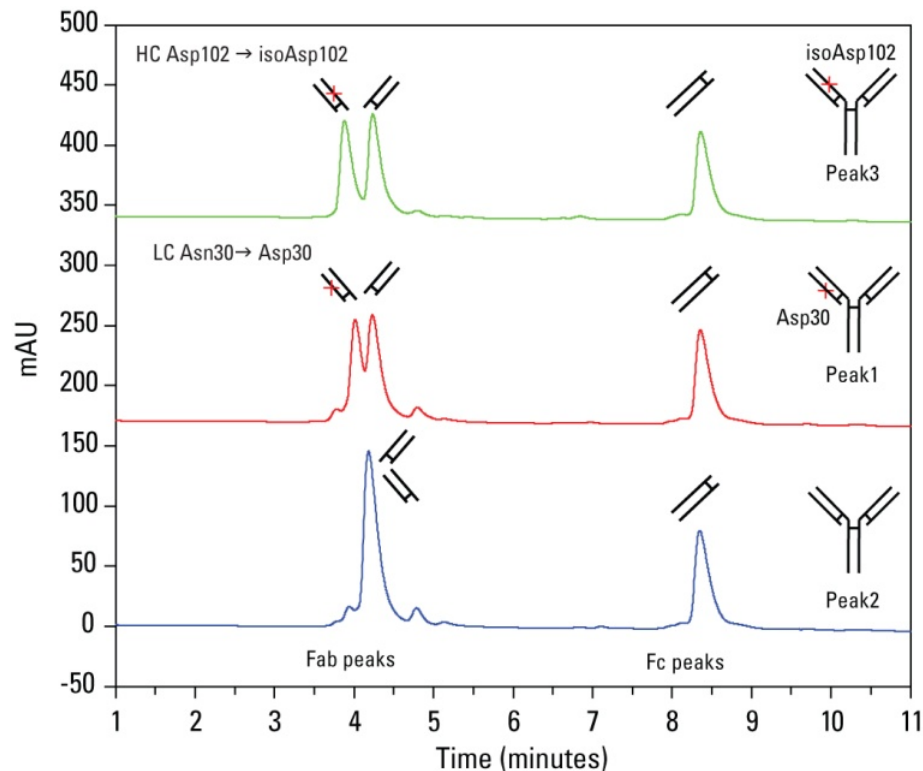


Analysis of therapeutic antibody (Papain digested)

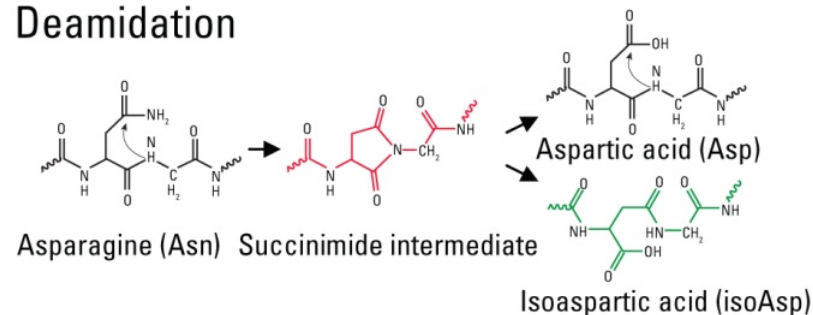


Papain: from papaya latex
Sigma P4762, 14 units/mg protein

Analysis of therapeutic antibody (Papain digested) using a TSKgel Butyl-NPR column



Deamidation



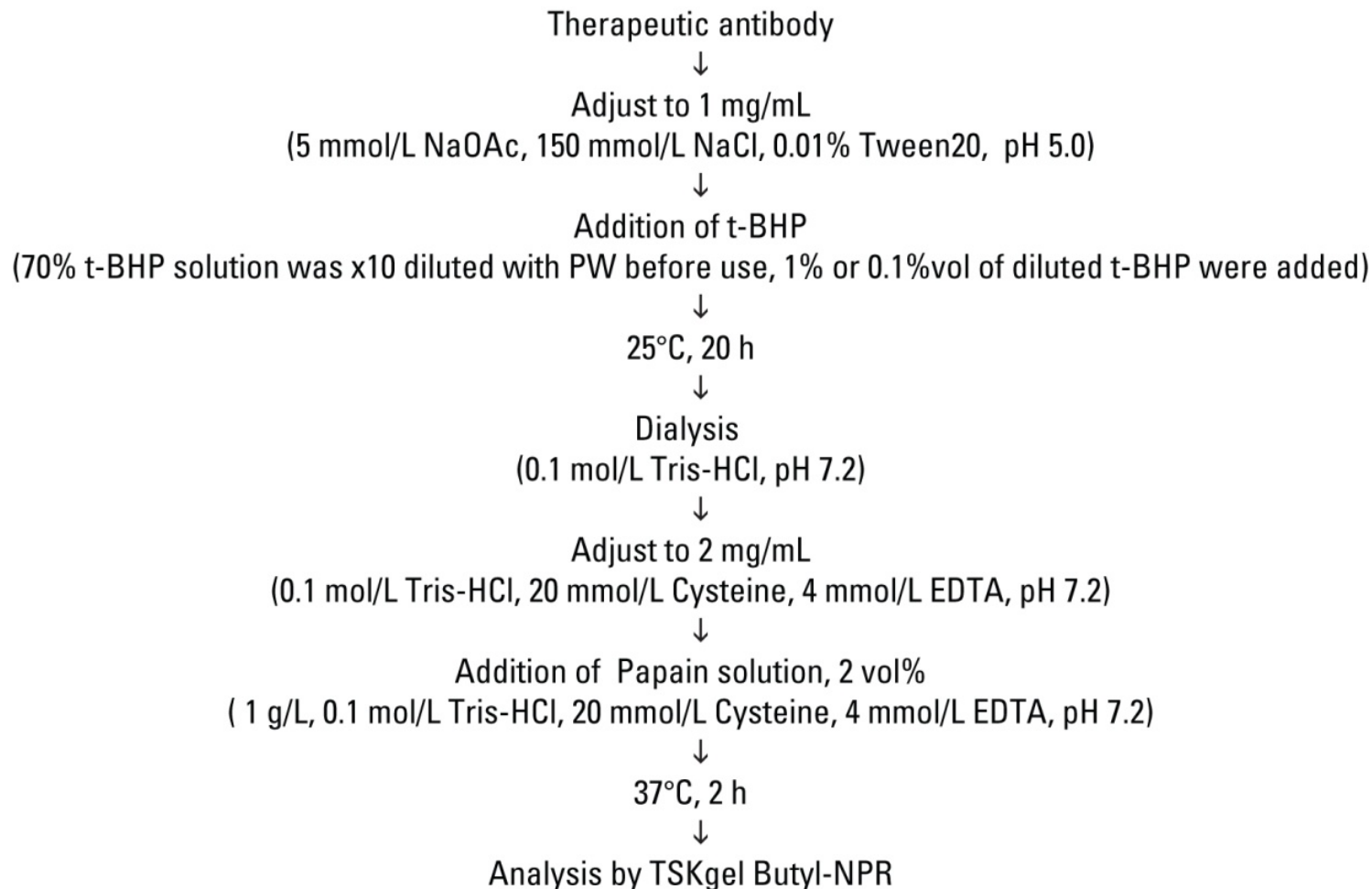
Column: TSKgel Butyl-NPR, 4.6 mm ID x 3.5 cm
 Mobile phase A: 20 mmol/L phosphate buffer containing 2 mol/L $(\text{NH}_4)_2\text{SO}_4$, pH 7.0
 Mobile phase B: 20 mmol/L phosphate buffer, pH 7.0
 Gradient: 10 min linear gradient from 25 to 60% B
 Flow rate: 1.0 mL/min
 Detection: UV@214 nm
 Temperature: 25°C
 Injection vol.: 2 μL (4 μg)

Peak	Structural difference	at LC Asn30	at HC Asn55	at HC Asp102
1	Deamidated (to Asp) at Asn30 of one light chain	Asn/Asp	Asn/Asn	Asp/Asp
2	Main peak form	Asn/Asn	Asn/Asn	Asp/Asp
3	Isomerized (to isoAsp) at Asp102 of one heavy chain	Asn/Asn	Asn/Asn	Asp/isoAsp

TSKgel Butyl-NPR is also a superior tool for detecting even one residue difference of the proteins.



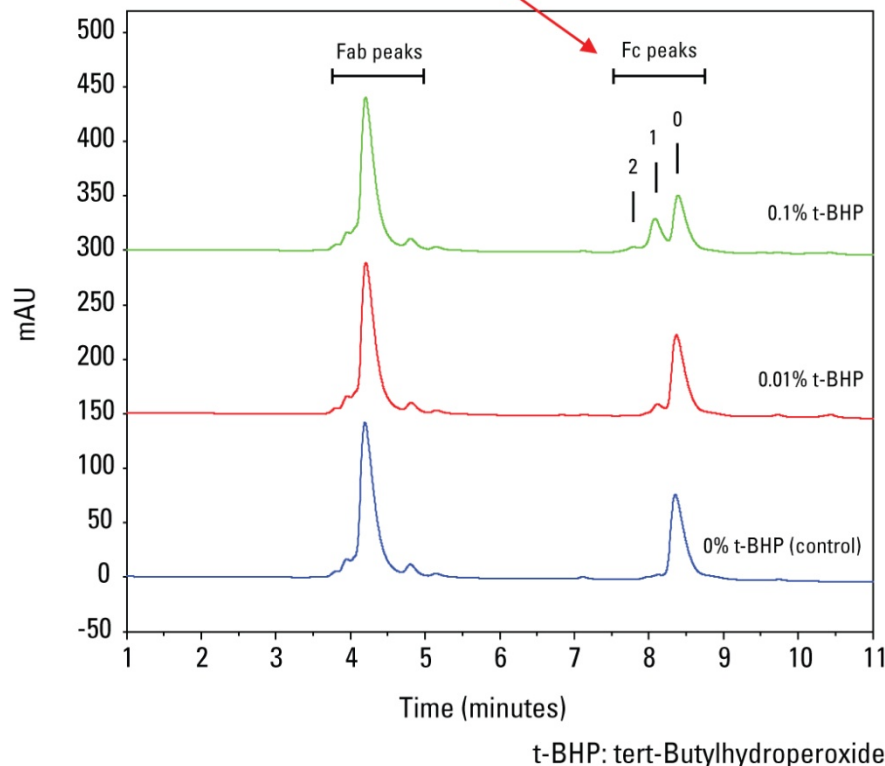
Analysis of therapeutic antibody (Papain digested)



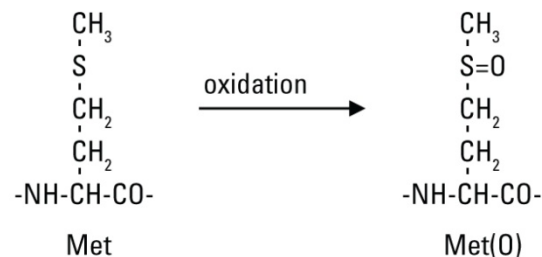


Analysis of therapeutic antibody (Papain digested) using a TSKgel Butyl-NPR column

Number of oxydated methionine (Met-255 → Met(O)-255
and/or Met-431 → Met(O)-431)



Methionine oxidation



Column: TSKgel Butyl-NPR, 4.6 mm ID x 3.5 cm
Mobile phase A: 20 mmol/L phosphate buffer containing 2 mol/L $(\text{NH}_4)_2\text{SO}_4$, pH 7.0
Mobile phase B: 20 mmol/L phosphate buffer, pH 7.0
Gradient: 10 min linear gradient from 25 to 60% B
Flow rate: 1.0 mL/min
Detection: UV@214 nm
Temperature: 25°C
Injection vol.: 2 μL (4 μg)

Therapeutic antibody was oxidated by incubating the antibody solution with t-BHP and analyzed with TSKgel Butyl-NPR after papain digestion.

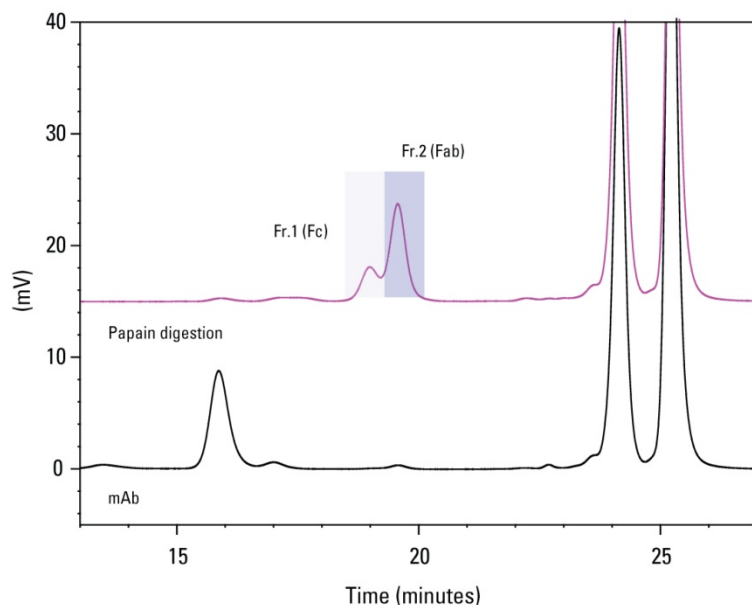
Methione residue of Fc was oxidated, and as a result multiple peaks were observed. Whereas no change was observed with Fab peaks.



4. Reversed Phase Chromatography

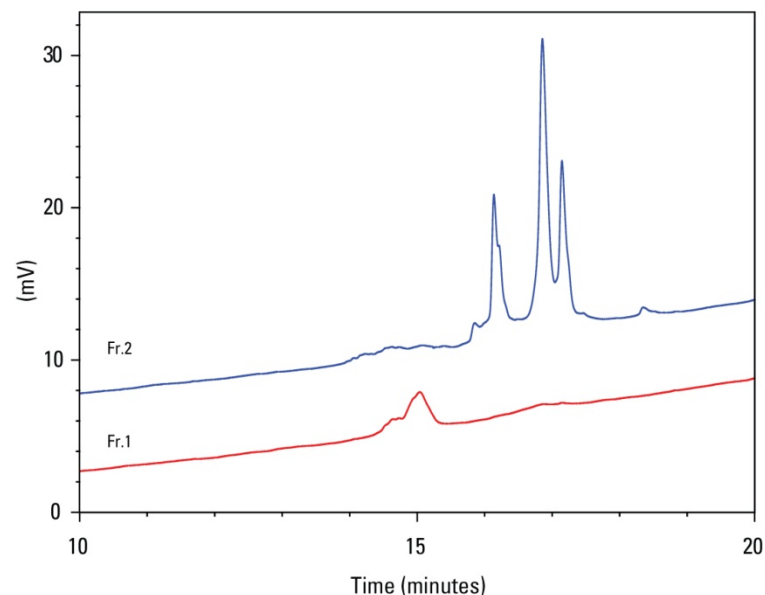


Analysis of antibody fragment using a TSKgel Protein C4-300 column



Conditions for SEC

Column: TSKgel G3000SW_{XL}, 7.8 mm ID x 30 cm x 2
Mobile phase: 20 mmol/L phosphate buffer, pH 7.0 + 0.3 mol/L NaCl
Flow rate: 1.0 mL/min
Temperature: 25°C
Sample: monoclonal antibody (human IgG₁)



Conditions for RPC

Column: TSKgel Protein C4-300, 4.6 mm ID x 15 cm
Mobile phase A: 0.05% TFA in water
Mobile phase B: 0.05% TFA in acetonitrile
Gradient: 5% B (0 min), 50% B (20 min)
Flow rate: 1.0 mL/min
Temperature: 70°C
Sample: monoclonal antibody (human IgG₁)

Human antibody was papain digested and separated with SEC. Two fractions were obtained and each fraction was analyzed with the TSKgel Protein C4-300 column. Three peaks were observed with the analysis of Fab.



Conclusions

- Therapeutic antibodies and recombinant proteins are now widely used for therapeutic treatment and evaluation of their heterogeneity is essential for the development, stability testing, and in the quality control of the final product.
- High resolution TSKgel nonporous resin columns for Ion Exchange Chromatography and Hydrophobic Interaction Chromatography can detect even one residue difference of the proteins.
- Size exclusion chromatography is suitable for detecting aggregates, fragments, and PEGylated proteins.
- The reversed phase chromatography column, TSKgel Protein C4-300, which has a large pore size of 300 Å, is applicable for the protein analysis.

Generous support was received from Dr. Hitoshi Kakidani, Sagami Chemical Research Institute.