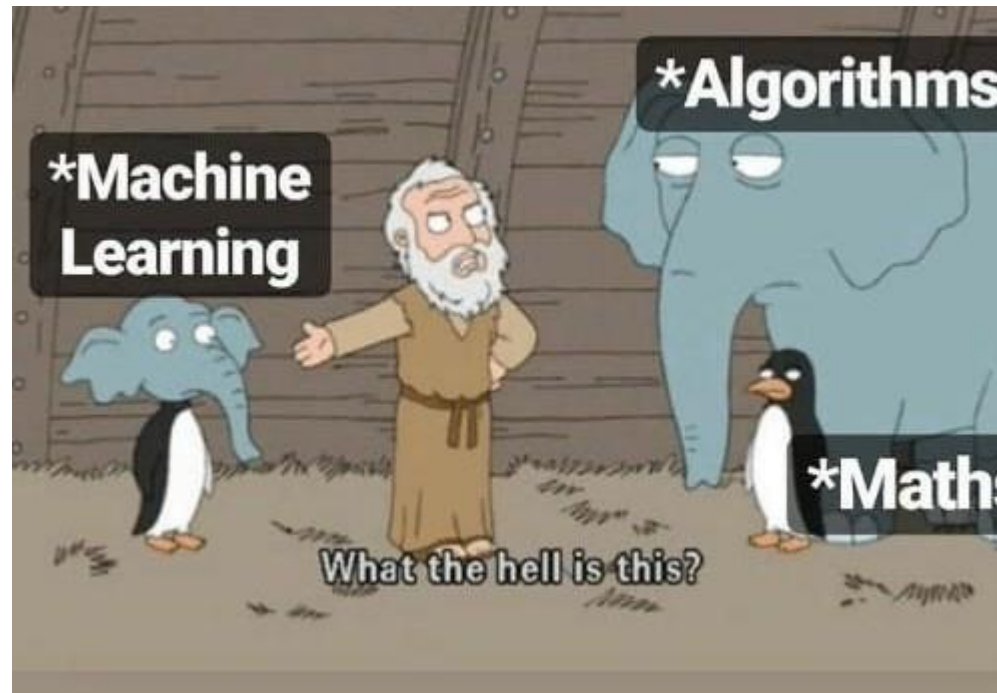


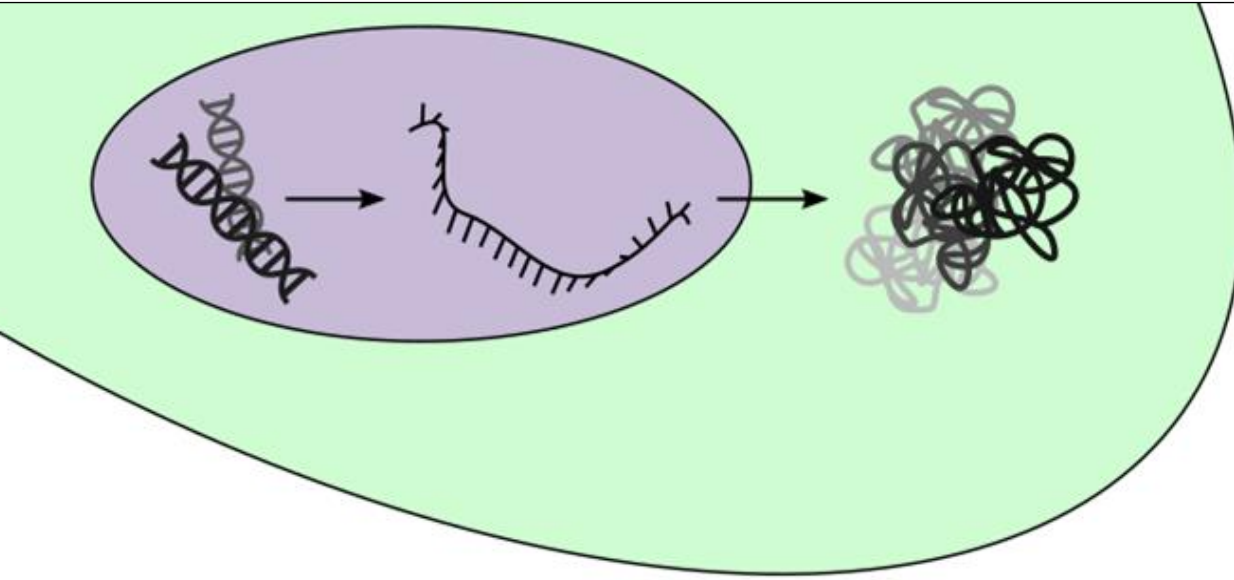
What can we learn about our health using machine learning?

Anna Sophie Welter
Proteome Dynamics
MDC, Berlin

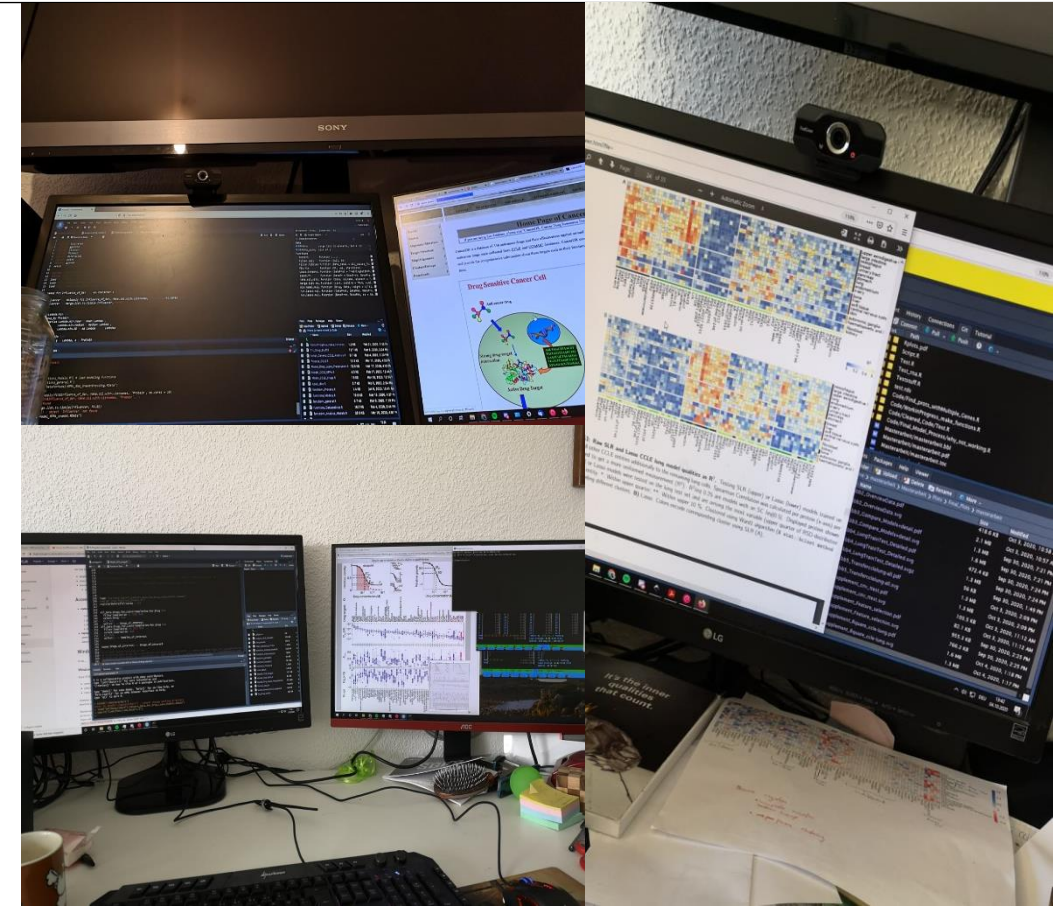
annasophie.welter@mdc-berlin.de



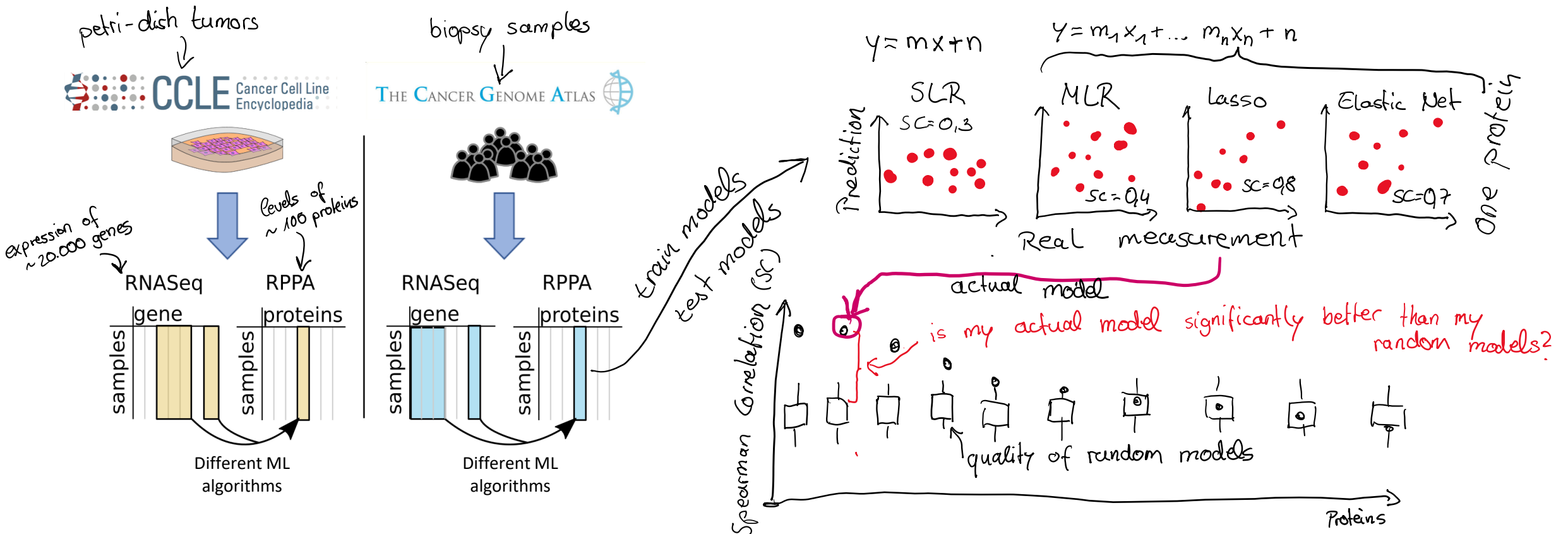
Master project – Predicting protein levels in cancer



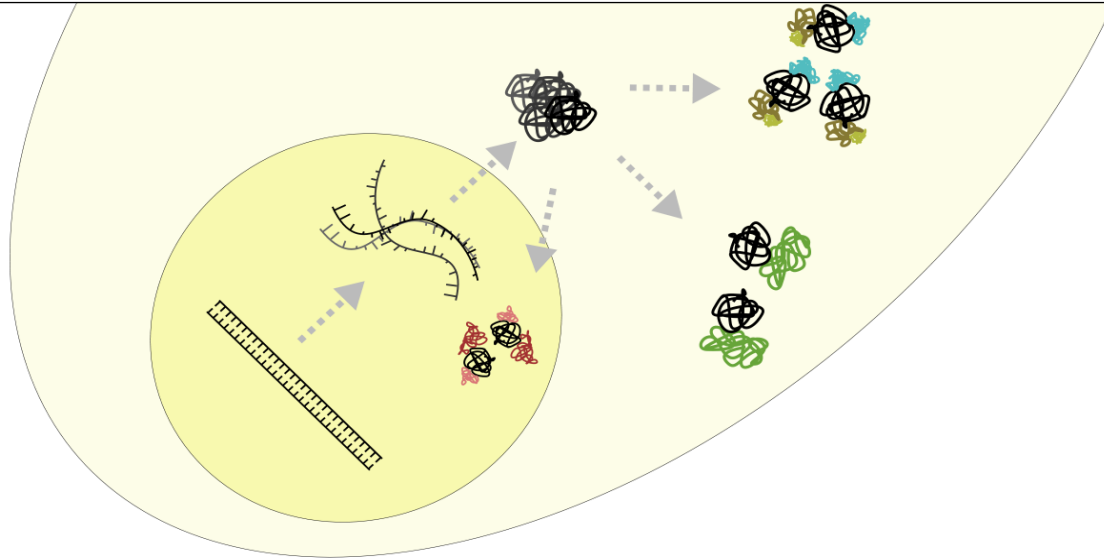
- Drugs act on (phospho)protein levels which are not measured
- Poor correlation between expression of (phospho)protein and encoding mRNA¹



Master project – Predicting protein levels in cancer



PhD project – Towards high-throughput analysis of protein-protein-interactions



- Protein-Protein-Interactions regulate cellular signaling and are aberrant in many cancers
- Understanding change in PPI might lead to discovery of new drug targets^{1,2}
- Implementing a screening in clinical settings might allow better treatments
- Understanding disease needs high amounts of data of different levels so we aim to implement mass spectrometry (MS)

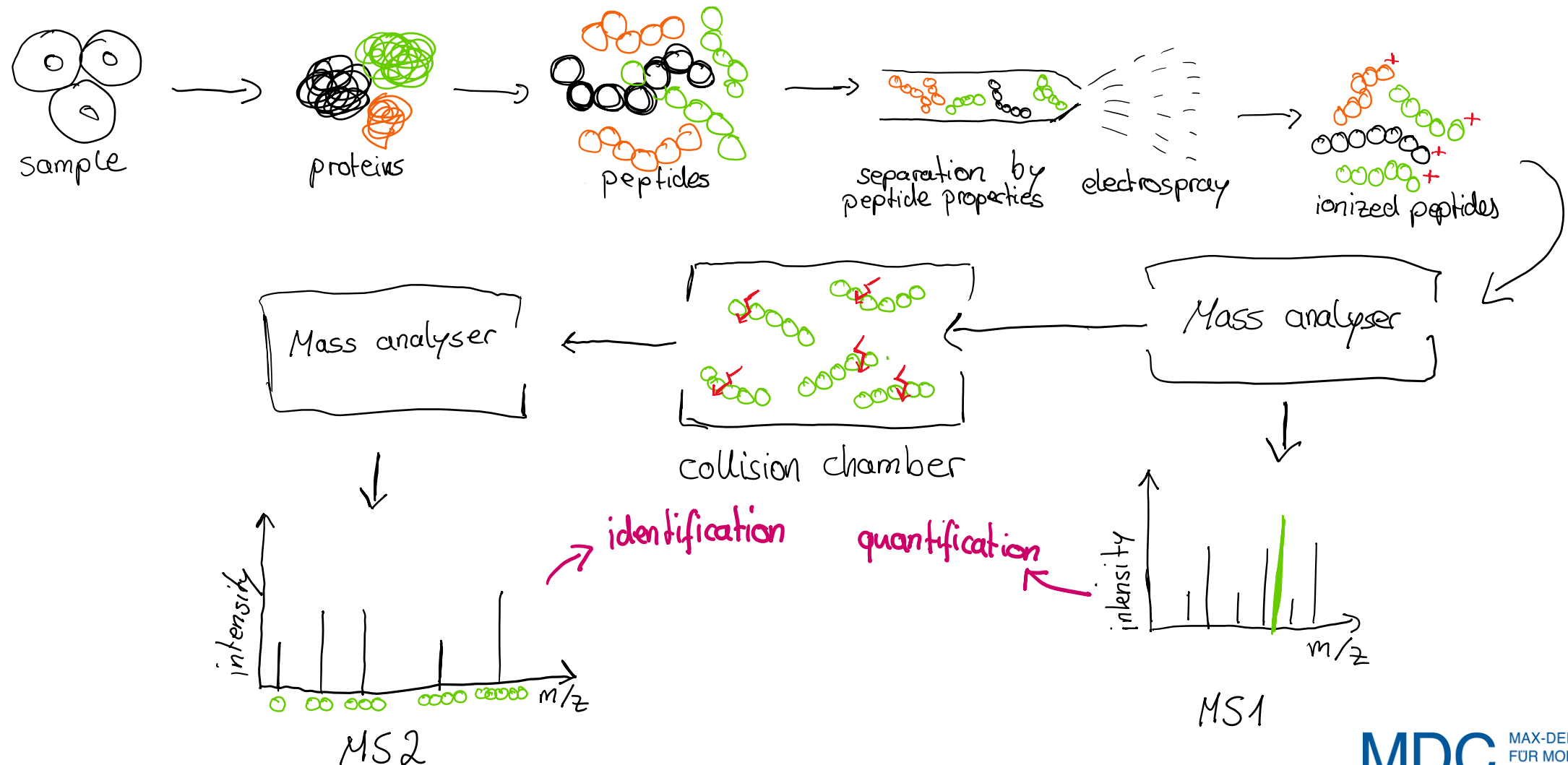
German consortium aiming to implement MS in clinics:
MSCoreSys - Massenspektrometrie in der Systemmedizin
<https://www.mscoresys.de/>



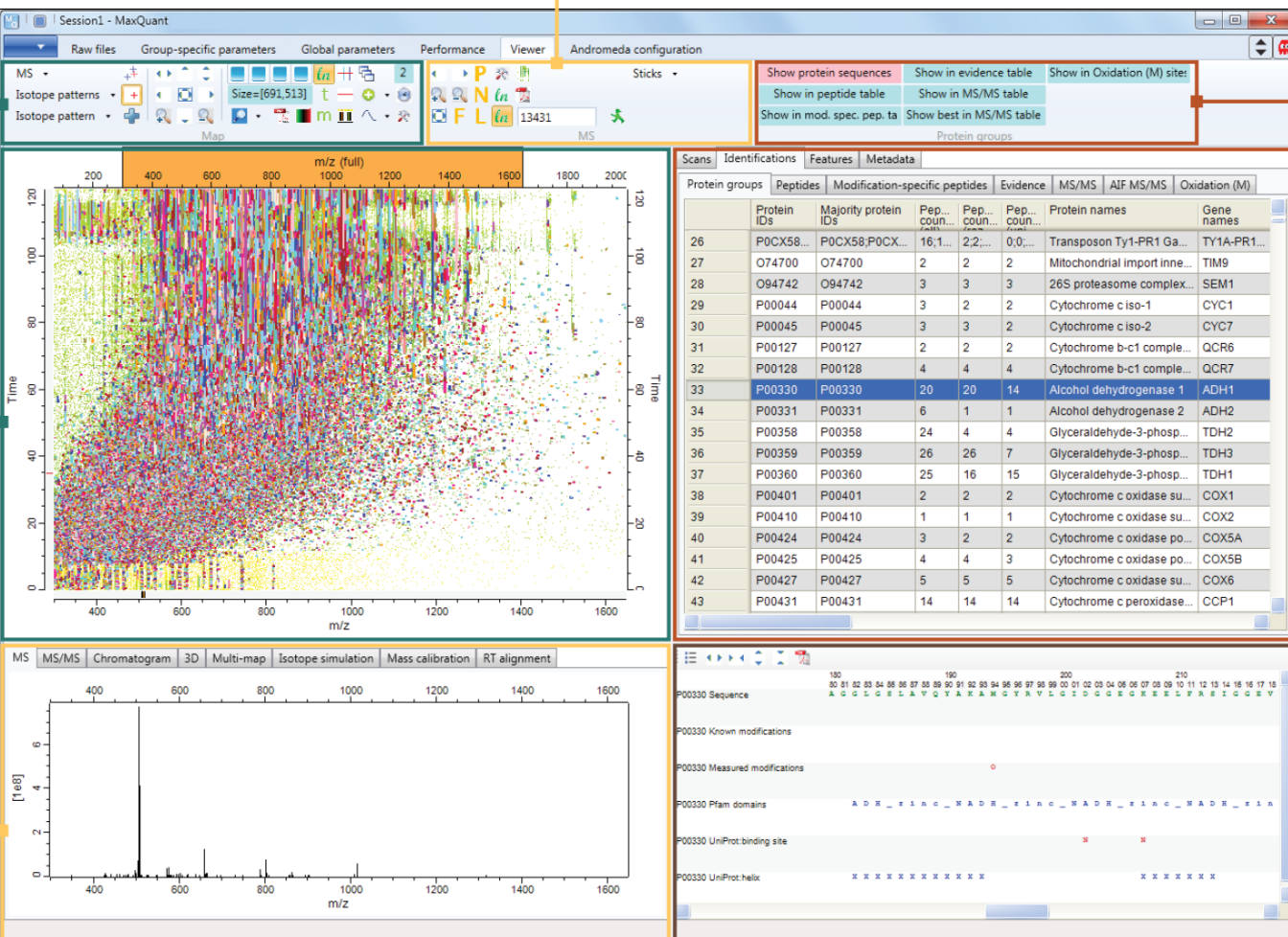
¹Lu et al. (2020): Recent advances in the development of protein-protein interactions modulators: mechanisms and clinical trials.

²Socinski et al. (2018): Atezolizumab for First-Line Treatment of Metastatic Nonsquamous NSCLC.

Using Mass Spectrometry to analyse proteins



Using Bioinformatics to deal with large amounts of data



- Powerful tools to identify the ~ 20.000 peptides and deduct the proteins they build
- Even more effort to quantify the proteins (peak intensity isn't directly correlated to amount of protein)
- Use statistics and ML to analyze and integrate the data

Take Home Messages



- You don't need to know what you want to do after finishing school.
- Use every opportunity you have to check out different fields!
- It's never too late to learn programming.