

Unraveling proteoform complexity by native liquid chromatography-mass spectrometry

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Stellingen behorend bij het proefschrift getiteld

Unraveling proteoform complexity by native liquid chromatography-mass spectrometry

- While biotherapeutics often have predictable PTM profiles, industrial enzymes keep surprising with the presence of rare modifications. this thesis
- Not every question requires the use of state-of-the-art technologies. Sometimes a simple SDS-PAGE gel provides you with the answers you need. this thesis
- 3. To ensure protein functionality at the end of the workflow, the effect of each step should be examined, from sample preparation to fraction collection.

 this thesis
- Maximizing separation power does not guarantee elution of a single protein species, potentially hampering the (confident) determination of structurefunction relationships.
- 5. There is more to hyphenation than connecting two parts.
- 6. BAC shows great potential to assess glycation. Nevertheless, deeper insights into the retention mechanism of proteins are needed.
- 7. There is no universal method for achieving complete protein characterization, but rather a combination of multiple analytical techniques is required.
- 8. Synonyms of boring: dull, tiresome, monotonous, manual fraction collection.
- 9. Data visualization is a form of art.
- 10. Starting with a clear project description leads to more efficient research.
- 11. If PhD research is a party, you have to make sure to decorate it festively.