

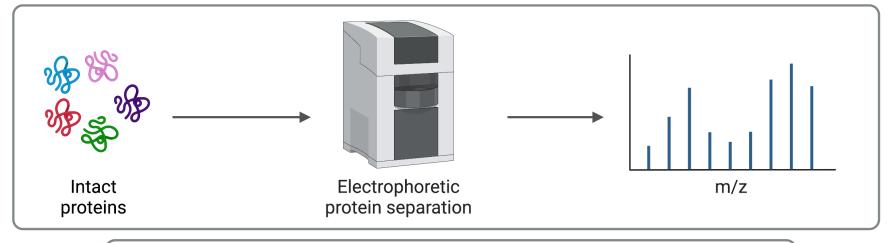
Desing of a capillary isoelectric focusing electrospray ionization-mass spectrometry interface for high resolution proteoform analysis

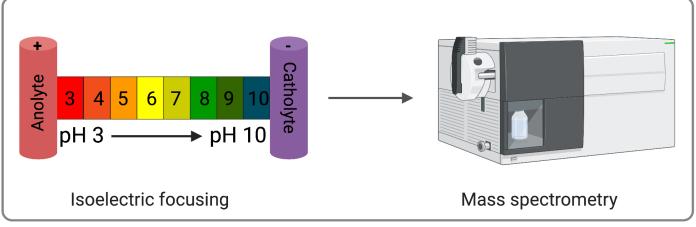
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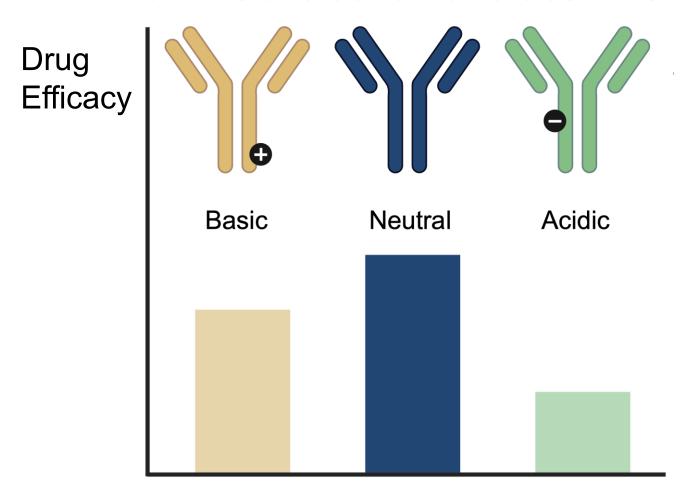


Using electrophoretic separations coupled to mass spectrometry



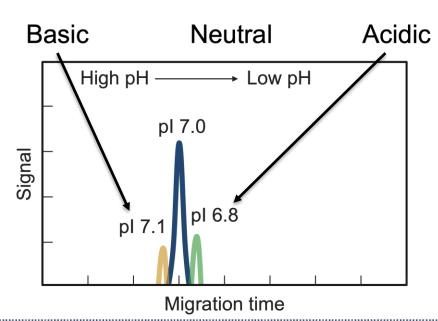


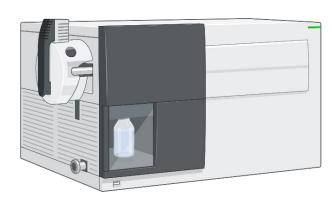
Protein charge variants alter function and are difficult to characterize



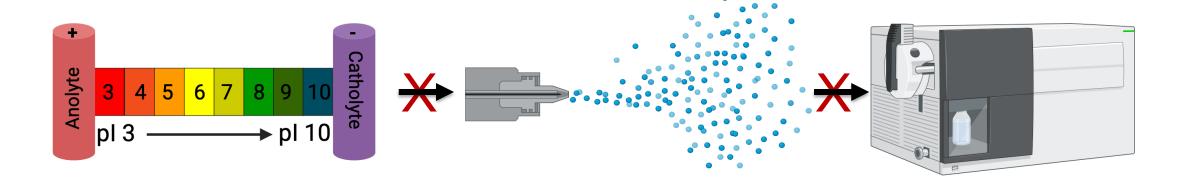
Protein charge variants can be separated with cIEF







cIEF coupled to mass spectrometry is a compromised process

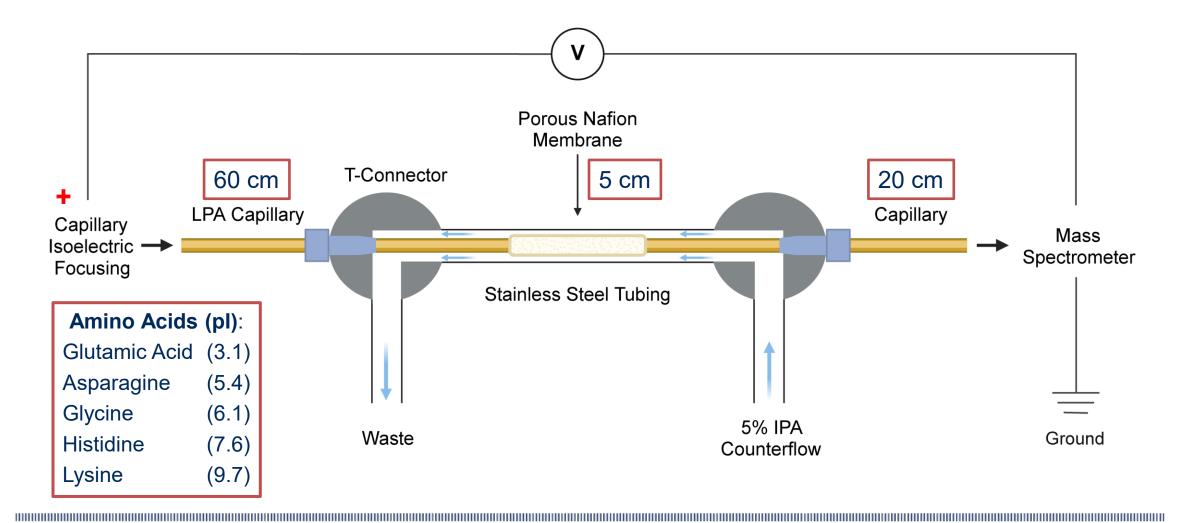


- 1 Capillary Isoelectric Focusing
- 2 Electrospray Ionization
- 3 Mass Spectrometry

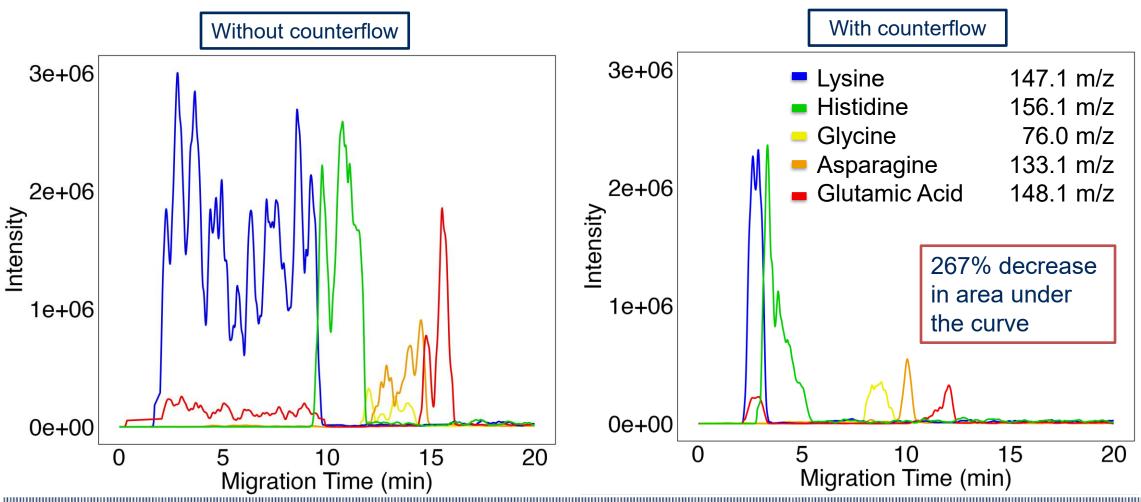
Issues with cIEF-MS:

- Ionization suppression
- Mass spectrometry contamination

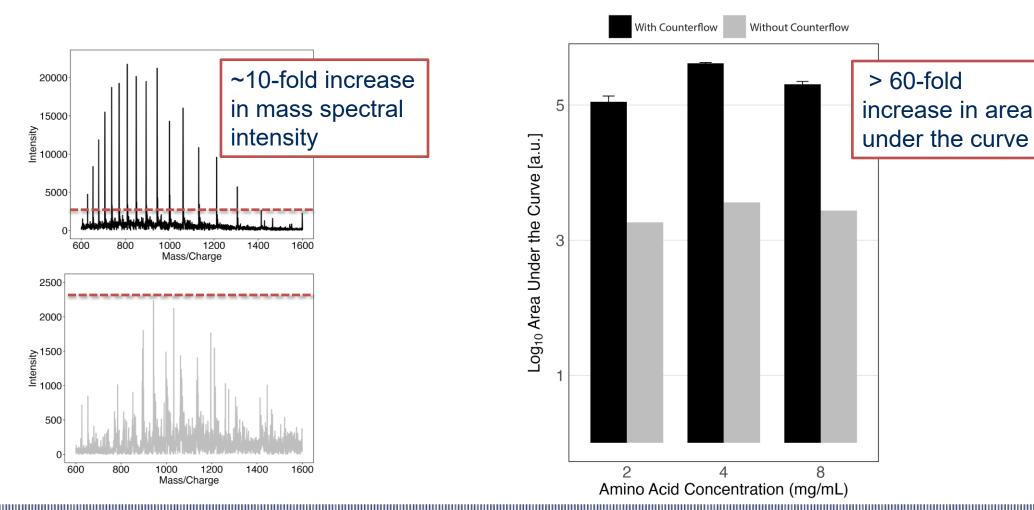
The interface reduces ampholyte concentration prior to MS



Extracted ion electropherograms of amino acid ampholytes signal with the counterflow decreases

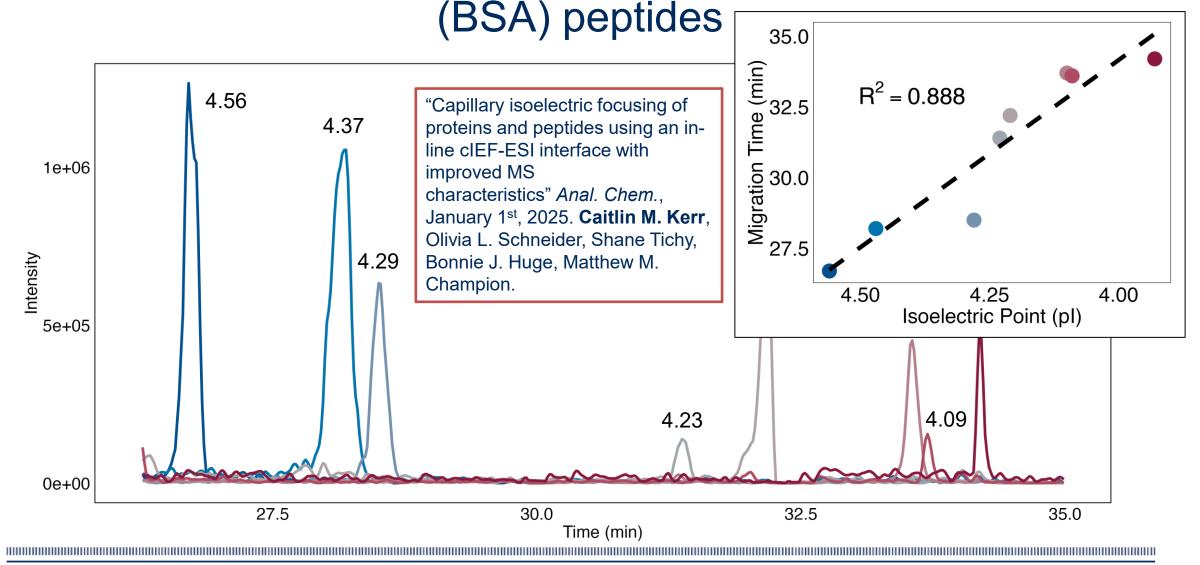


Interface increases the electropherogram area under the curve and mass spectra intensity of myoglobin



100 ng myoglobin injections

High resolution cIEF-Nafion-MS of bovine serum albumin



CIEF-Nation-MS conclusions

Conclusions

- The interface reduces the amino acid concentration prior to MS
- The interface decreases ionization suppression
- The amino acids can be optimized for high pl protein separations
- BSA peptides can be separated with high resolution
- The interface can be applied to clean up/desalt other analytes

Acknowledgements

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 - Bristol Myers Squibb
 - Dr. Tara Enda
- Some images created on Biorender and R Studio





<u>Group</u>: Dr. Taylor Lundgren, Dr. Daniel Hu, Hannah Marietta, **Dr. Bonnie Huge**, Dr. Sadie Schultz, **Sacheela Wanigasinghe**, Dr. Matthew Champion, Dr. Simon Weaver, and Christine DeRosa, **Olivia Schneider**, Dr. Bruce Mousseau, and Elizabeth Chang. <u>Not Pictured</u>: Angela Good, Emily Bravo, Anna McCartan

Questions?