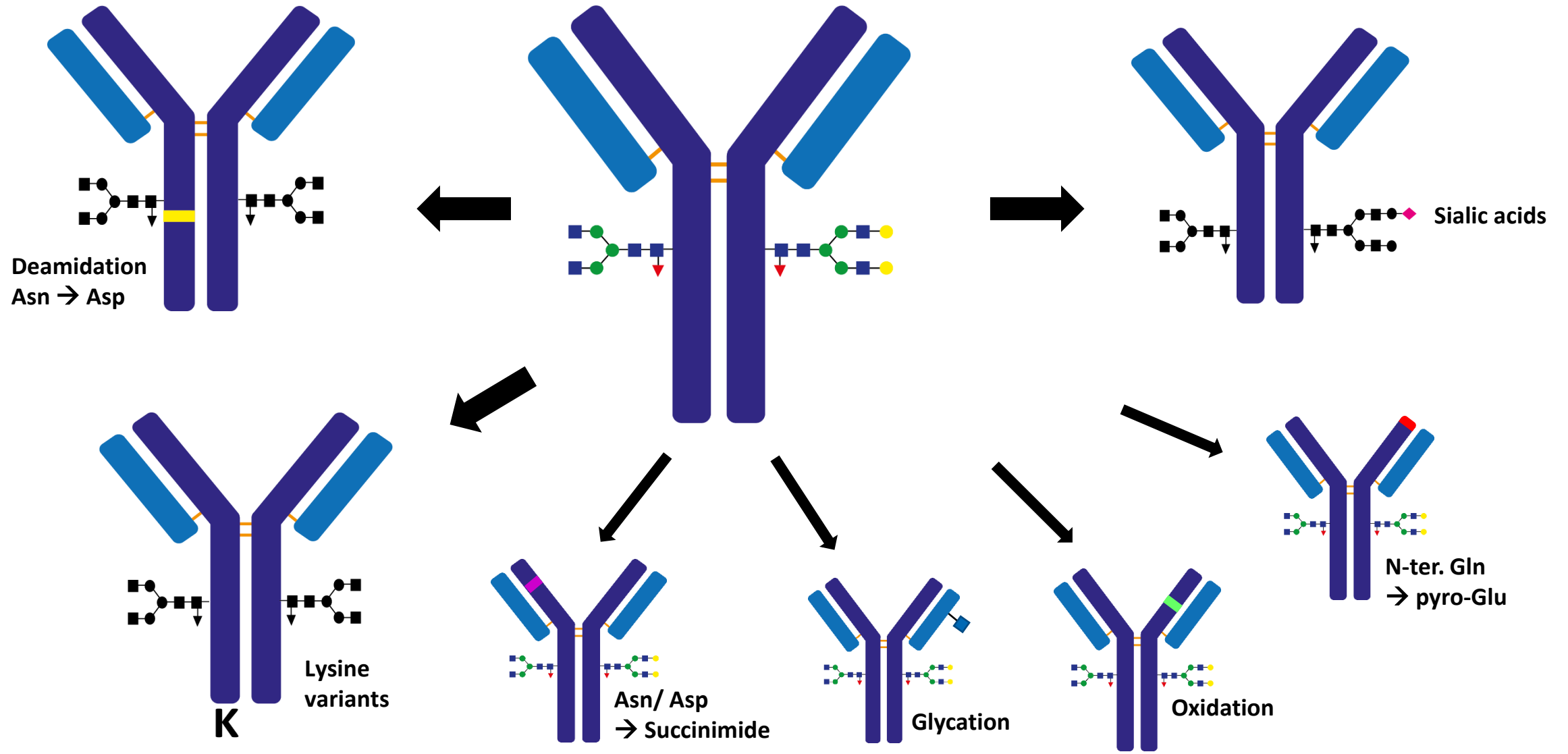


High-Performance Native Separation of mAb Proteoforms by CZE-MS under Native and Denaturing nanoESI Conditions for Ultrasensitive Charge Variant Characterization

Ann-Katrin Schwenzer, Christian Neusüß

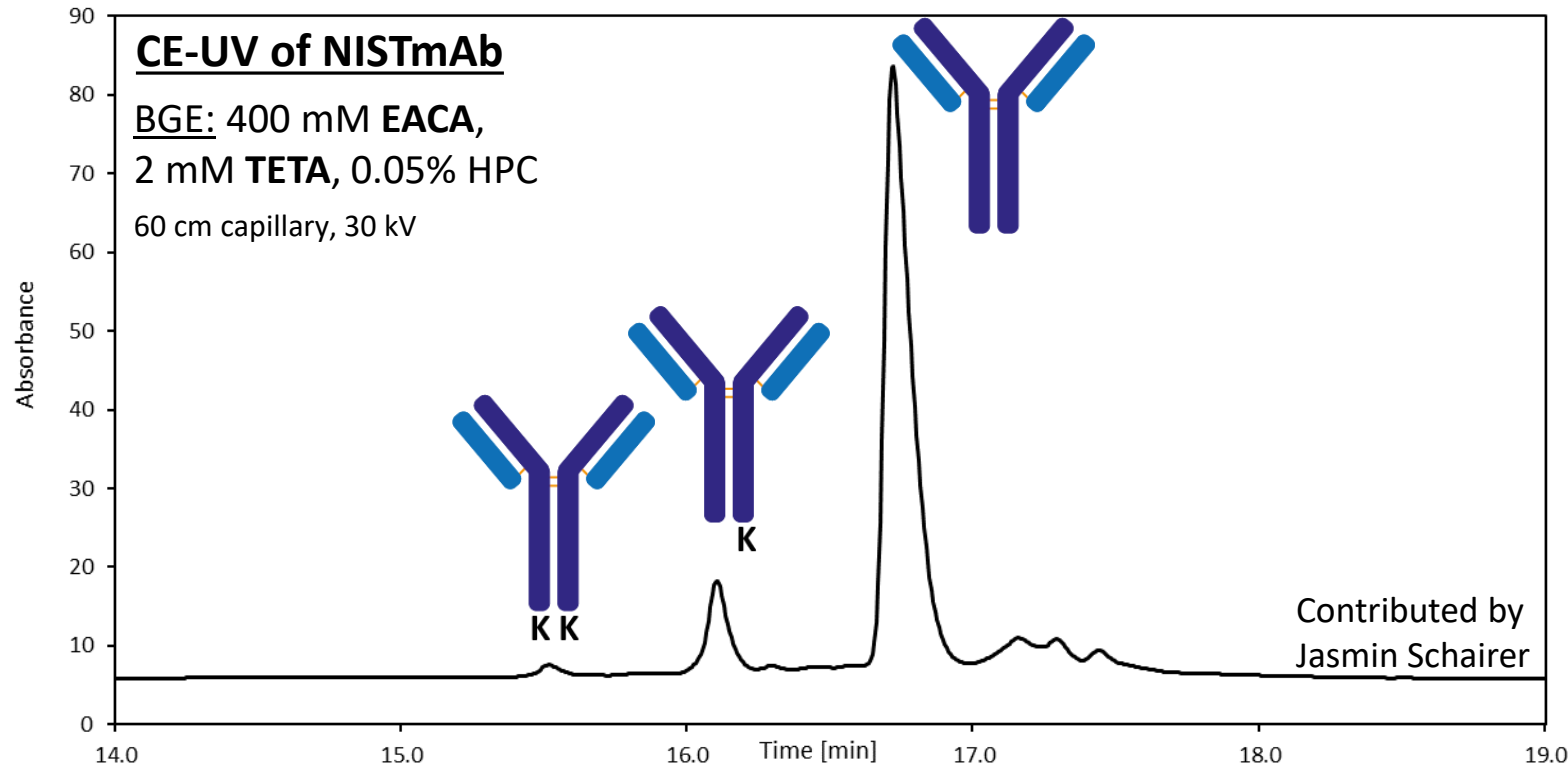
Aalen University, Beethovenstraße 1, 73430 Aalen

Therapeutic antibodies – a mixture of proteoforms



Some **charge variants** might influence **potency** or **safety** of the biopharmaceutical → should be **monitored**

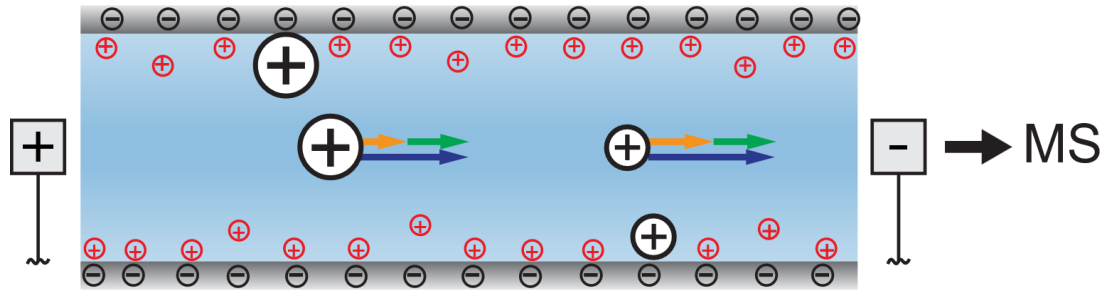
- Commonly applied **CZE-UV** method for charge variant separation involves **EACA** and **TETA** in the electrolyte:



→ Very good separation but not MS compatible (non-volatile electrolyte)

➡ How can we do CZE-MS analysis of mAb charge variants?

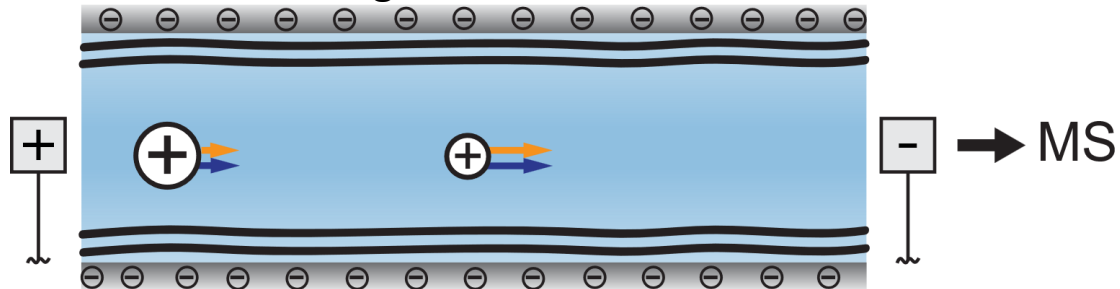
- **Uncoated:**



→ Decreases, broadens, or even eliminates analyte signals 👎



- **Neutral coating:**

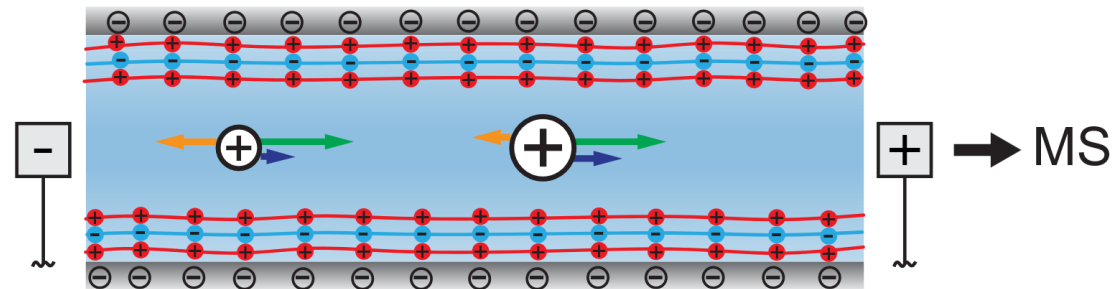


→ Reduction of protein interactions with capillary wall 👍

→ Supression of the EOF



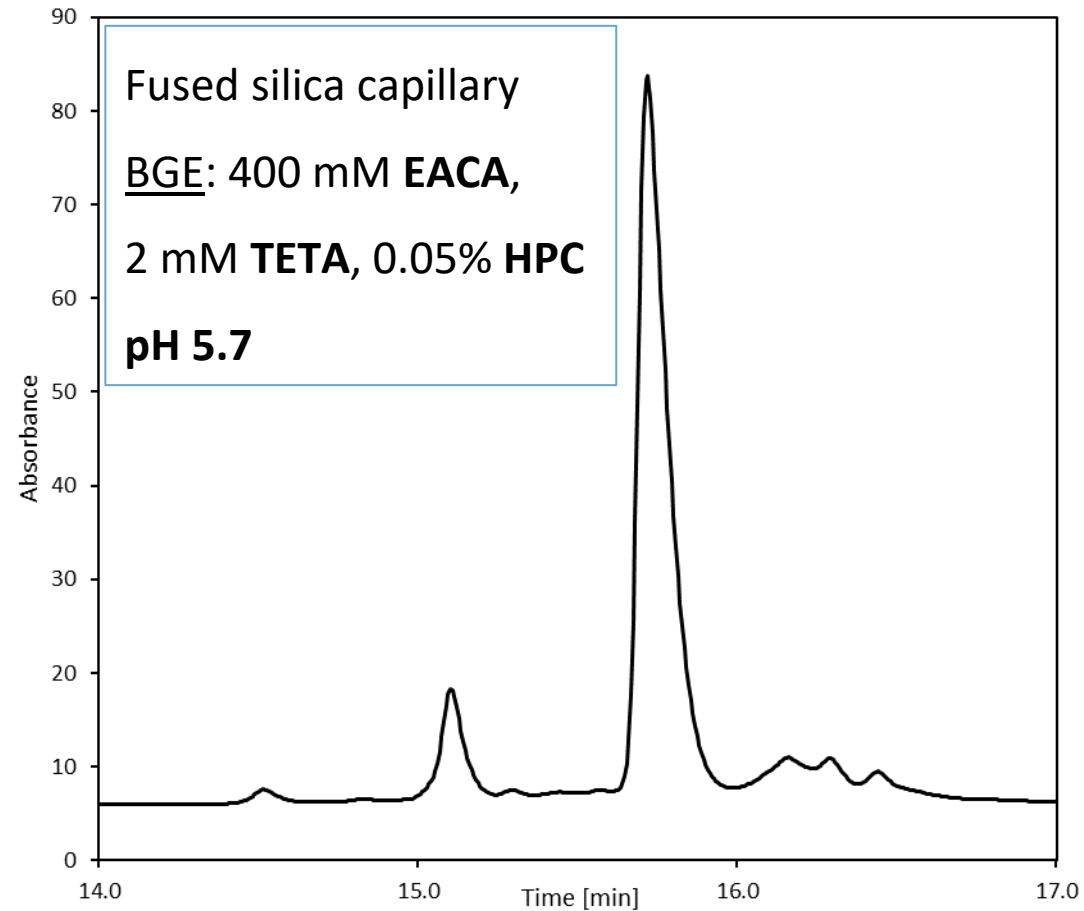
- **Cationic coating:**



→ Reduction of protein interactions with capillary wall 👍

→ Reversed EOF

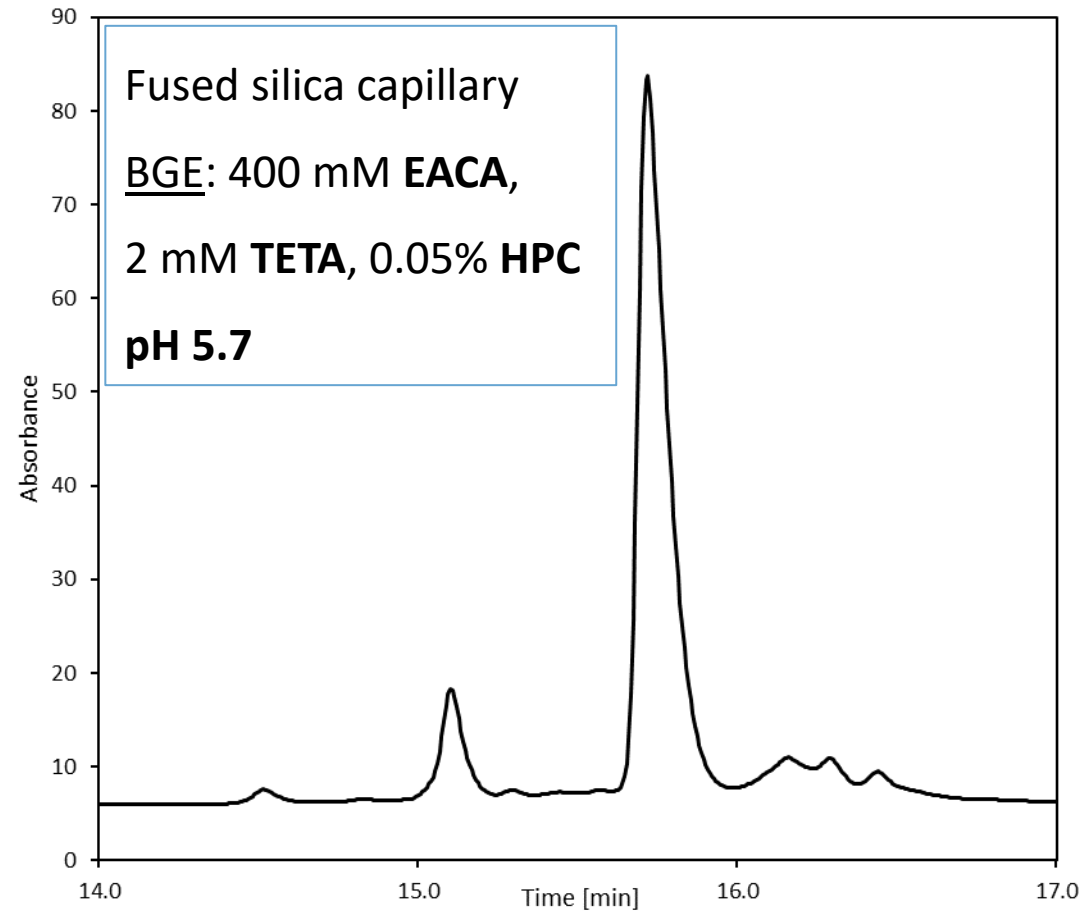
EOF → Analyte mobility → Apparent mobility →



→ Efficient separation of charge variants ✓

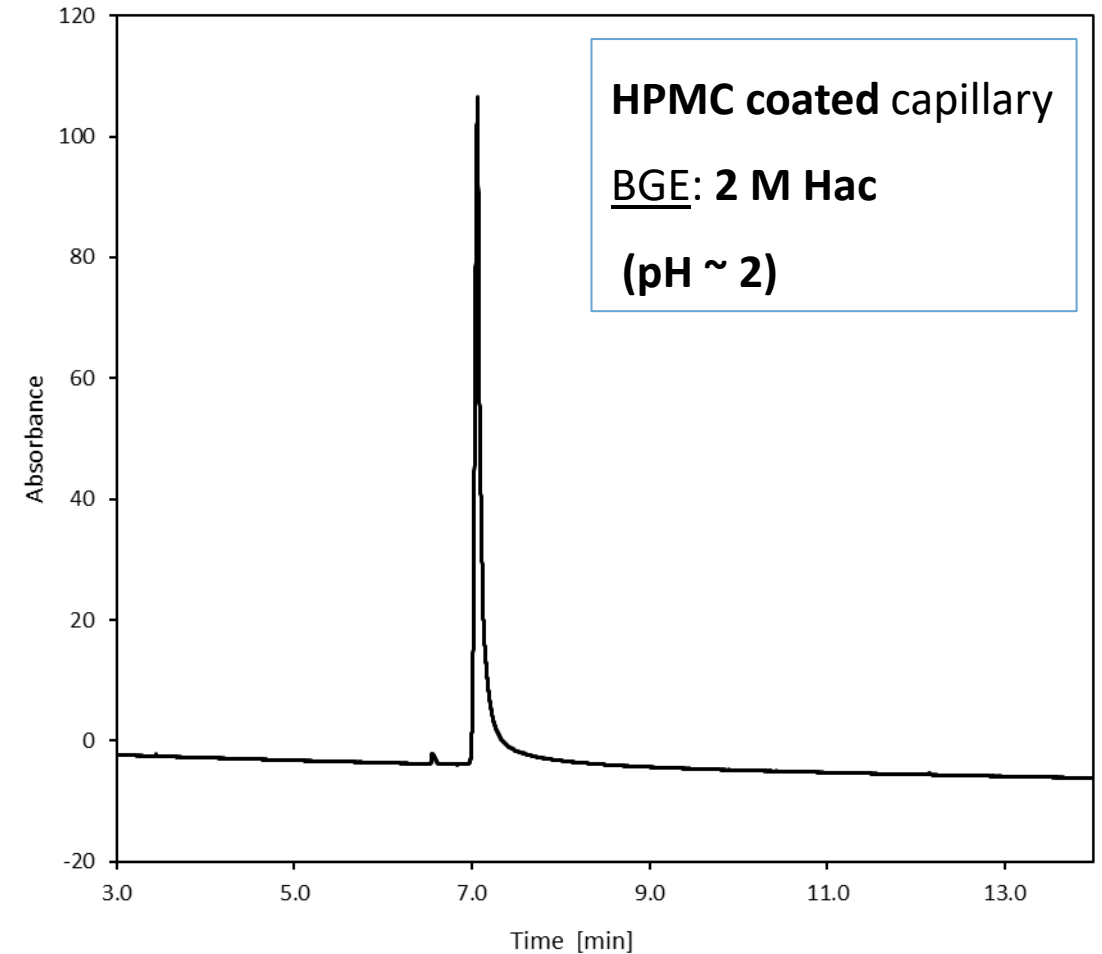
→ But: **Not** MS compatible ✗

CE-UV of NISTmAb using different electrolytes



→ Efficient separation of charge variants ✓

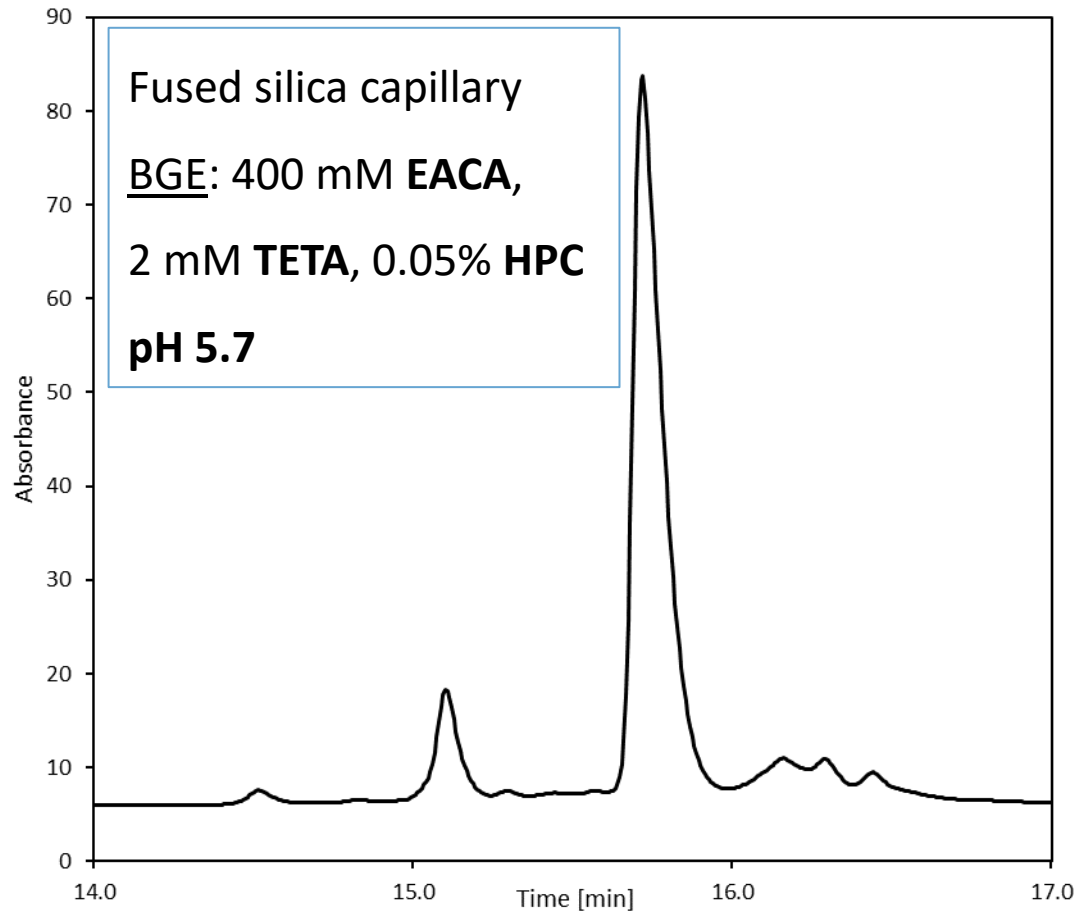
→ But: **Not** MS compatible ✗



→ MS compatible ✓

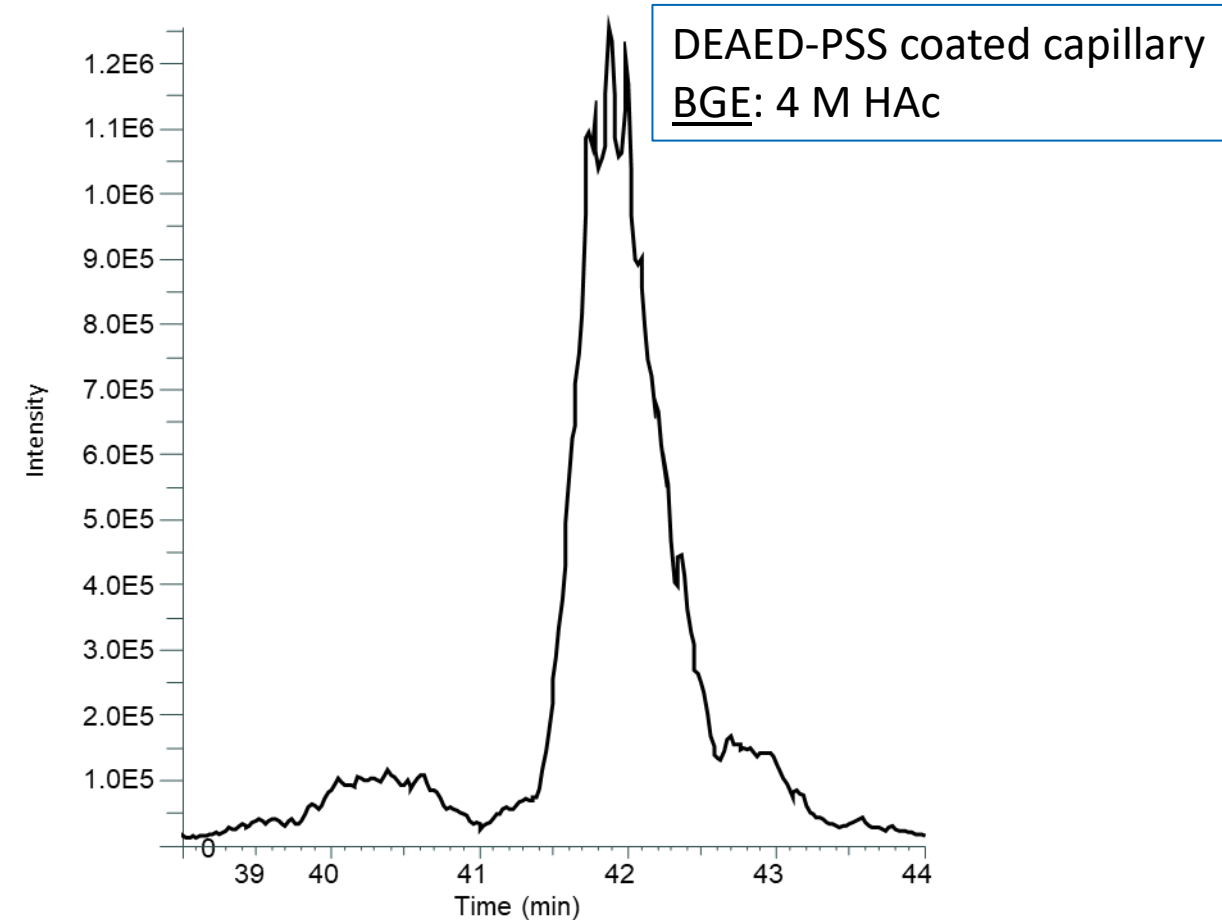
→ **No separation** of charge variants ✗

CE-UV of NISTmAb using different electrolytes



→ Efficient separation of charge variants ✓

→ But: **Not** MS compatible ✗

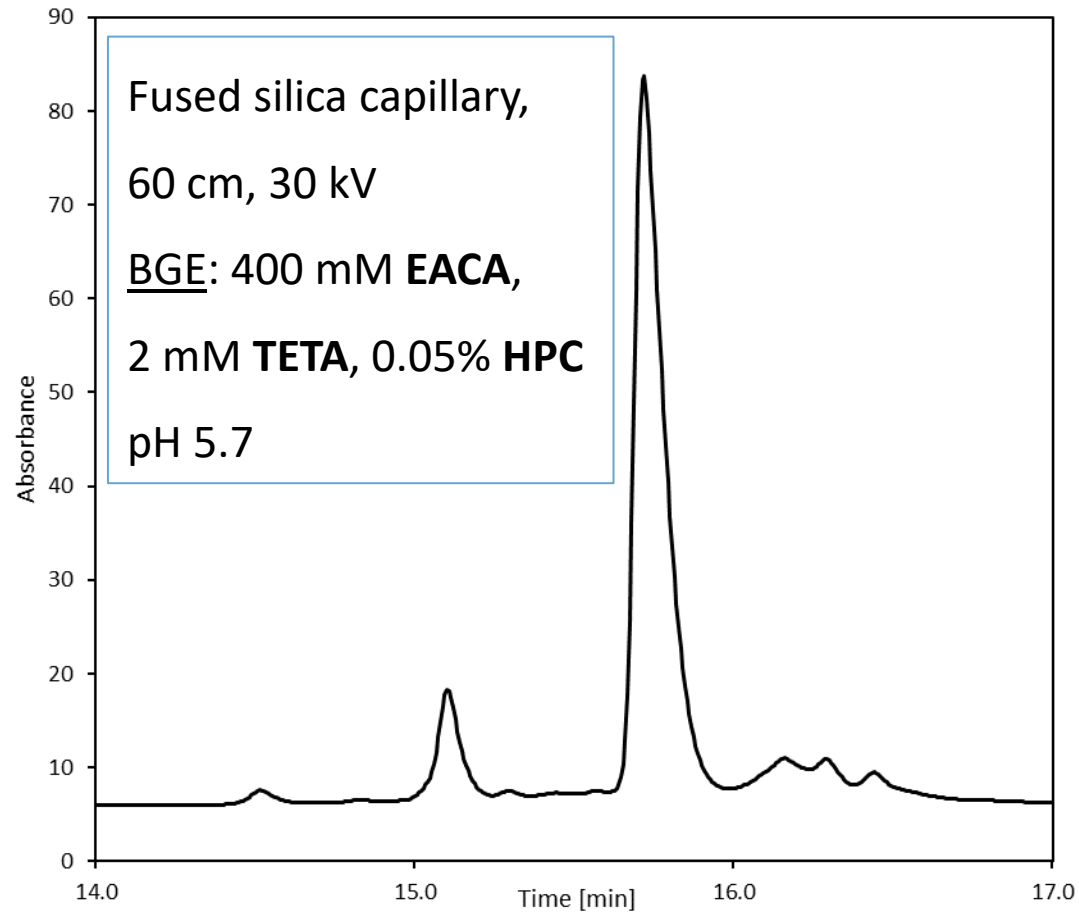


→ MS compatible ✓

→ **Slight separation** of charge variants

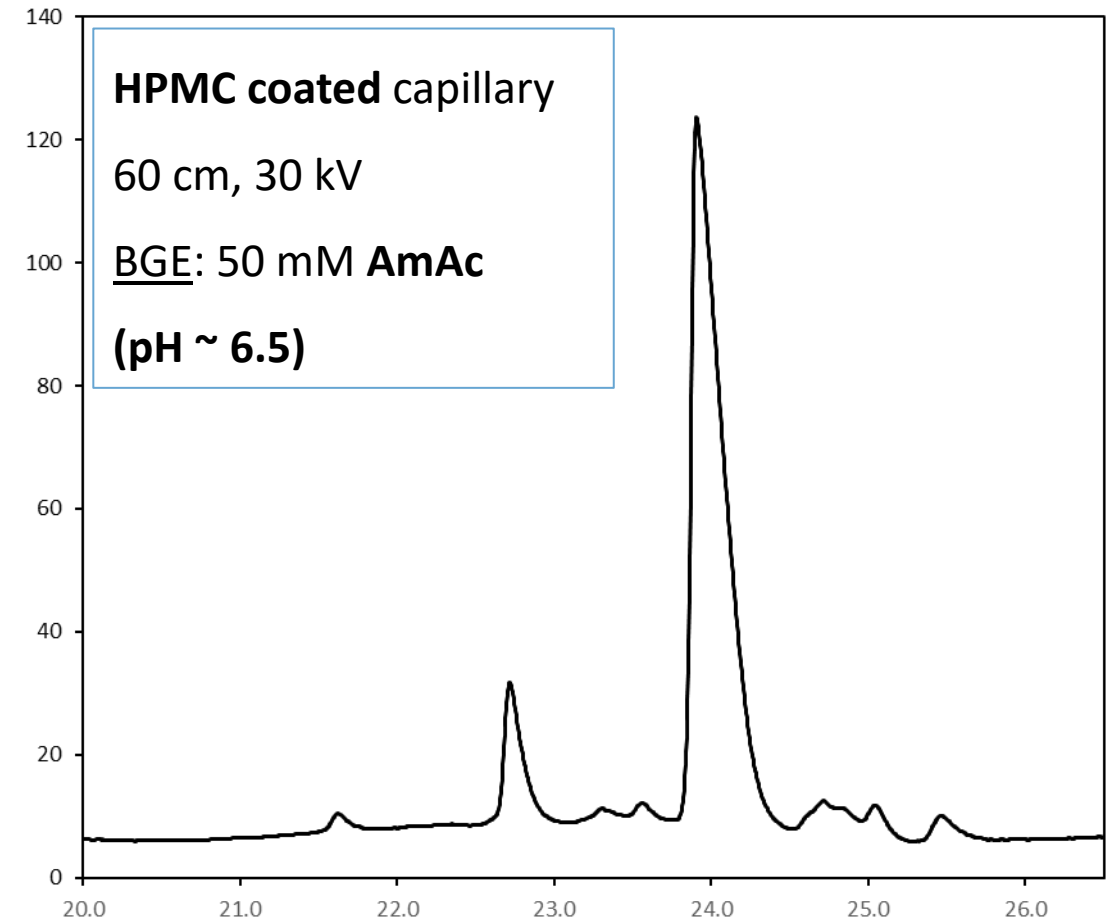
Höchsmann, A. et al., Charge variant analysis of monoclonal antibodies by CZE-MS using a successive multiple ionic-polymer layer coating based on diethylaminoethyl-dextran. *Electrophoresis* **2025**

CE-UV of NISTmAb using different electrolytes



→ Efficient separation of charge variants ✓

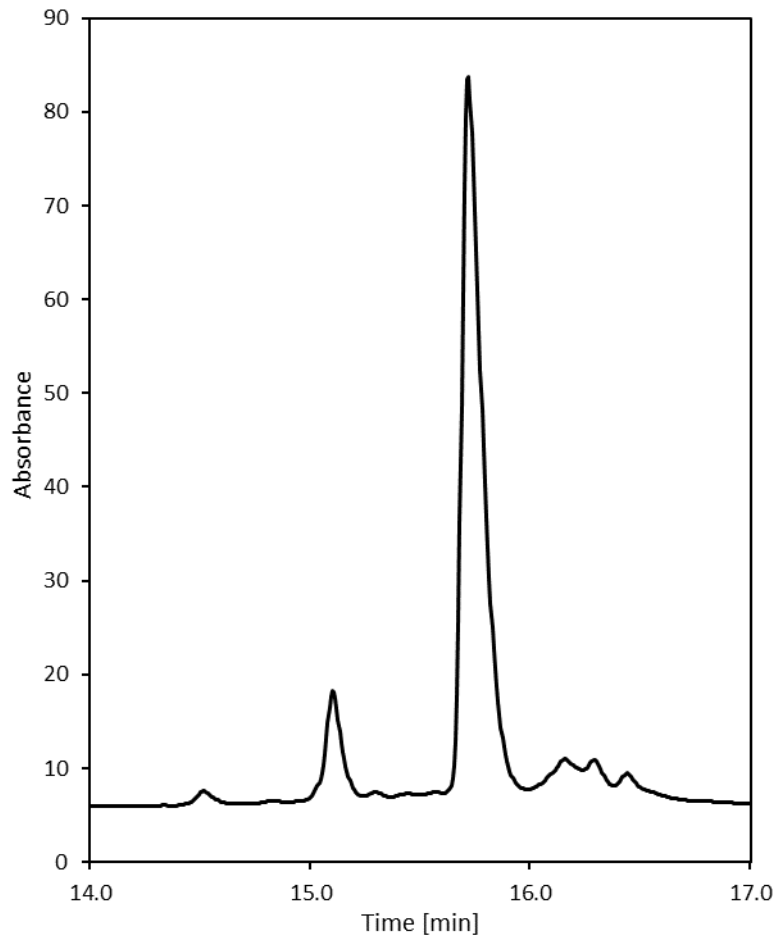
→ But: **Not** MS compatible ✗



→ Efficient separation of charge variants ✓

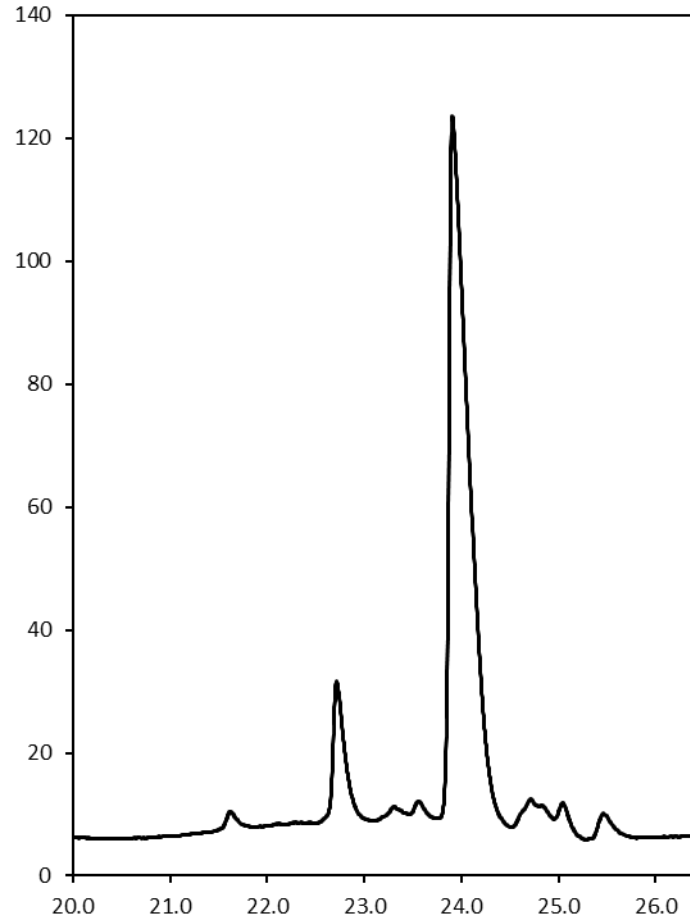
→ **MS compatible** ✓

CE-UV (EACA/TETA method)



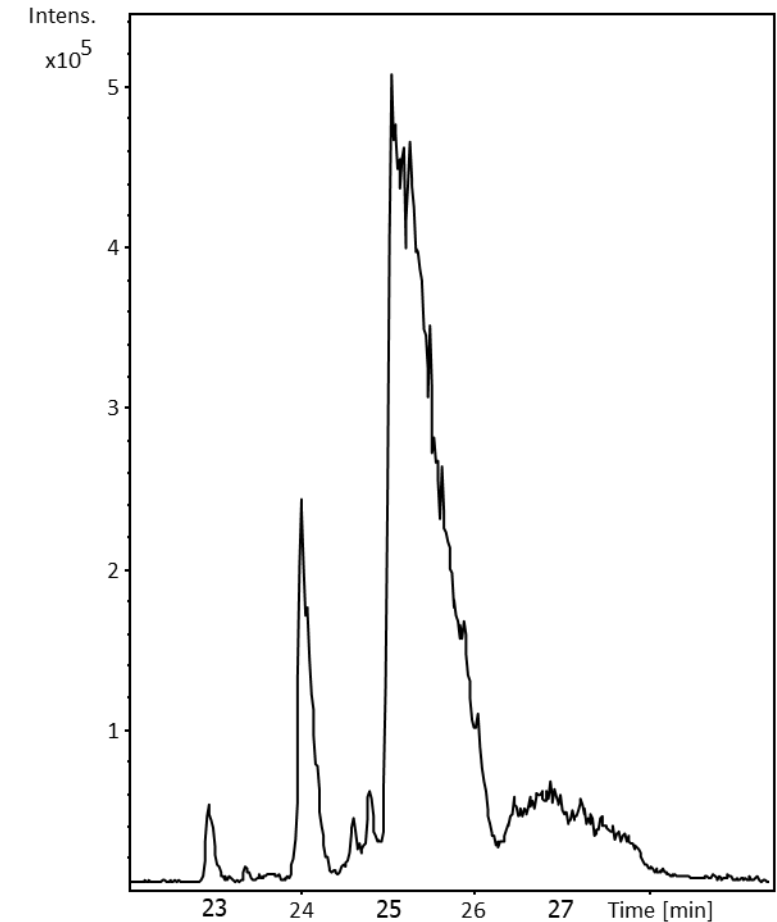
Fused silica capillary, 60 cm, 30 kV
BGE: 400 mM **EACA**, 2 mM **TETA**,
0.05% **HPC**; pH 5.7

CE-UV (MS compatible method)

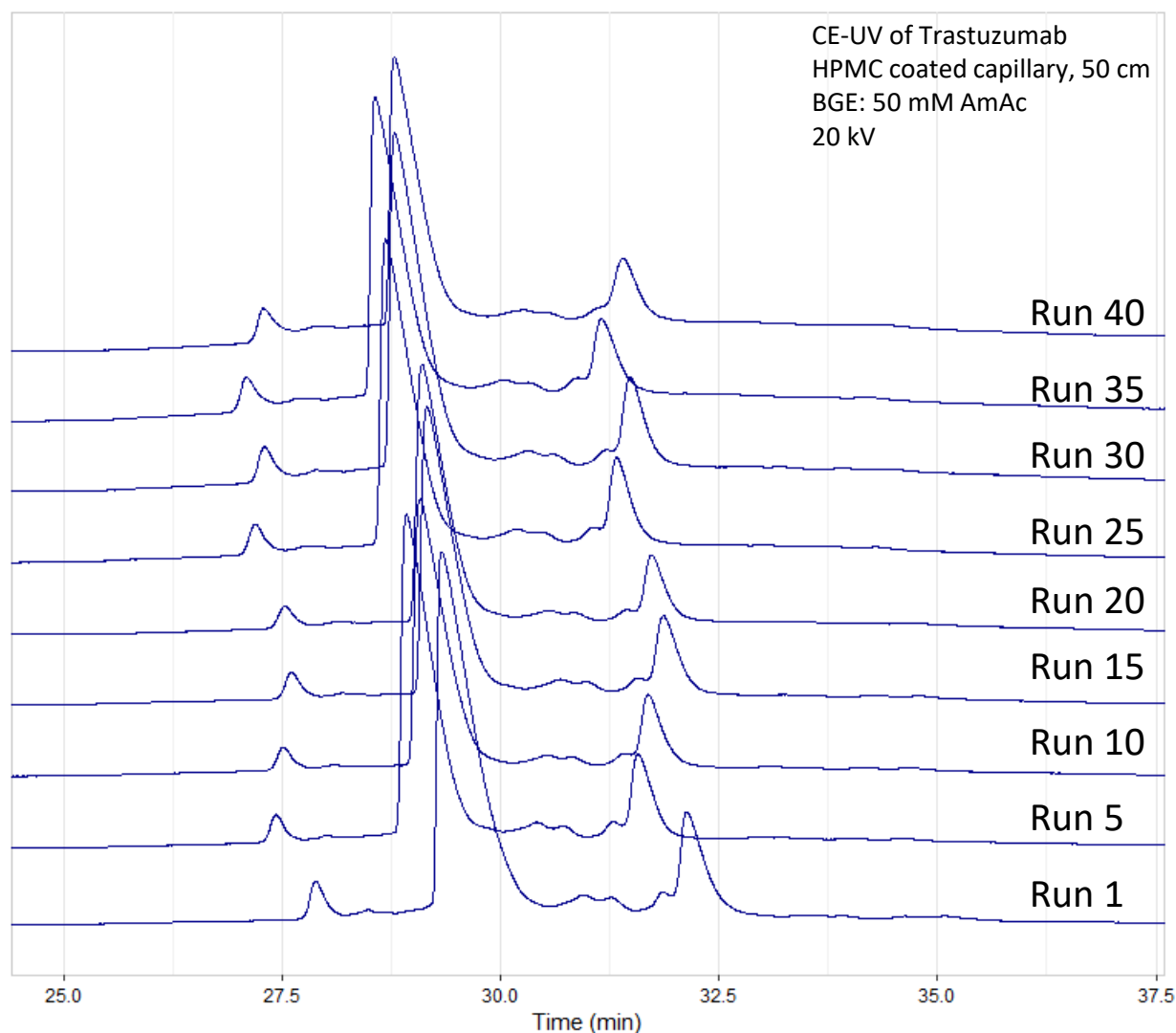


HPMC coated capillary, 60 cm, 30 kV
BGE: 50 mM **AmAc** (pH ~ 6.5)

CE-MS (MS compatible method)

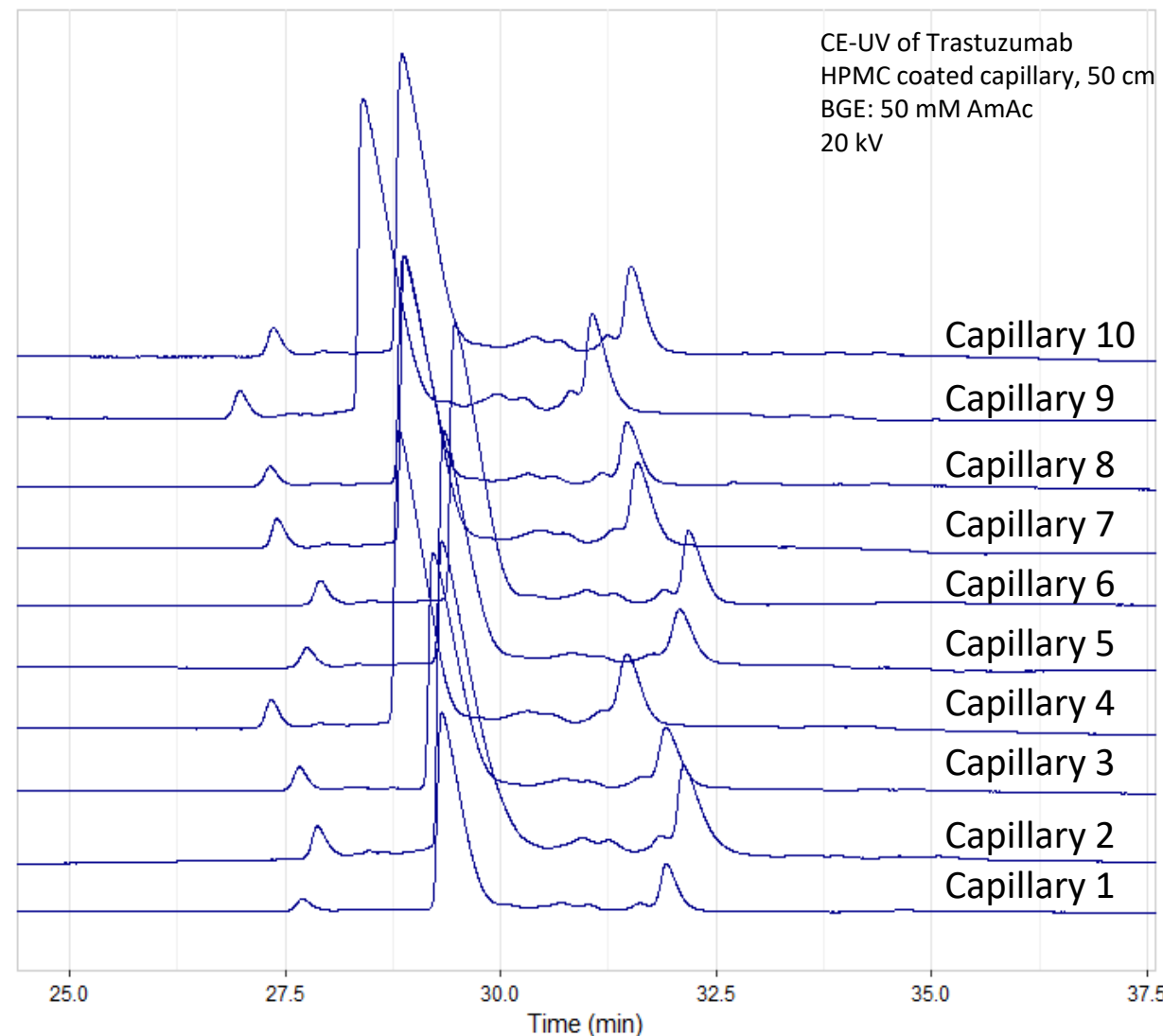


Repeatability (40 runs)



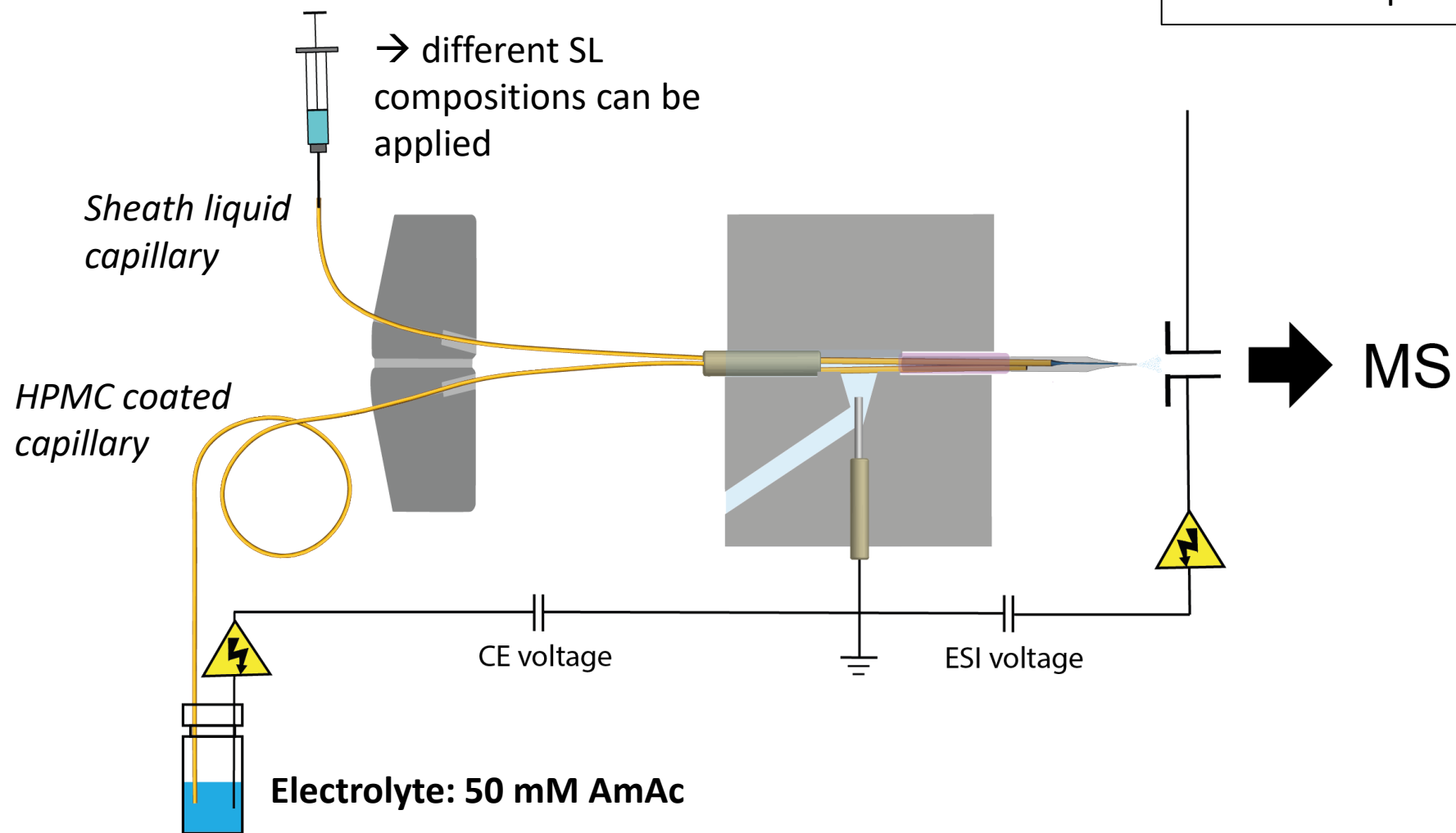
Relative standard deviation of migration times: 0.96%

Reproducibility (10 capillaries)

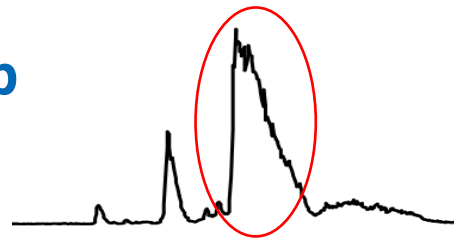


Relative standard deviation of migration times: 1.15%

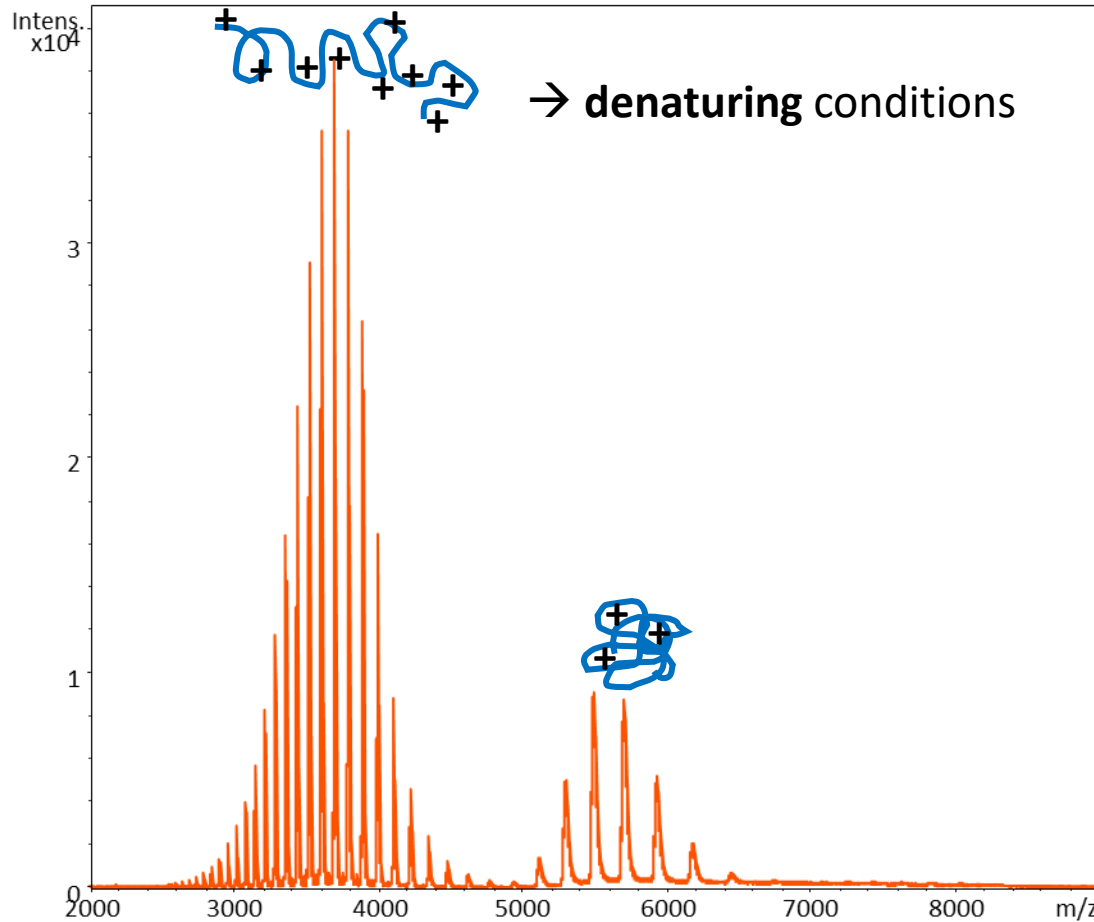
CE-MS using the nanoCEasy interface



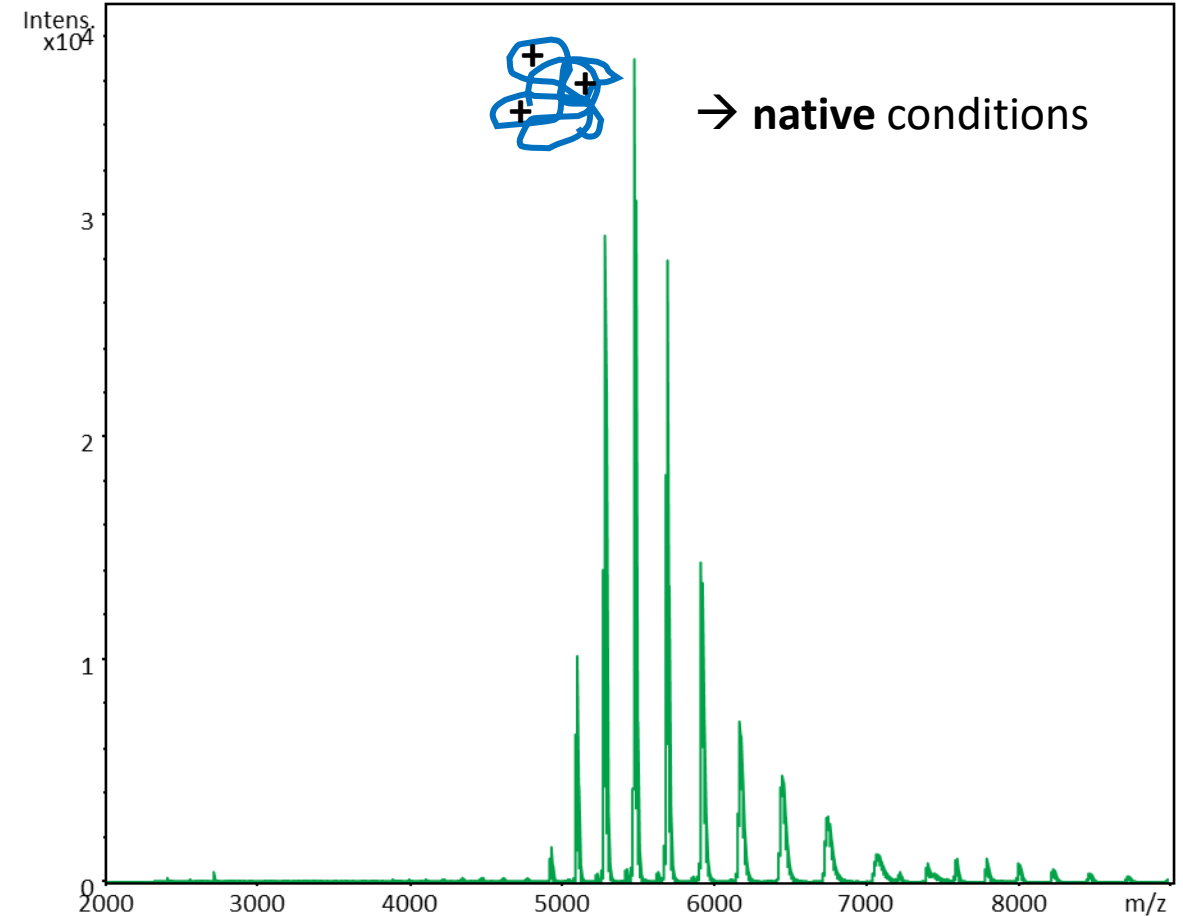
CE-MS of the main form of NISTmAb using different SL



IPA/H₂O (50:50), 5 % HAc

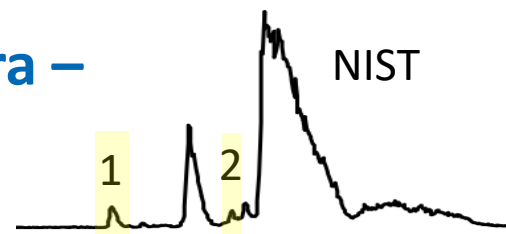


IPA/H₂O (30:70), 20 mM AmAc



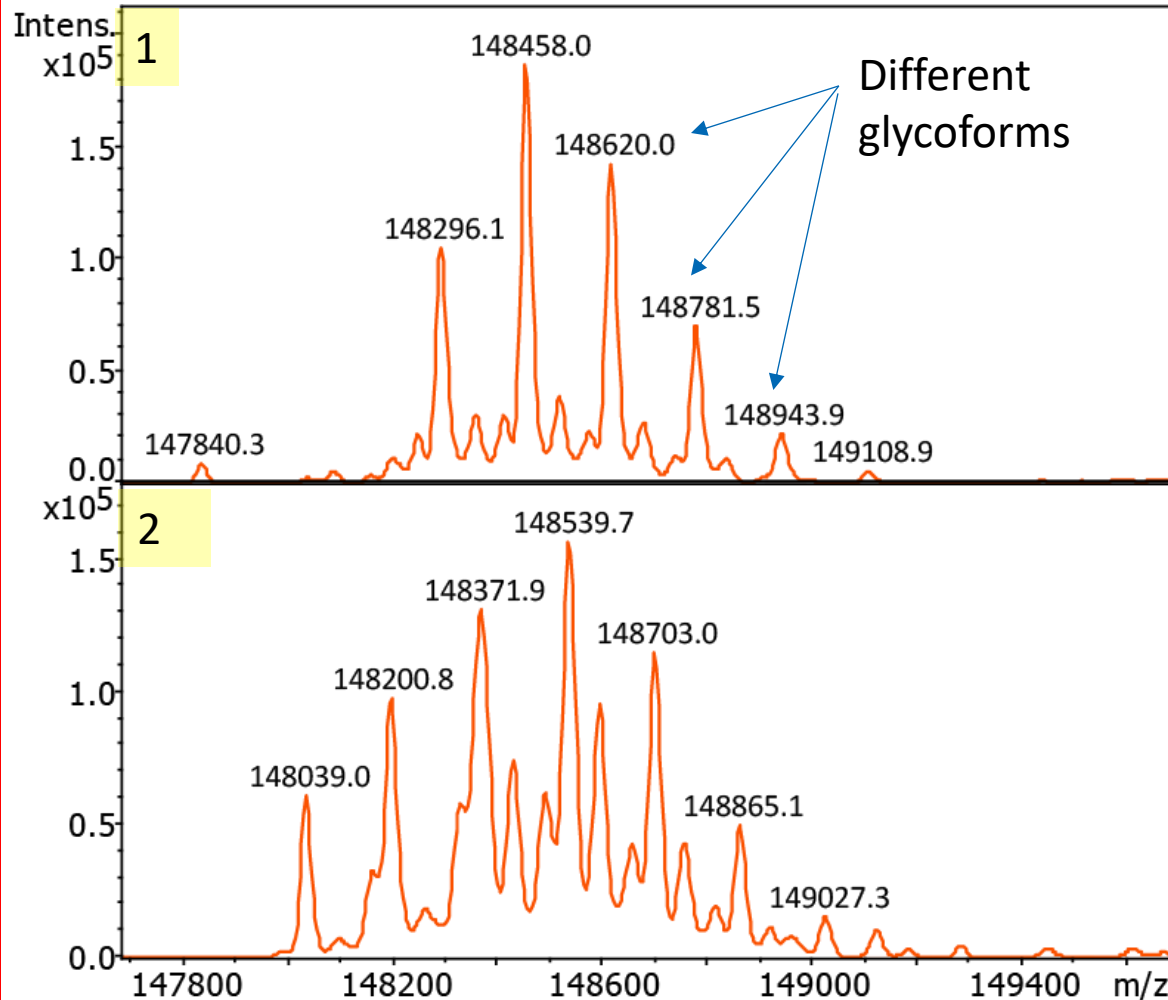
→ The nanoCEasy interface enables analysis of CE separated charge variants under **denaturing** and under **native nanoESI conditions**

Comparison deconvoluted mass spectra – denaturing vs. native ESI conditions



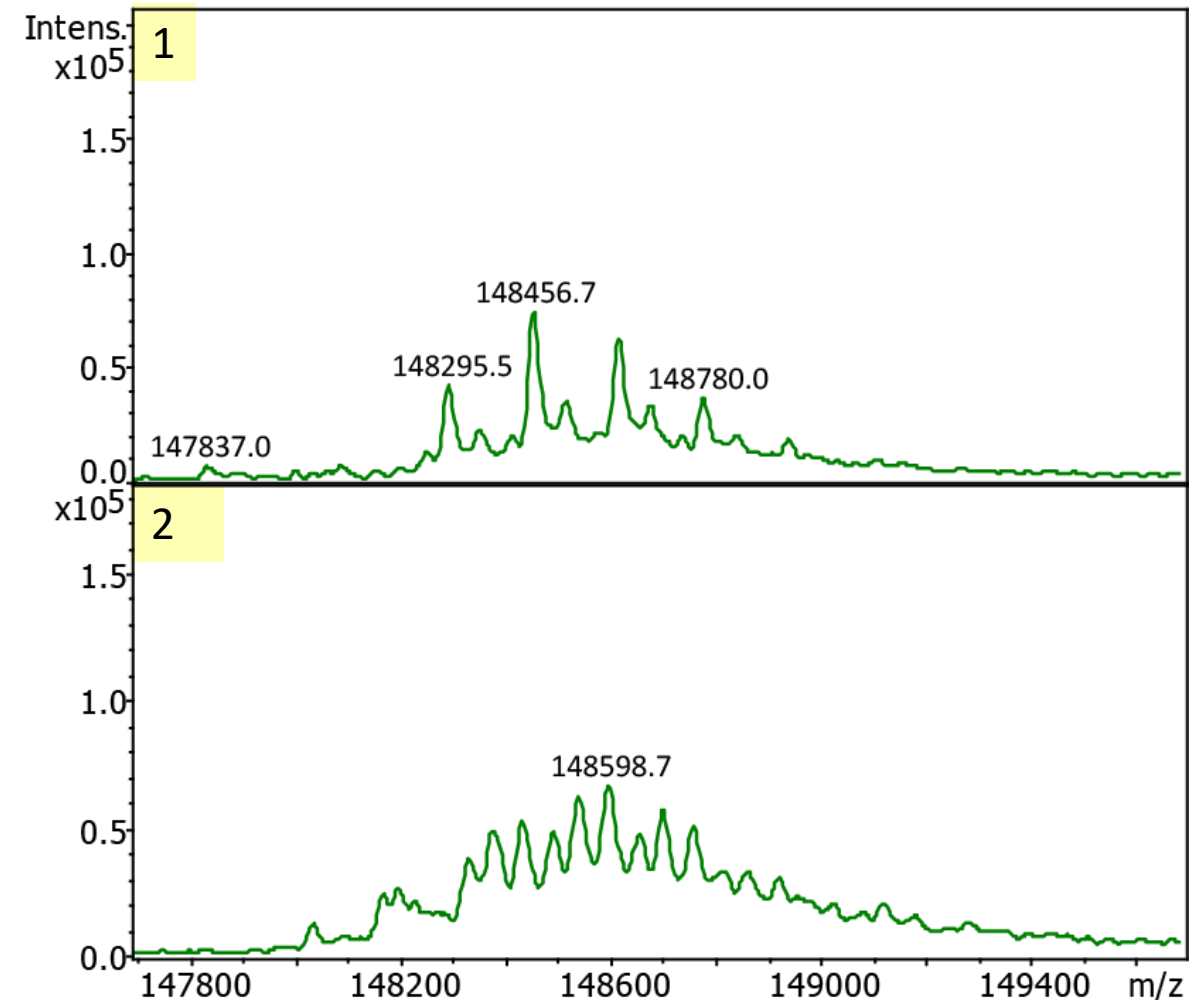
Denaturing

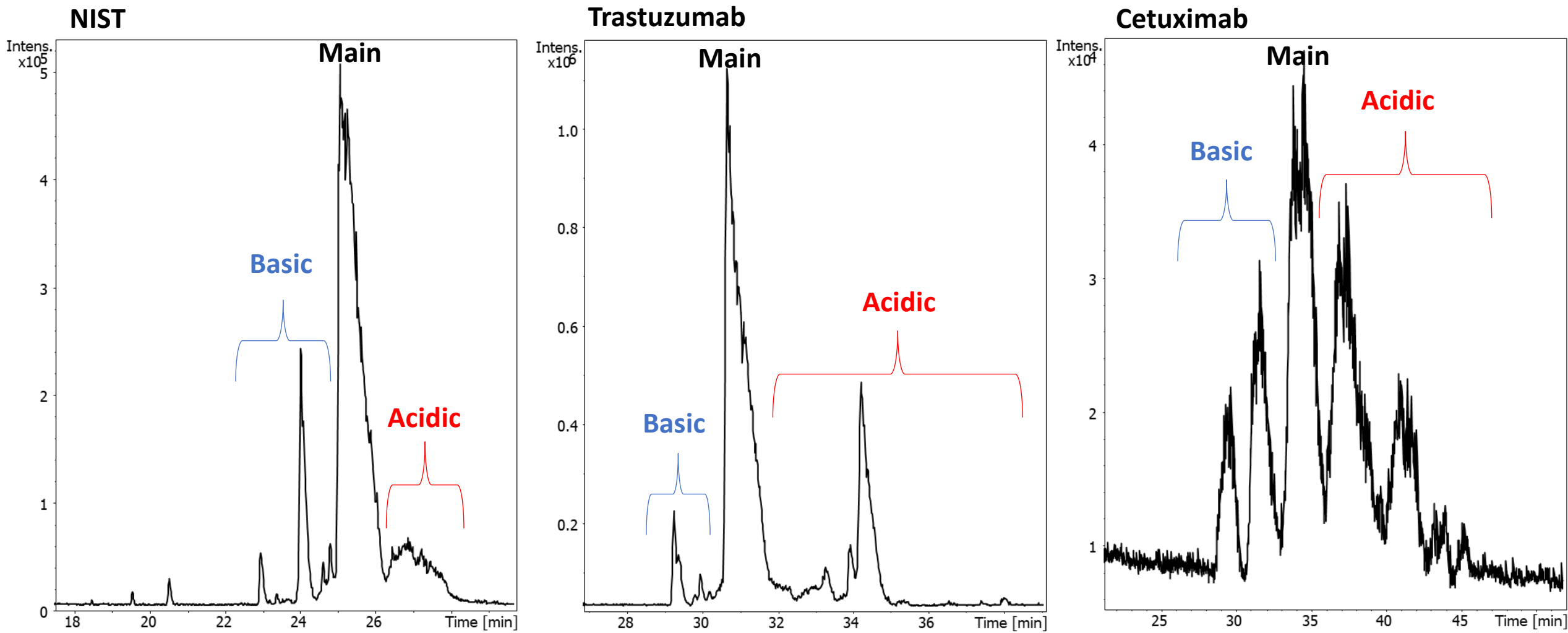
SL: : IPA/H₂O (50:50), 5% HAc



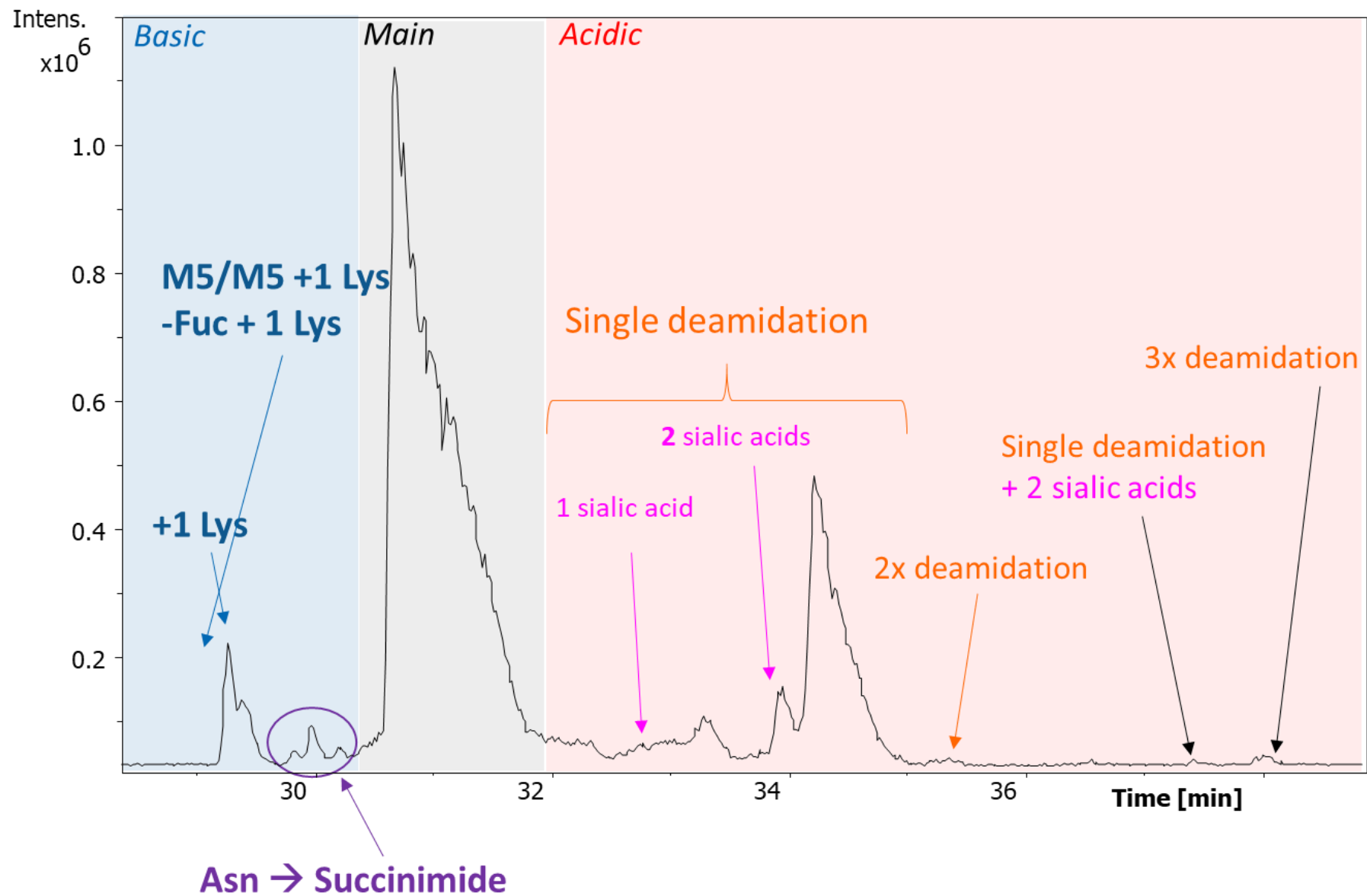
Native

SL: : IPA/H₂O (30:70), 20 mM AmAc

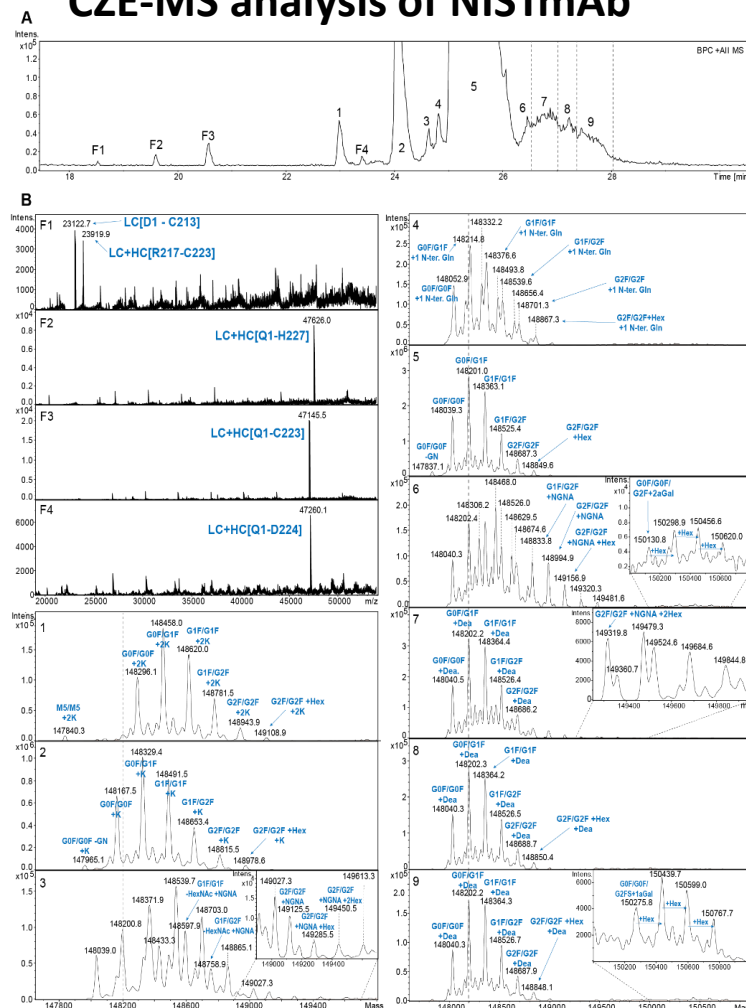




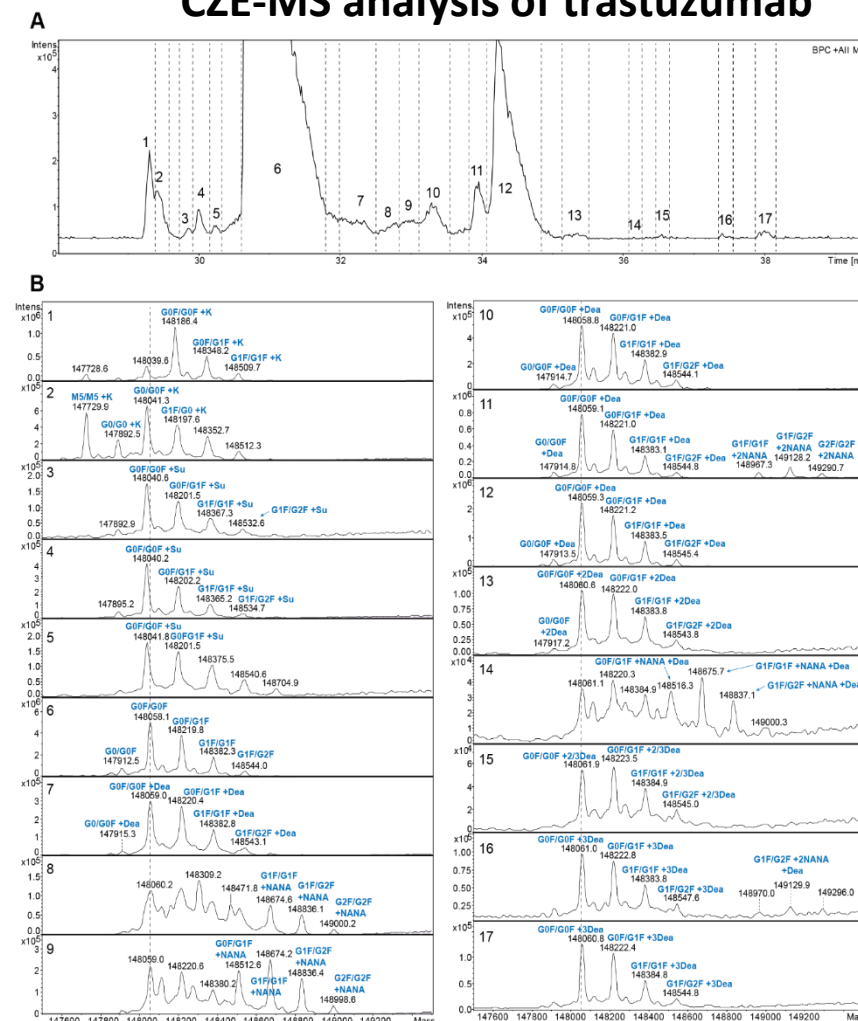
→ separation of basic, main and acidic variants



CZE-MS analysis of NISTmAb



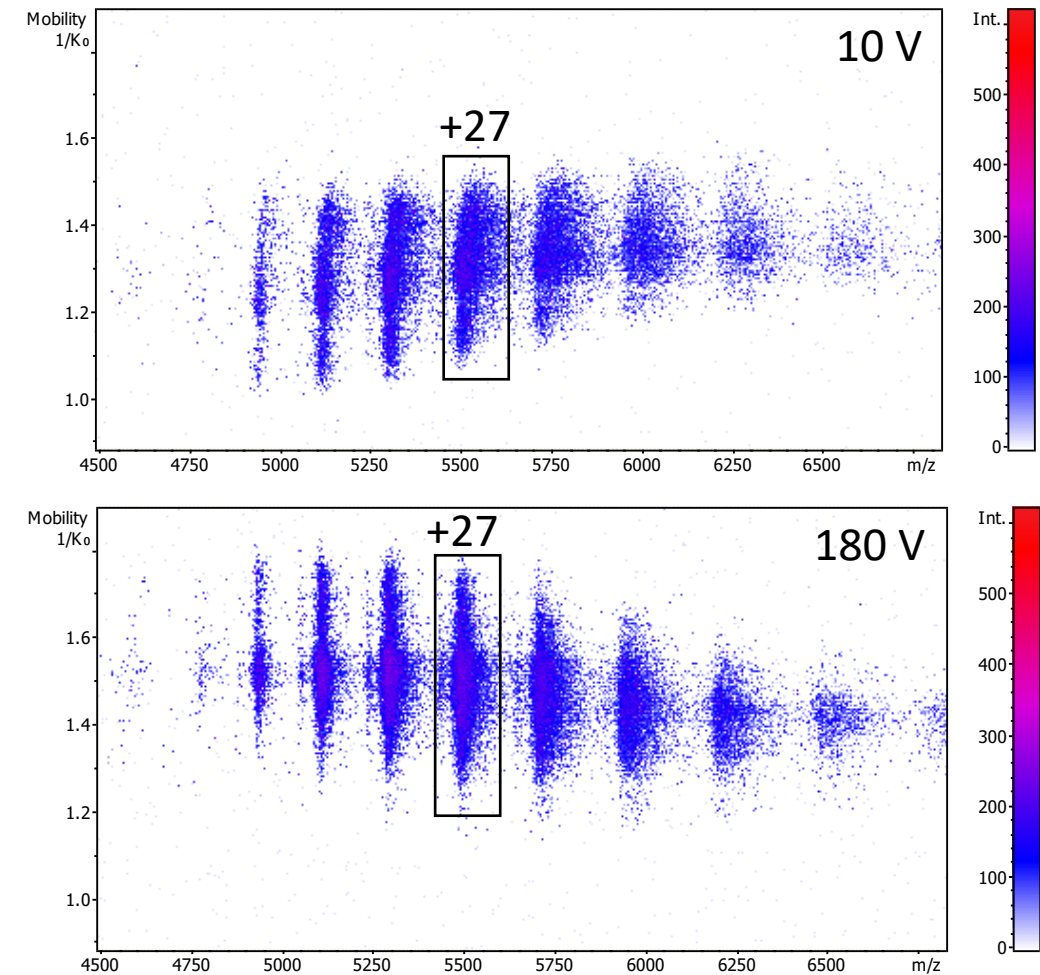
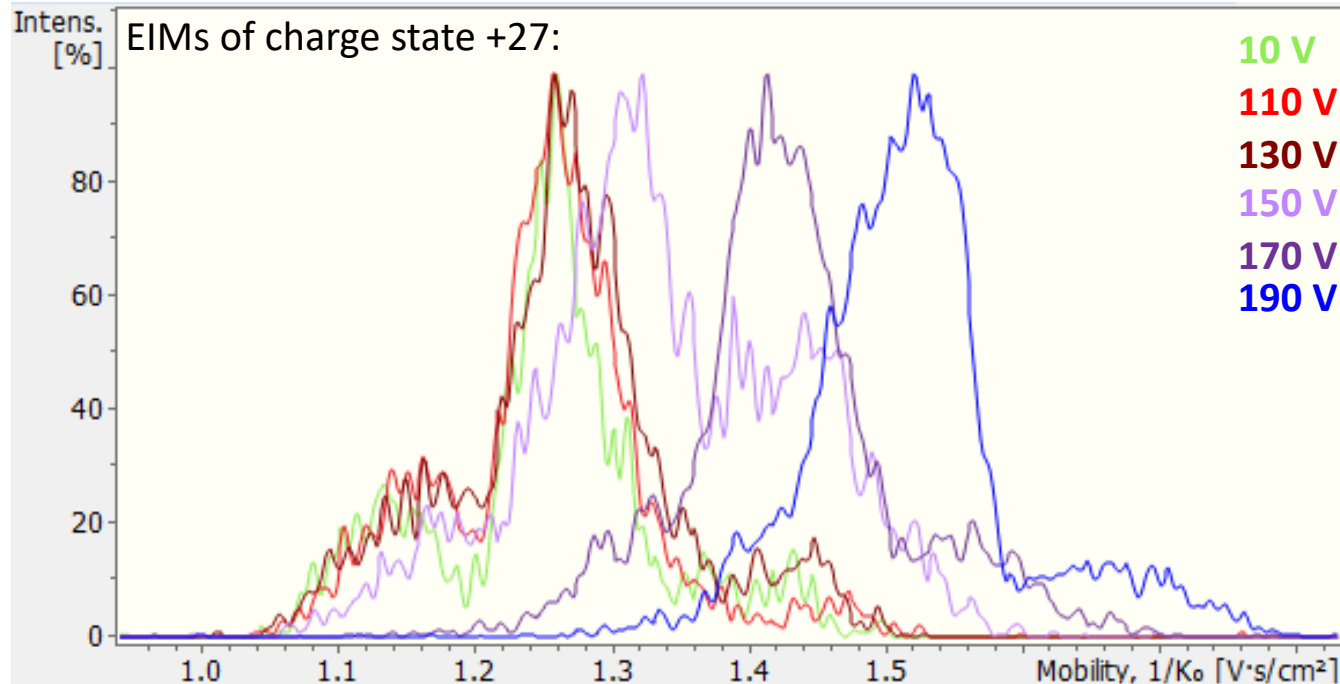
CZE-MS analysis of trastuzumab



→ Detection of **70** proteoforms for trastuzumab and **115** proteoforms for NISTmAb

Ion Mobility of native mAbs on modified timsTOF SCP

→ Collision Induced unfolding (CIU) of Trastuzumab:



Changes in structure of protein variants?

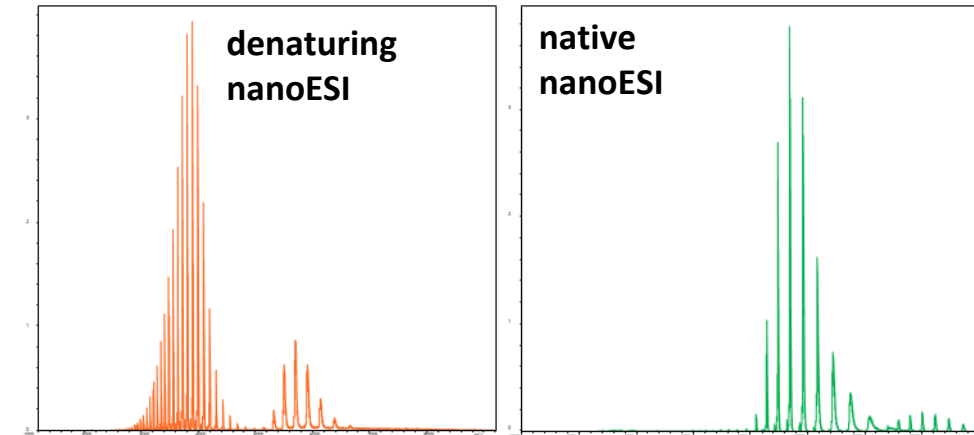
→ Differences in mobility

→ Differences in CIU

- Combination of native CE separation with ultrasensitive denaturing online MS analysis for charge heterogeneity characterization

→ Detection of **70 proteoforms** for trastuzumab and **115 proteoforms** for **NISTmAb**

- **Online MS analysis under denaturing or native ESI conditions**



Outlook:

- Native SL system enables **affinity CE with native MS**
 - Analysis of **antibody-antigen interactions** of **proteoforms**?
- Analysis of **changes in structure of protein variants** using ion-mobility MS?

Acknowledgement

Collaboration Partners:

abbvie

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Markus Hollmann



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Oliver Räther

Christian Albers



Laura
Stürzel

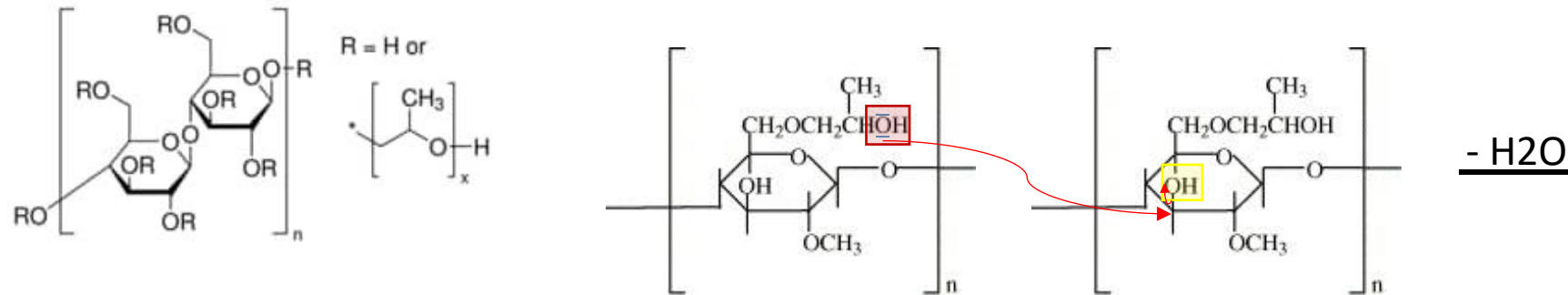
Jasmin
Schairer

Alisa
Höchsmann

Founding: German Federal Ministry for Education and Research (FKZ 13FH135KX0)



→ By heating, the **hydroxyl groups** on the substituted hydroxyalkyl chains can react with the **remaining hydroxyl groups** on the substituted cellulose chain to yield relatively pH stable cellulose ether bonds, as formed in the manufacture of the hydroxyalkyl-substituted celluloses. Shen et al. **suppose** that chemical **reactions** occur **between** the coated hydroxyalkyl-substituted **celluloses** and the **fused silica capillary** inner wall.



Comparison of CE currents for different SL:

