

Boosting mass spectrometry-based analytics for biopharma Gstöttner, C.J.

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Stellingen behorende bij het proefschrift

Boosting mass spectrometry-based analytics for biopharma

- 1. Intact and top-down approaches decrease the risk of introducing unintended modifications and provide a more accurate picture of the proteoform variability. (this thesis)
- 2. Biotechnologically produced proteins and their natural forms can differ enormously with respect to their proteoforms and, in consequence, to their functionality. (this thesis)
- 3. Multidimensional liquid chromatography has the potential to bridge the gap between manual and robotic sample preparation and analysis. (this thesis)
- 4. Affinity capillary electrophoresis hyphenated with mass spectrometry is a highly flexible tool, able to study affinity in a proteoform-specific manner, provides very detailed (higher-order) structural information and requires minute amounts of antibody and receptor. (this thesis)
- 5. Publishing in a suitable lower impact factor journal may help to better disseminate the results to the target audience than publishing in a non-suitable higher impact factor one.
- 6. While most conventional cancer immunotherapies allow only an extension of patients' lifetime, new antibody formats or therapies could hopefully allow to cure them.
- 7. If one or two dimensions are not enough, we should think in more dimensions.
- 8. Is capillary electrophoresis better than liquid chromatography or vice versa? I would say none is better, but they are complementary to each other.
- 9. Actively working on any machine/instrument increases the risk of damage.
- 10. Similarly, cleaning an instrument does not guarantee a better performance afterwards.
- 11. Collaboration between different groups can result in an excellent outcome, but that requires someone who keeps track of what everyone does.
- 12. Science is like hiking; every small step brings you closer to the tip from where you can enjoy the big picture and be proud of all the effort you made.