

# MS in the Faculty of Pharmacy - (single cell) proteomics and metabolomics

Jaakko Teppo, PhD (Pharm.)

University researcher

# Mass spectrometry and metabolomics research group

## People

- Jaakko Teppo ([jaakko.teppo@helsinki.fi](mailto:jaakko.teppo@helsinki.fi))
- Juho Heininen
- Essi Heinämäki
- Juha-Pekka Hieta
- Tapio Kotiaho
- Tom Sillanpää

## Laboratory



# Instruments

- Waters Xevo QTOF
- Waters TQ-S
- Thermo LTQ Orbitrap XL
- Bruker MicroTOF
- Agilent 6330 ion trap
- Waters Micromass Q-TOF Premier
- UPLC (3), capillary LC (2), GC (1)
  
- Thermo Orbitrap Fusion, UltiMate 3000 RSLC, EASY-nLC 1200
  - In Kumpula, co-owned by us and the Department of Chemistry

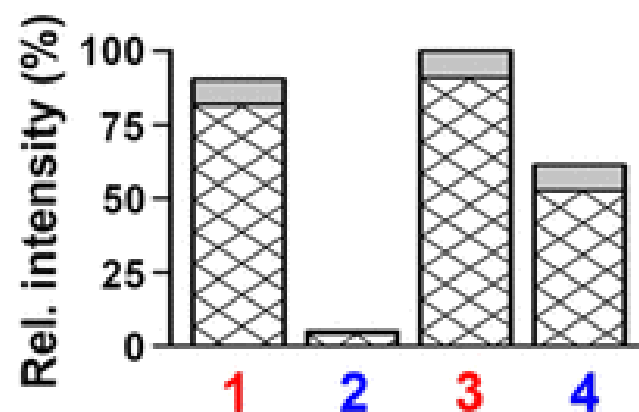
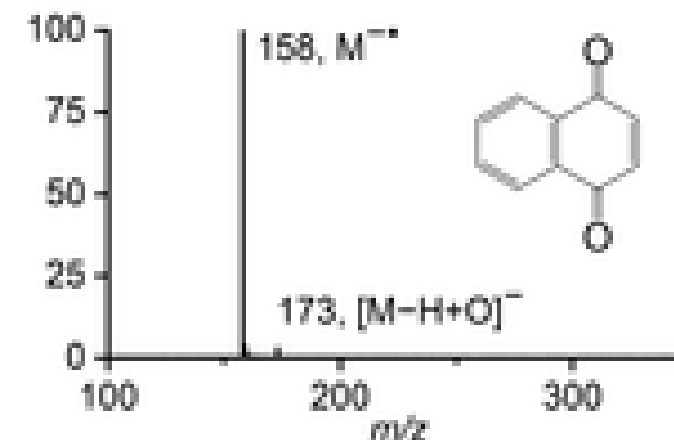
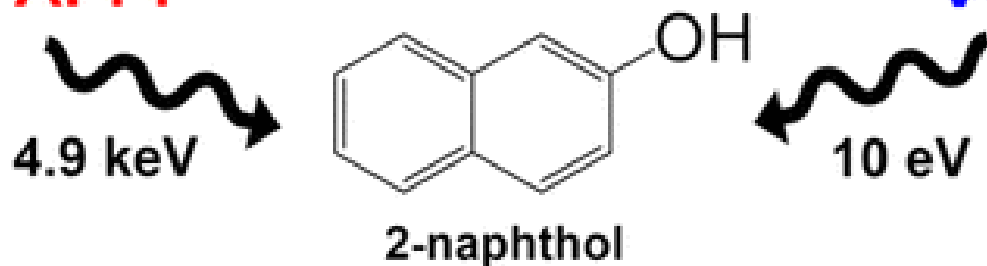
# Research

- Method development
- MS imaging
- (Single cell) proteomics and metabolomics

# Method development: photoionization

**Soft X-ray APPI**

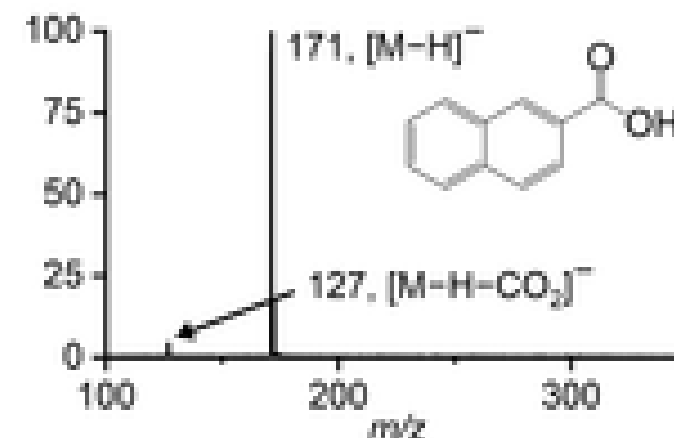
**VUV APPI**



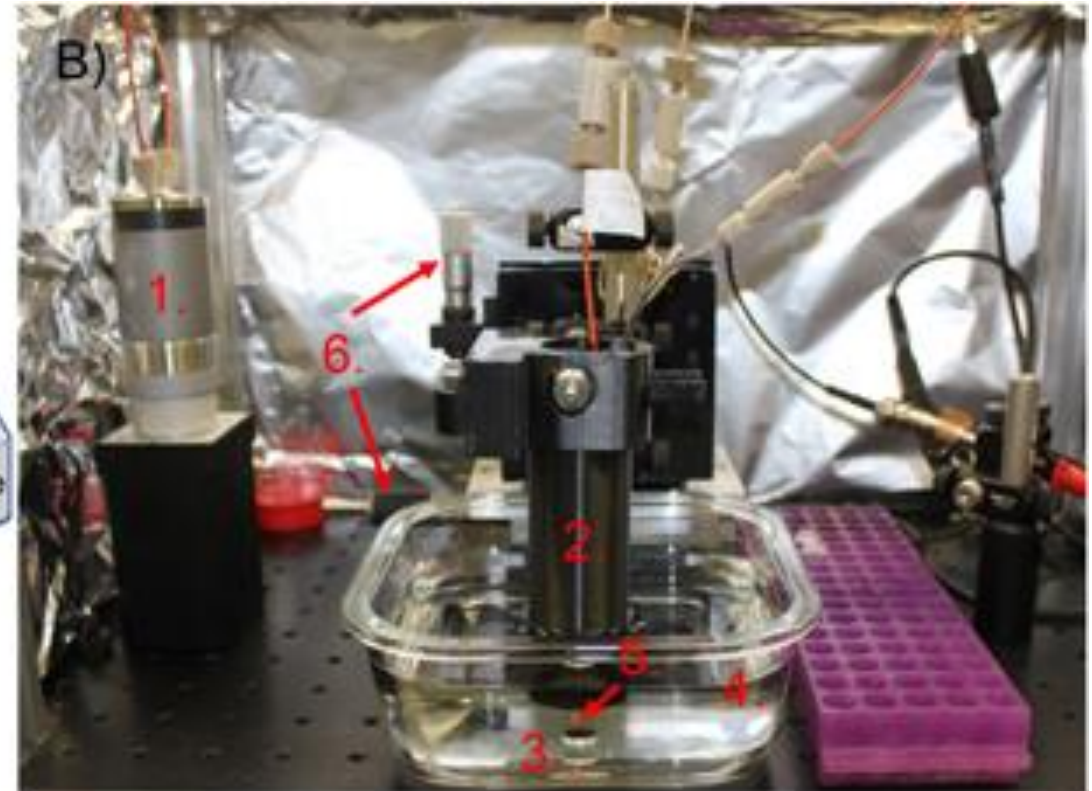
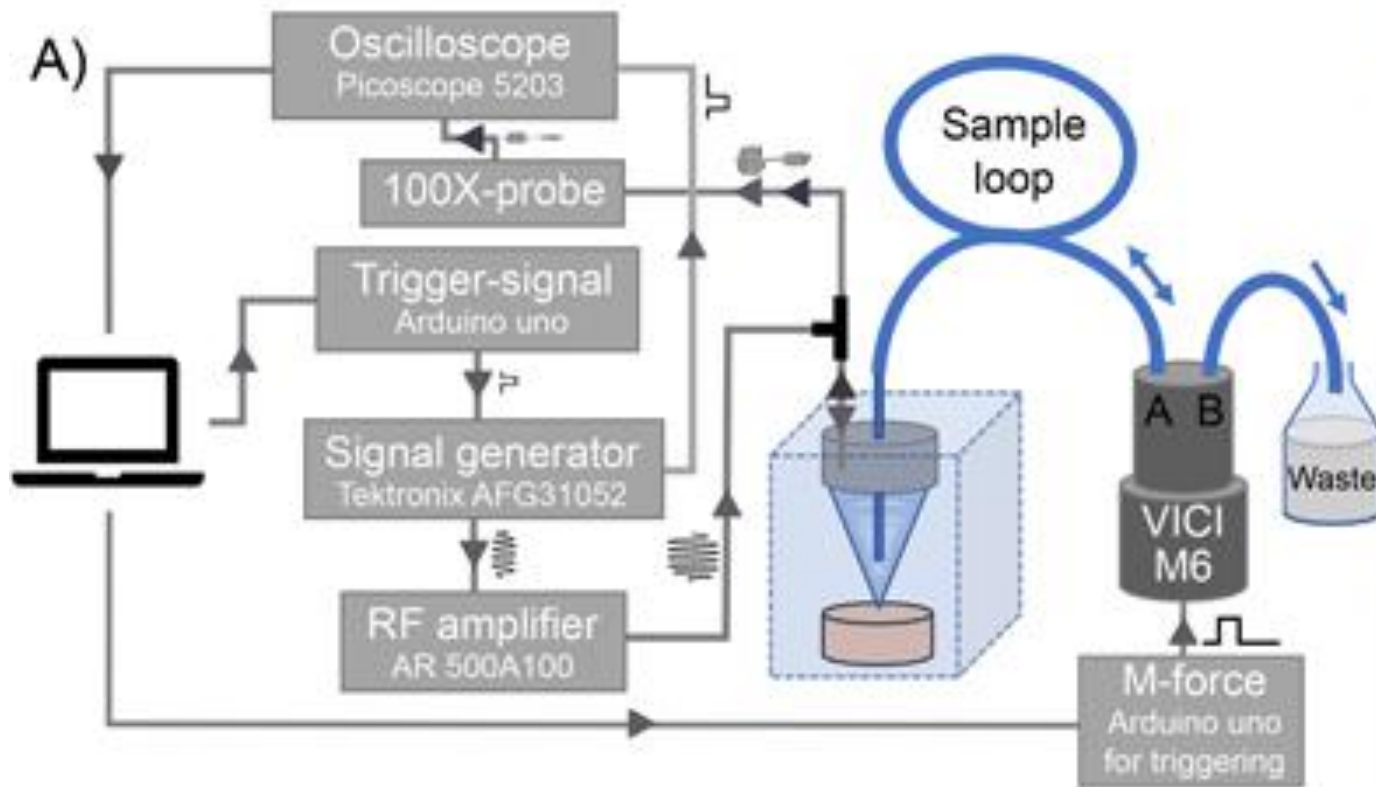
  $[M-H]^{-}$

 Substitution products

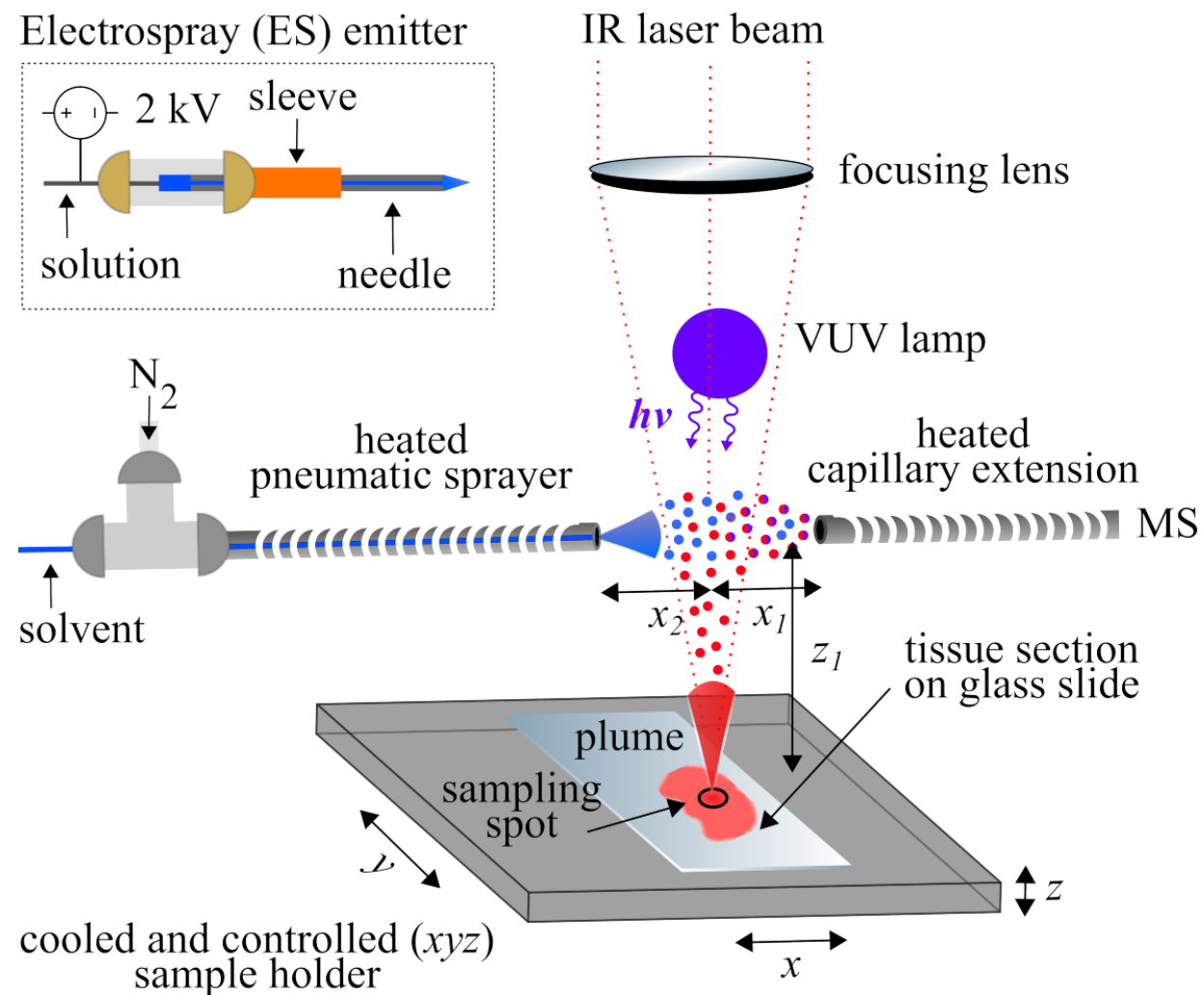
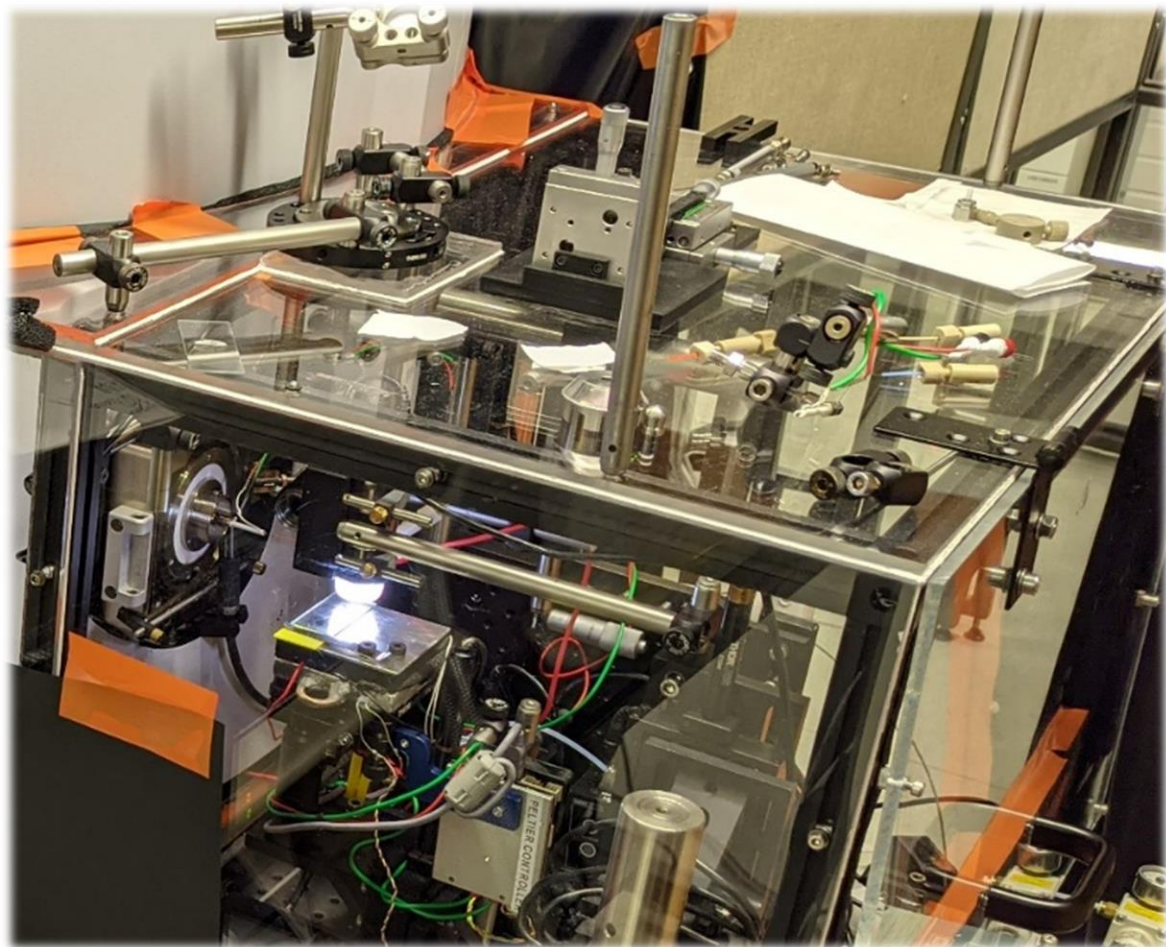
1 = Soft X-ray APPI  
2 = VUV APPI  
3 = Soft X-ray APPI + dopant  
4 = VUV APPI + dopant

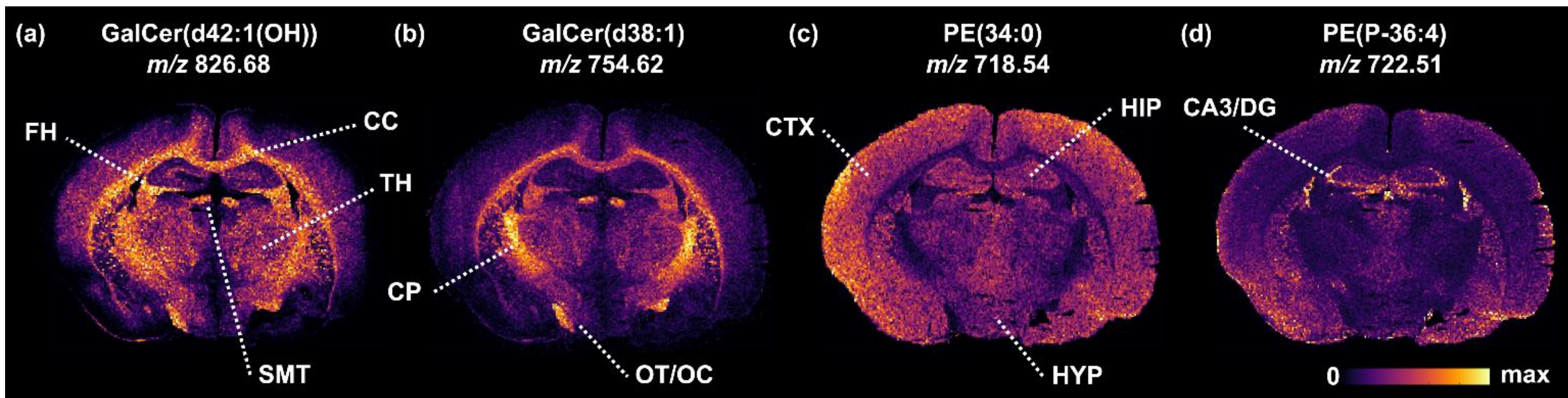


# Method development: ultrasound sampling

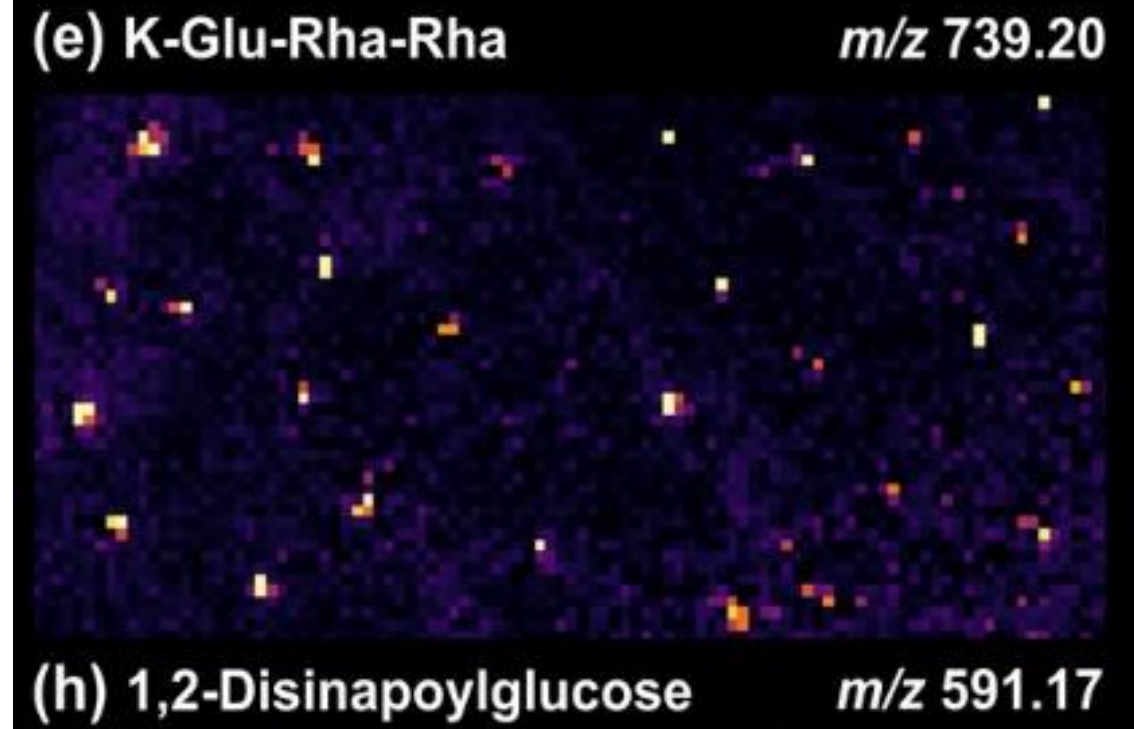
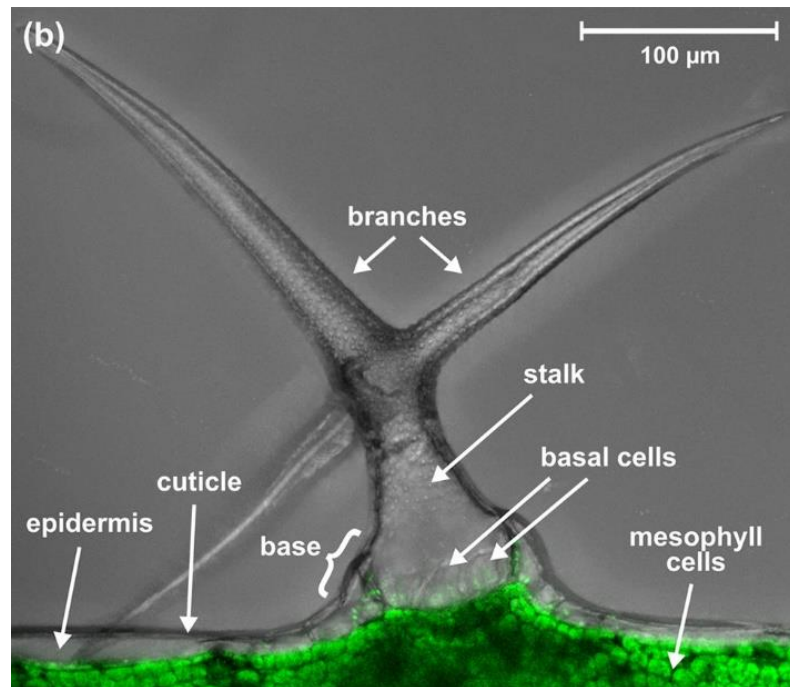
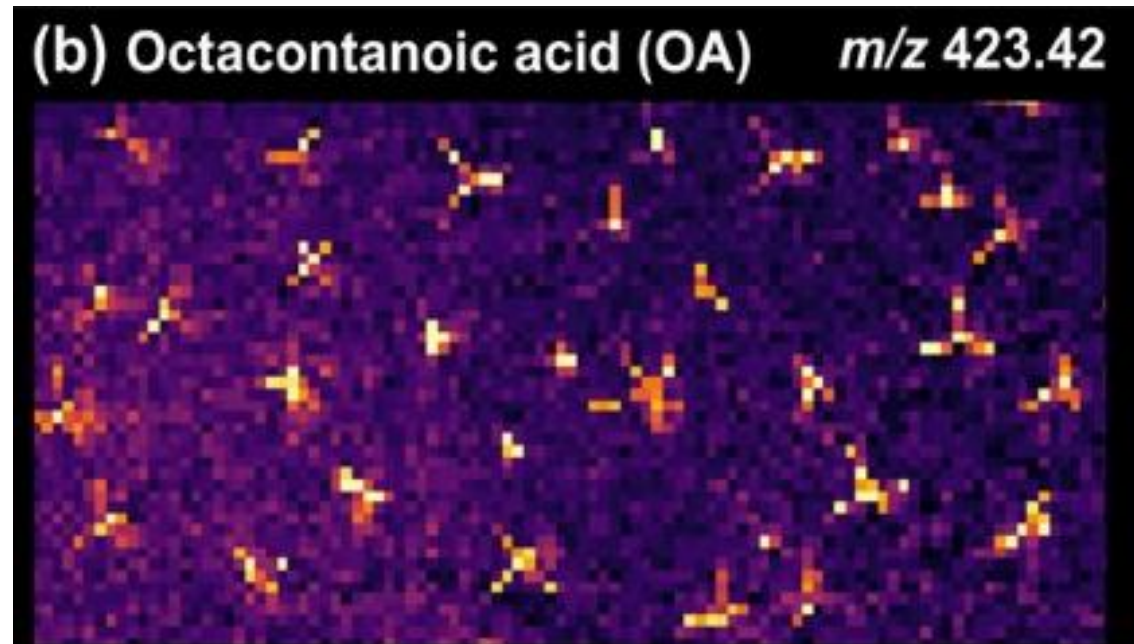


# MS imaging

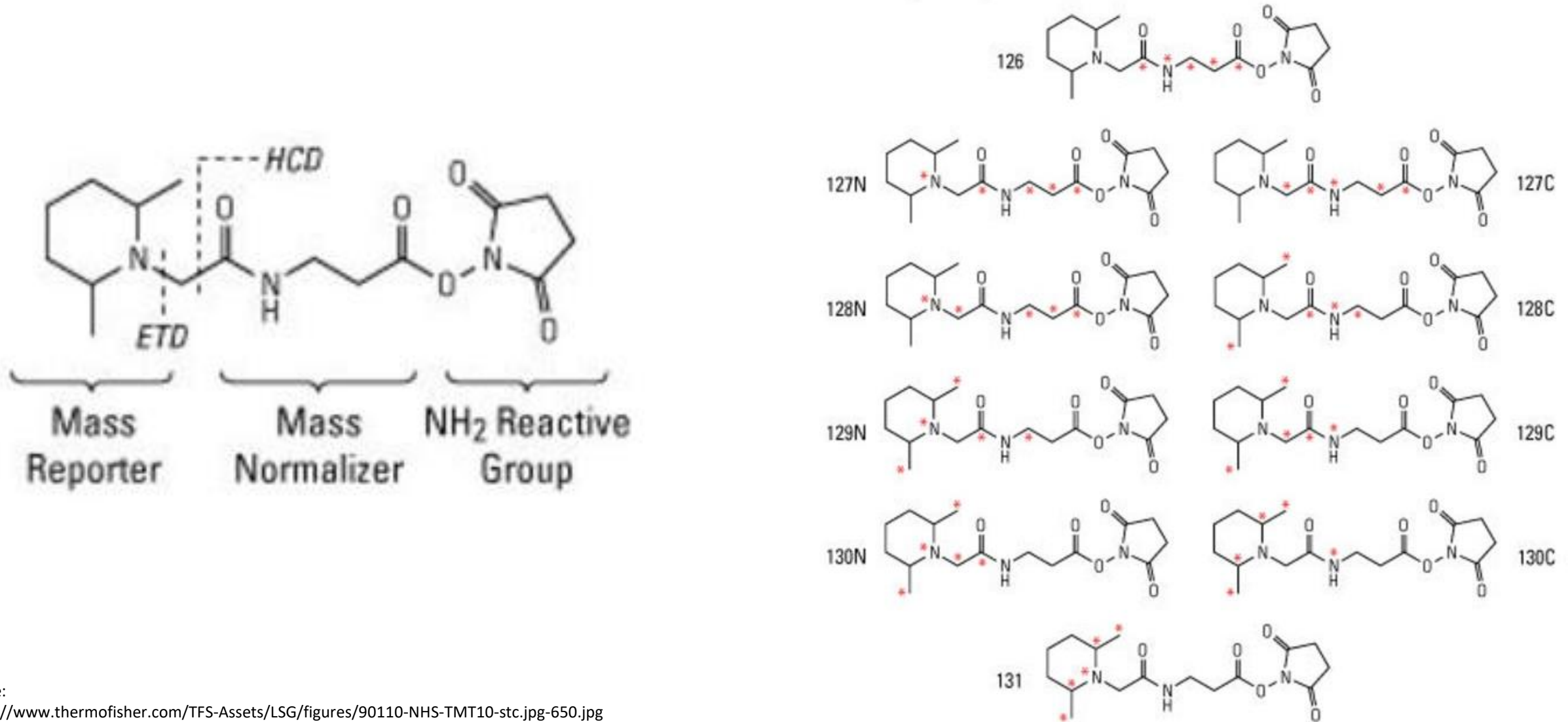




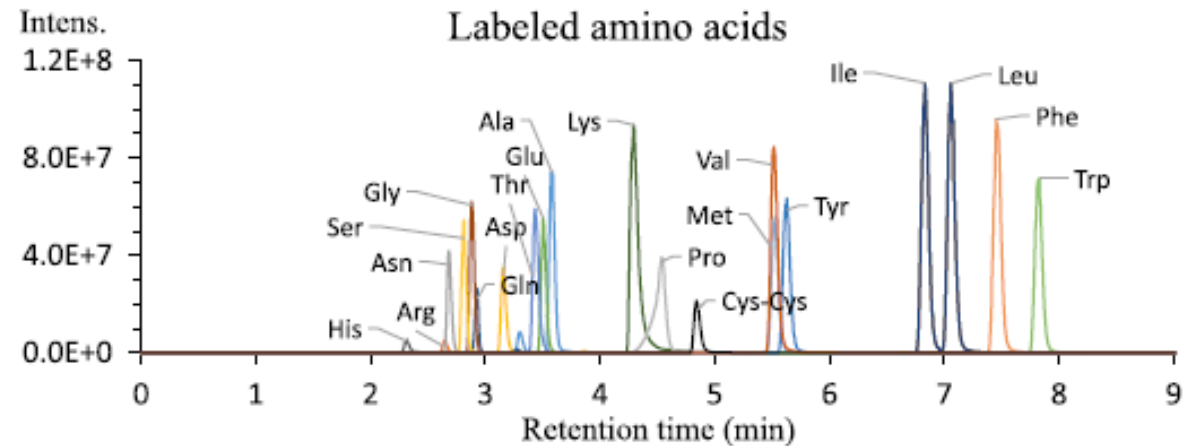
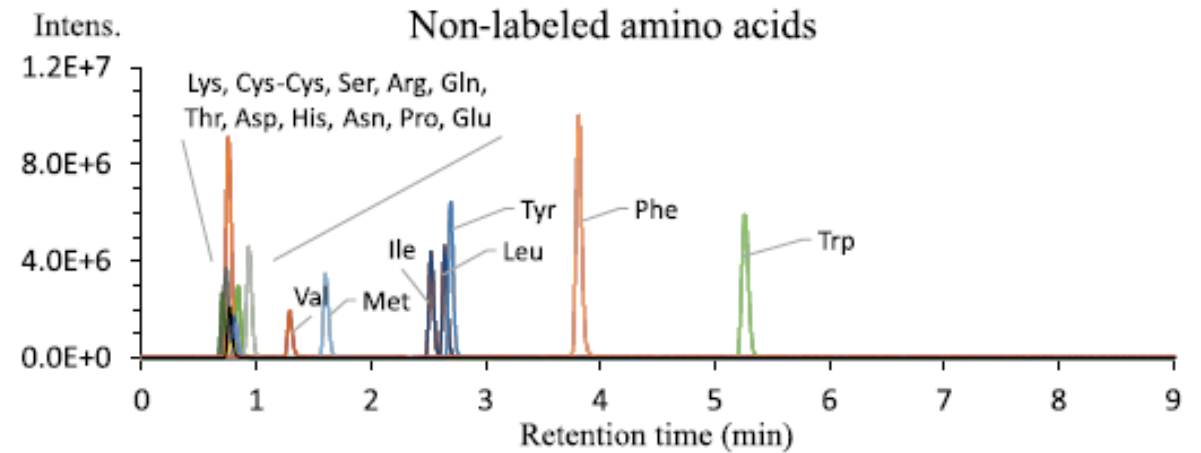
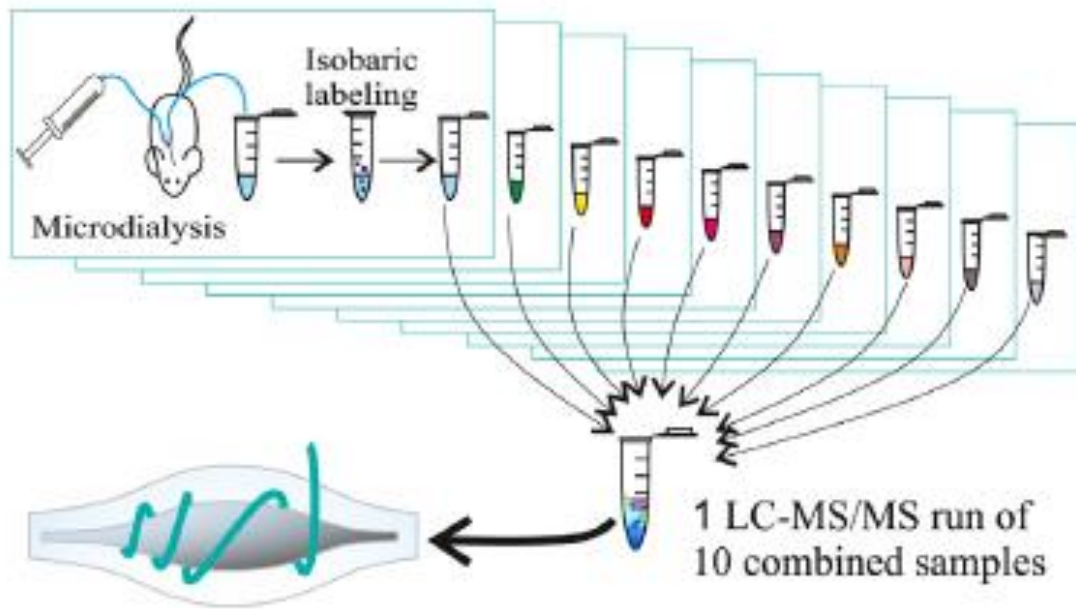




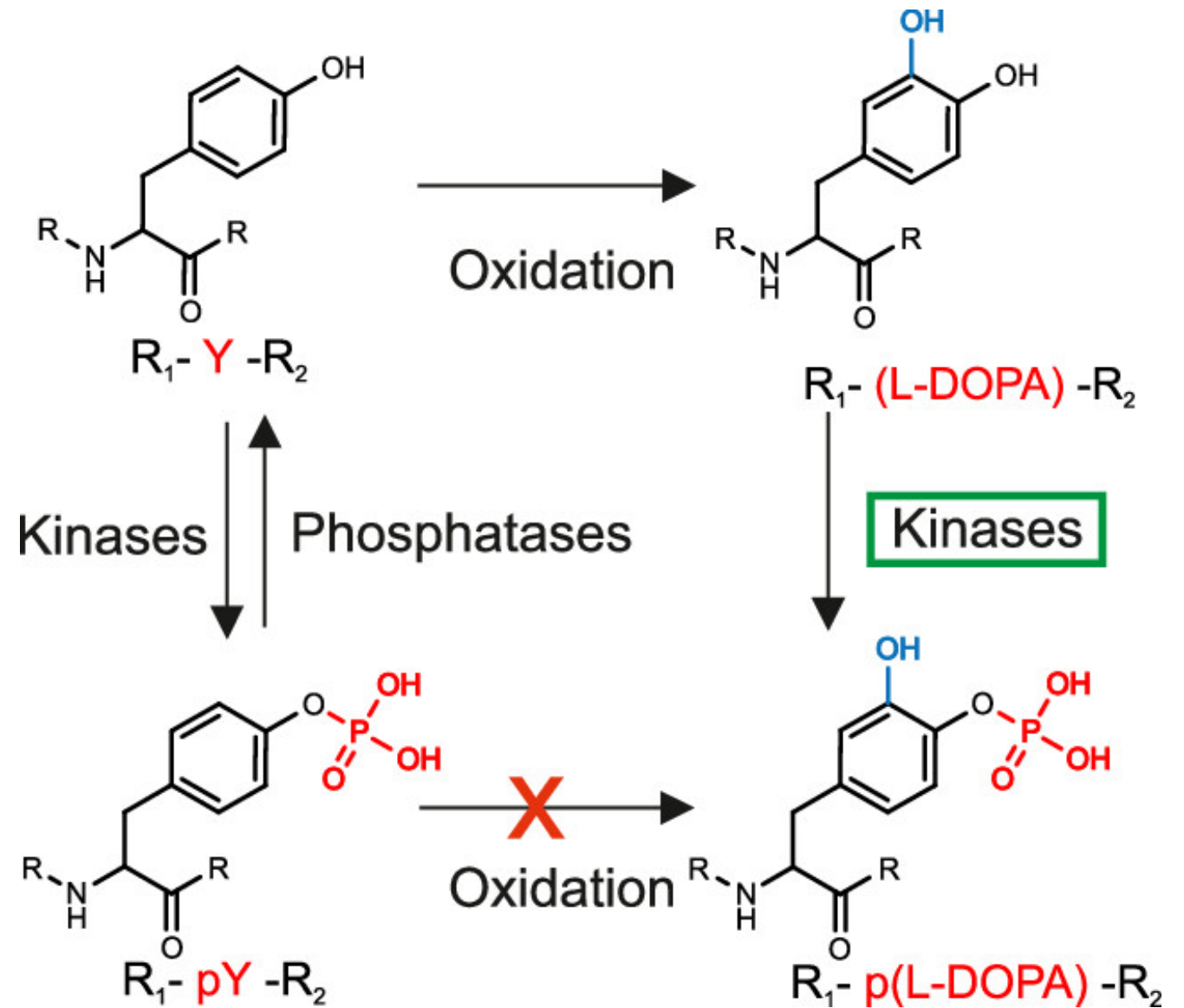
# Proteomics and metabolomics



# Metabolomics



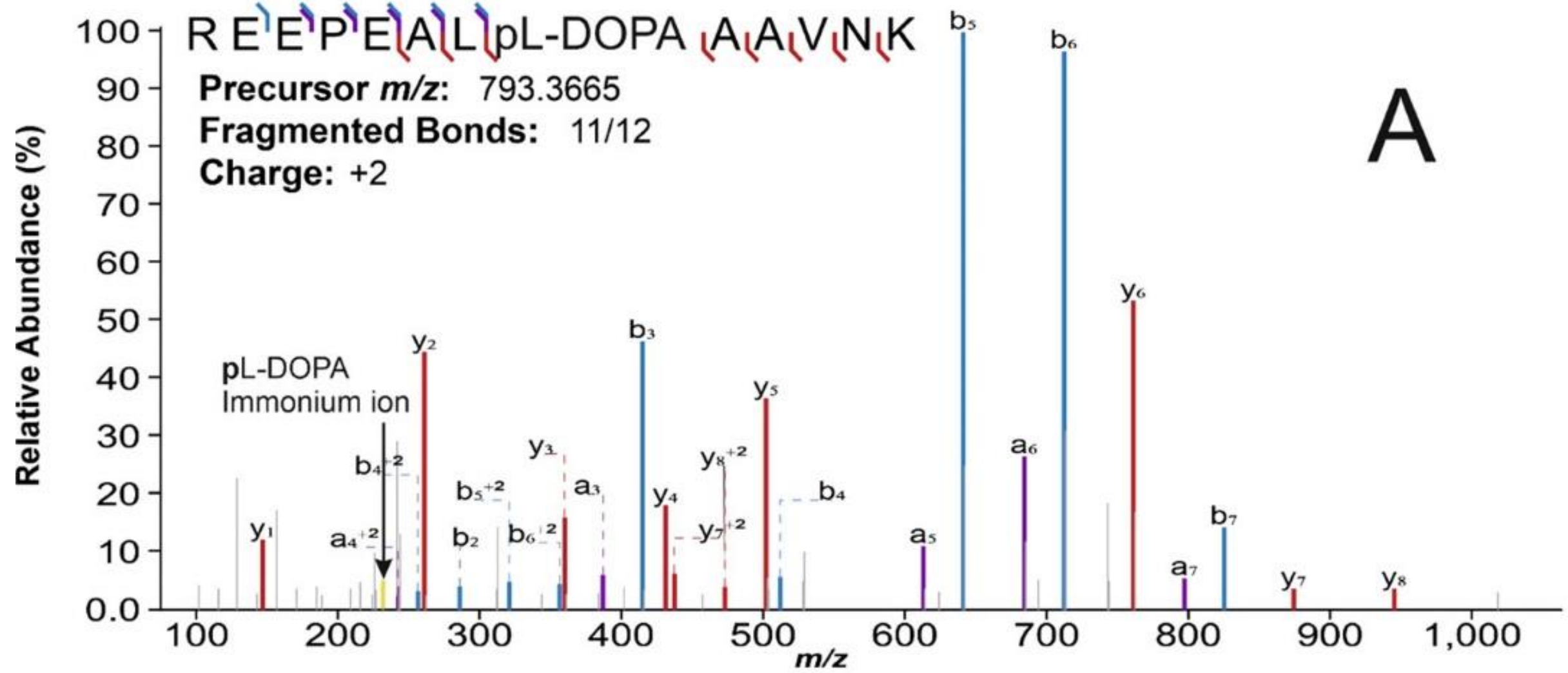
# Proteomics



R E E P E A L pL-DOPA A A V N K

Precursor  $m/z$ : 793.3665  
Fragmented Bonds: 11/12  
Charge: +2

A



Thank you!