



Aspect Analytics

Mass spectrometry imaging and machine learning
- a fruitful marriage.

AGENTSCHAP
INNOVEREN &
ONDERNEMEN



Vlaanderen
is ondernemen



Accelerated by
Health



INCEPTION PROGRAM



KU LEUVEN



Mass spectrometry imaging

A brief introduction

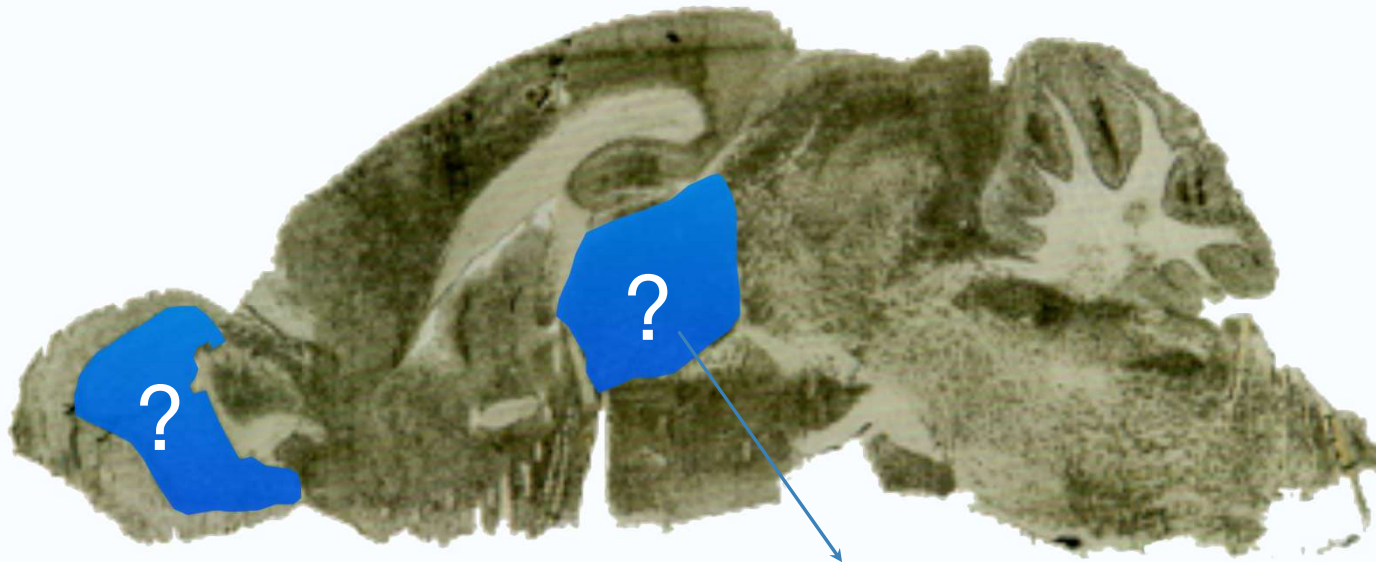
Mass spectrometry imaging workflow

Molecular imaging technology that
links chemical and spatial information together



Mass spectrometry imaging workflow

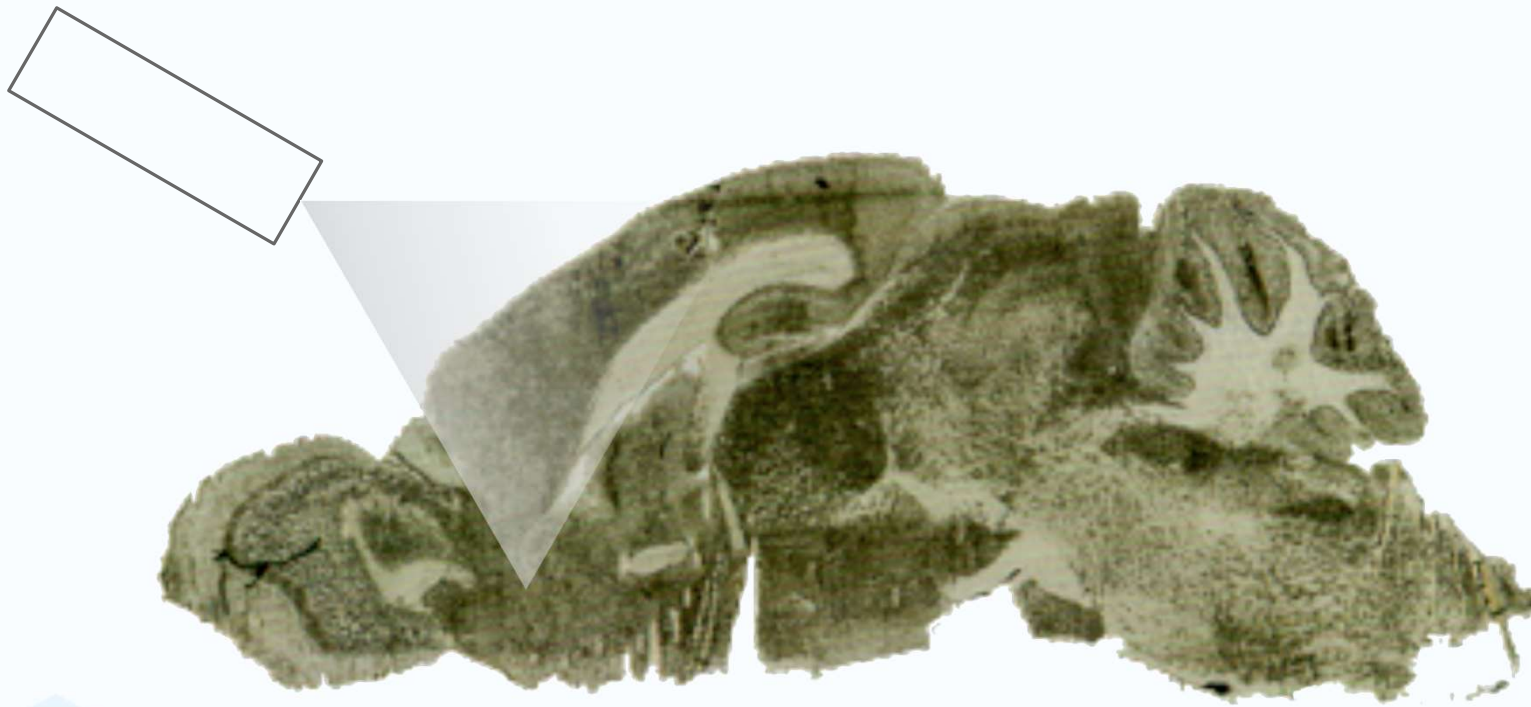
Molecular imaging technology that links chemical and spatial information together



Which biomolecules are located in this part of the mouse brain?

Mass spectrometry imaging workflow

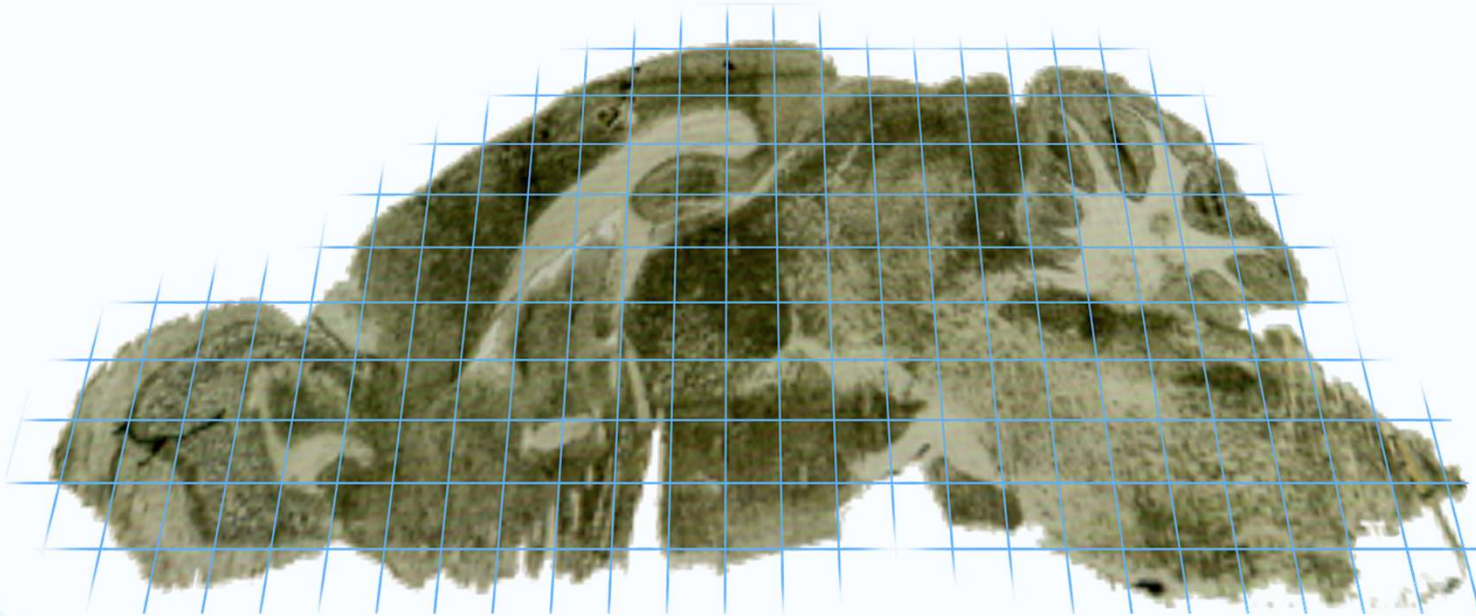
Matrix application



Mass spectrometry imaging workflow

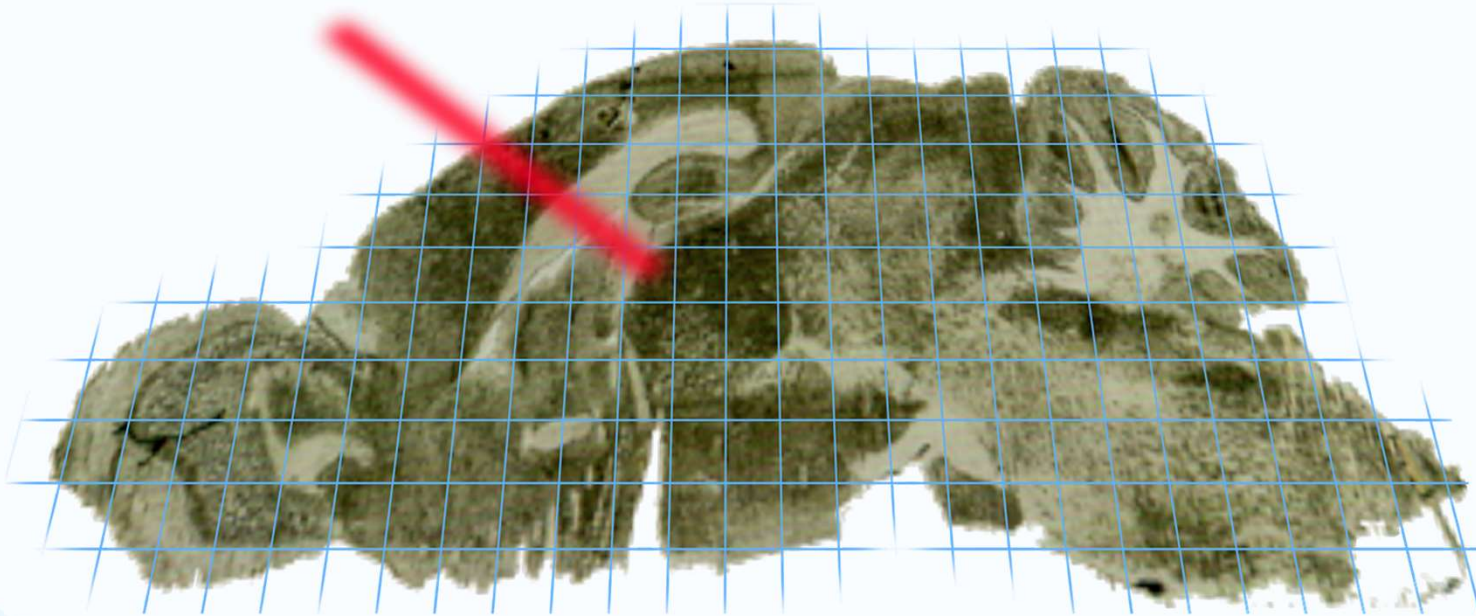


Overlay tissue slice with virtual grid



Mass spectrometry imaging workflow

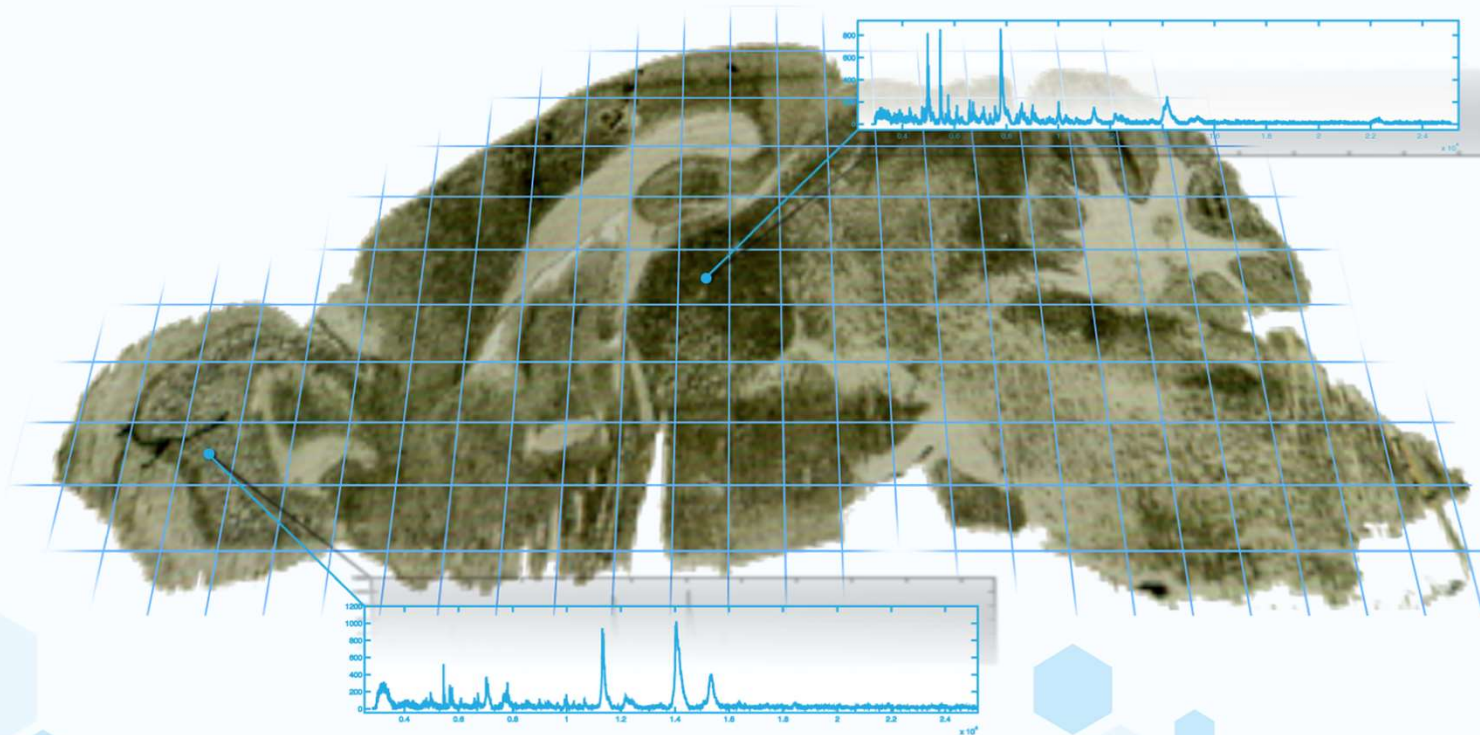
Perform MALDI mass spectrometry at each gridpoint

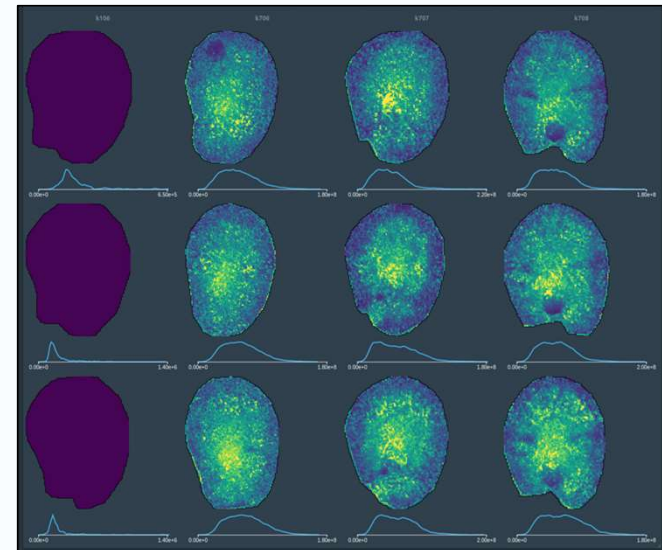
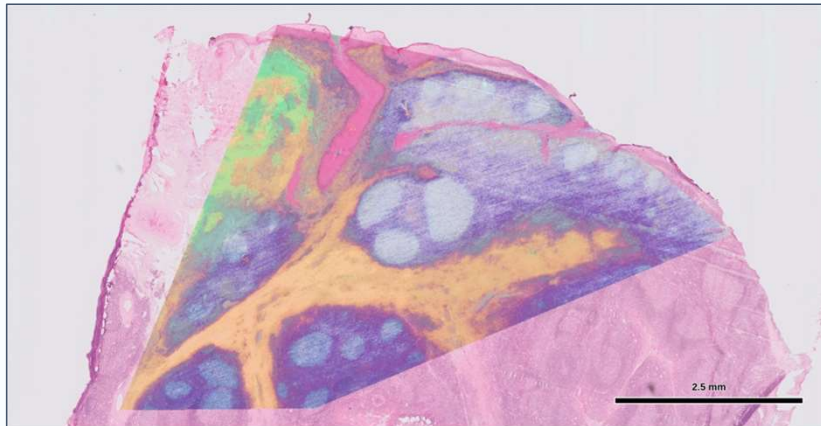


MALDI = Matrix Assisted Laser Desorption and Ionization

Mass spectrometry imaging workflow

Result: mass spectrum at each gridpoint



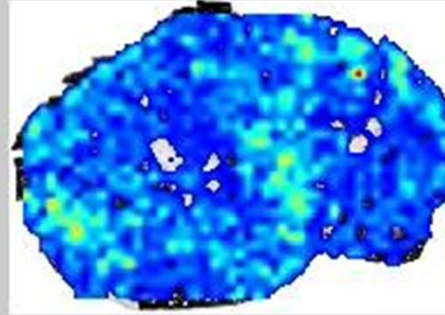


Biomarker discovery & clinical diagnostics

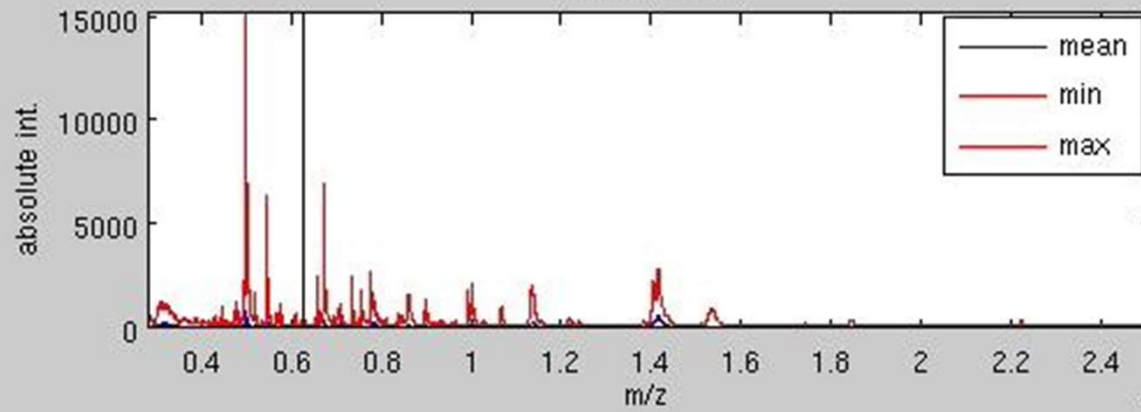
Drug discovery

How did I end up here?

6257.9604 m/z



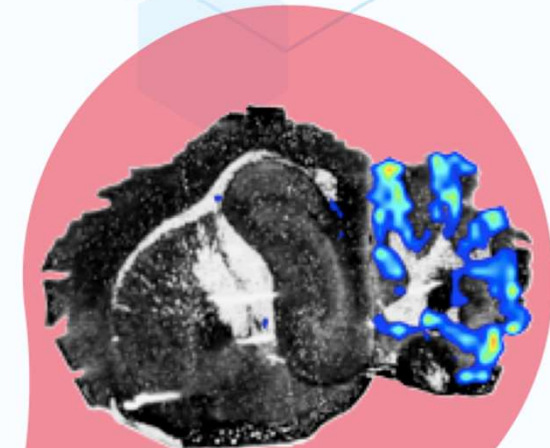
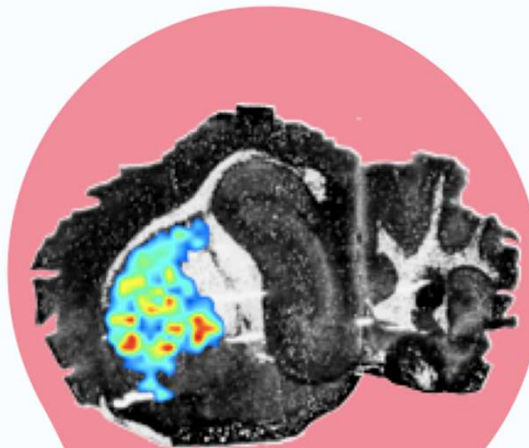
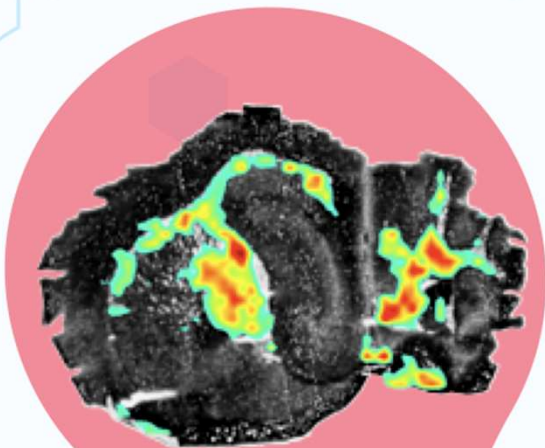
Mass spectra



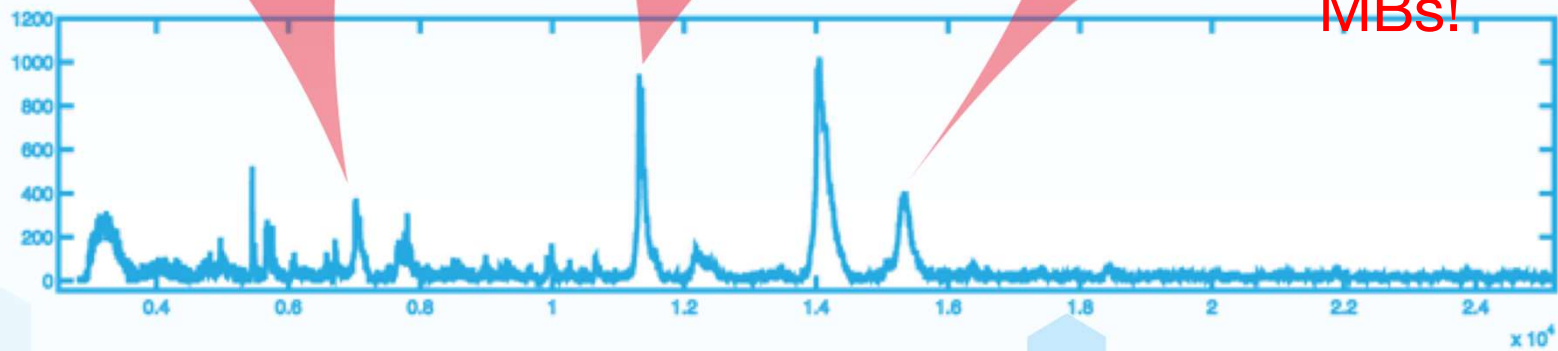
Prof. Raf Van de Plas



Prof. Etienne Waelkens



32 whole MBs!



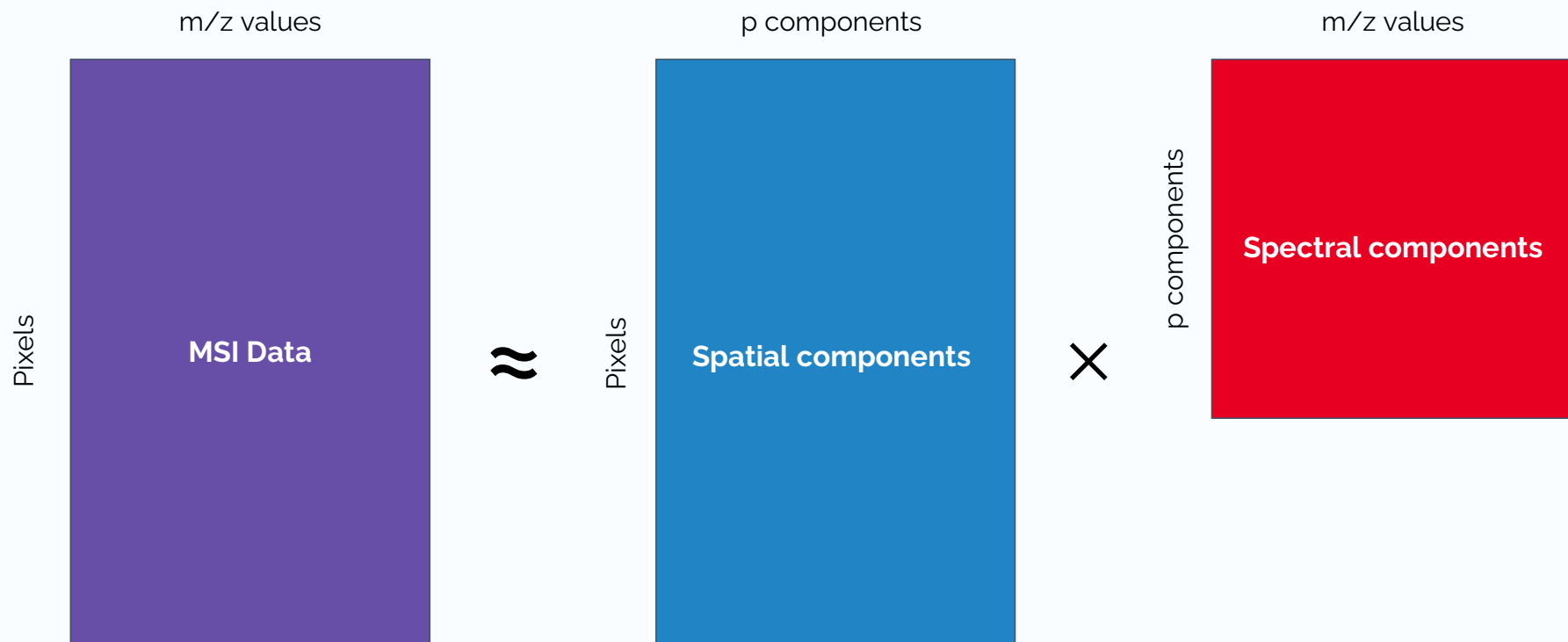


- 10,000's of ion images in a single experiment
- ~1.5 TB data in a single experiment

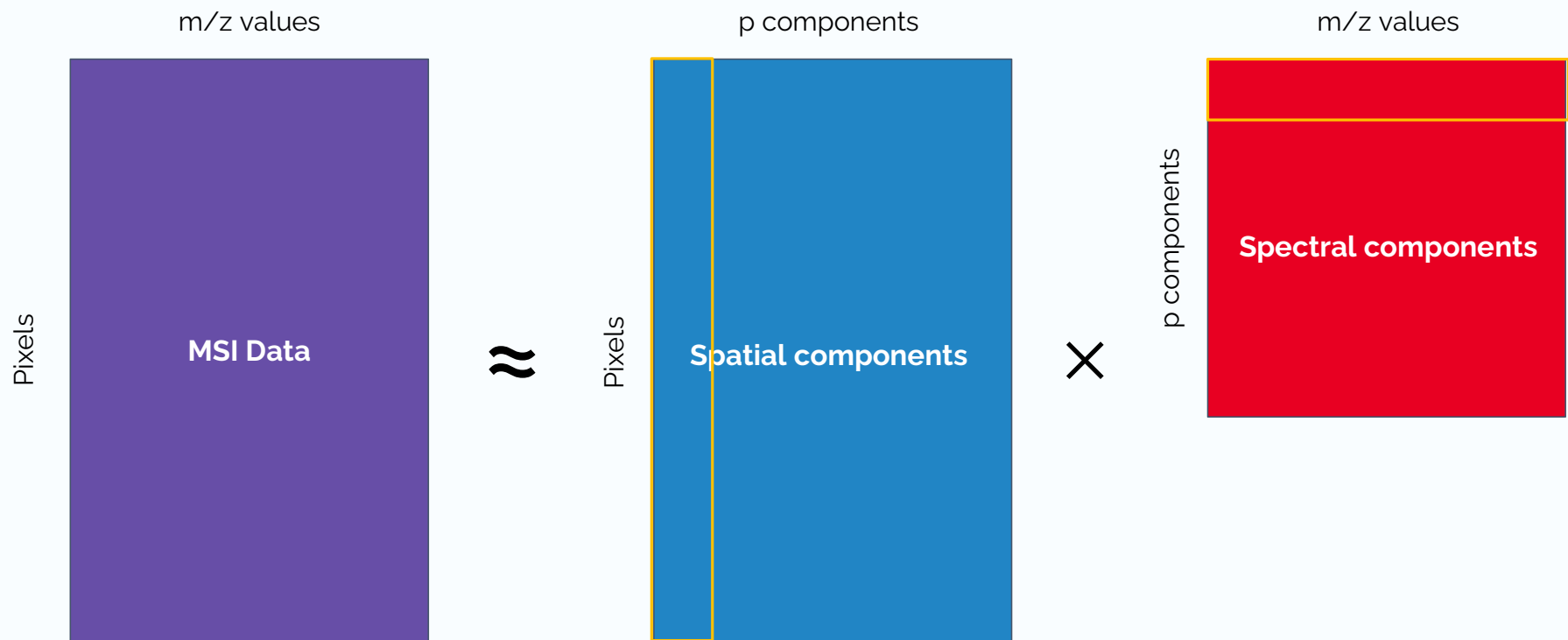


How to deal with all that data?

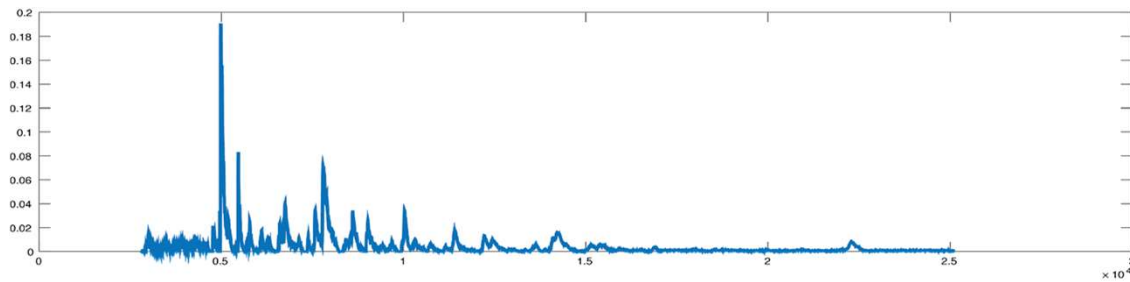
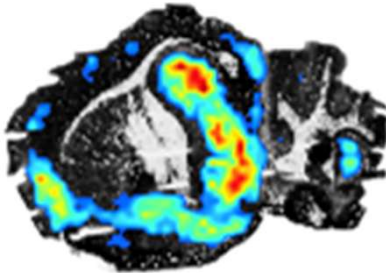
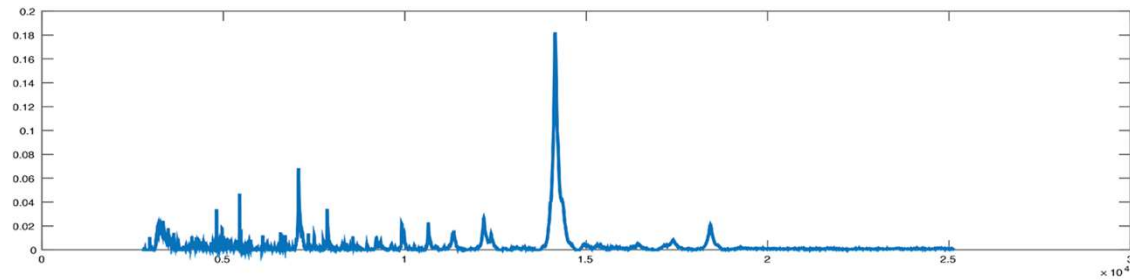
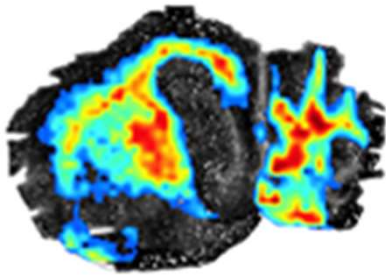
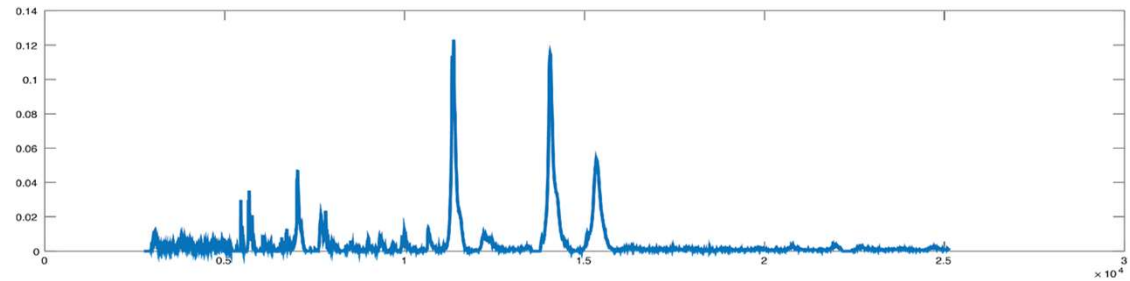
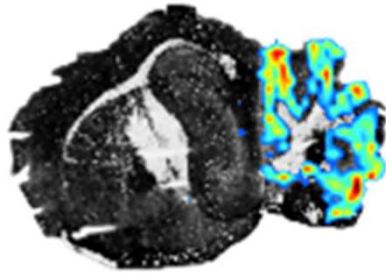
Matrix decomposition



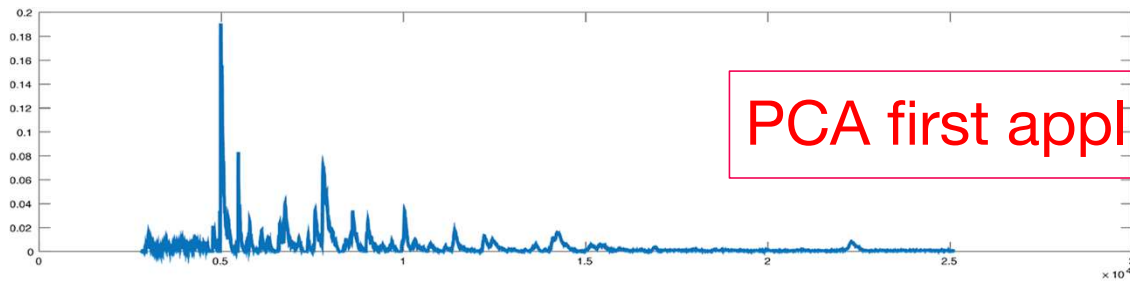
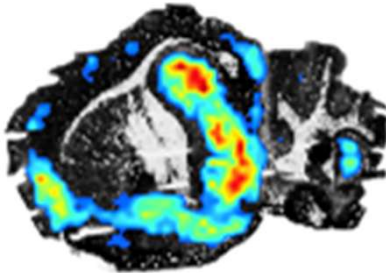
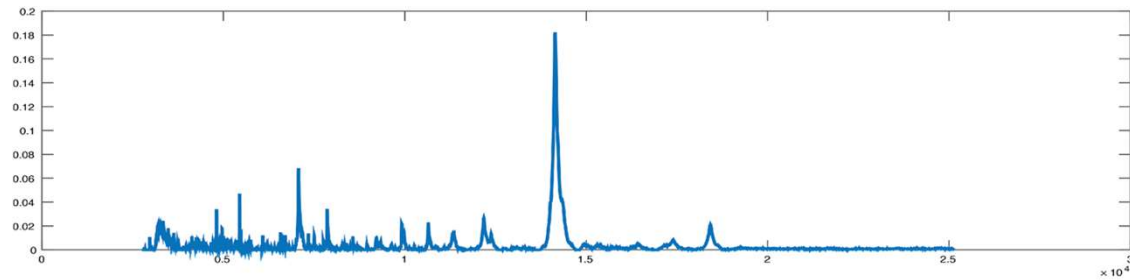
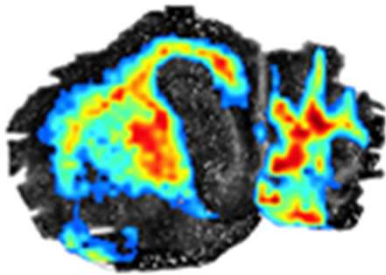
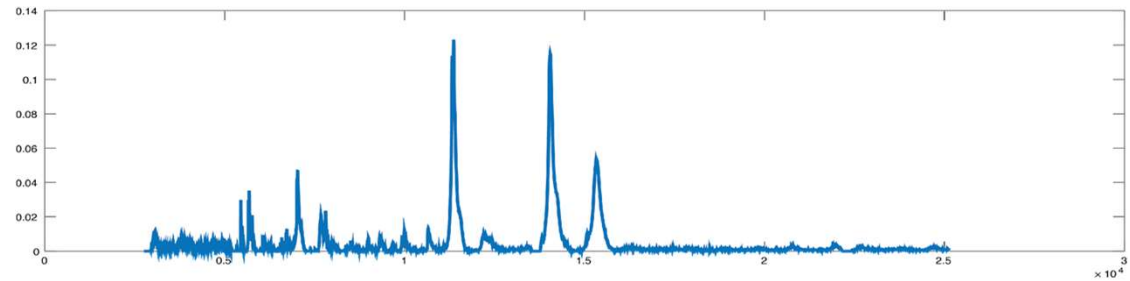
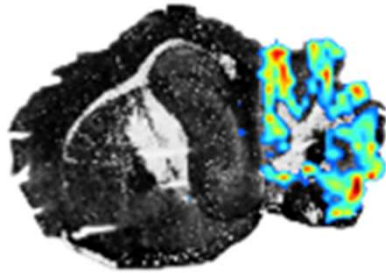
Matrix decomposition



Matrix decomposition (NMF)

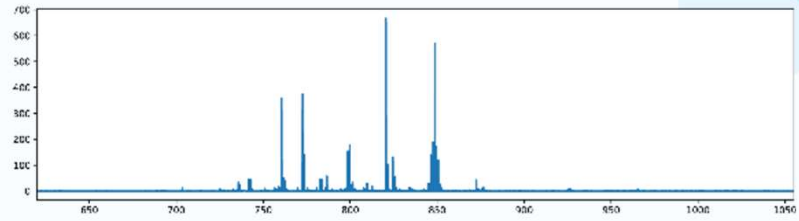
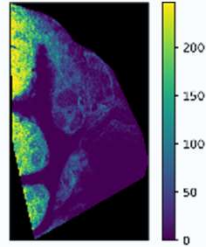


Matrix decomposition (NMF)

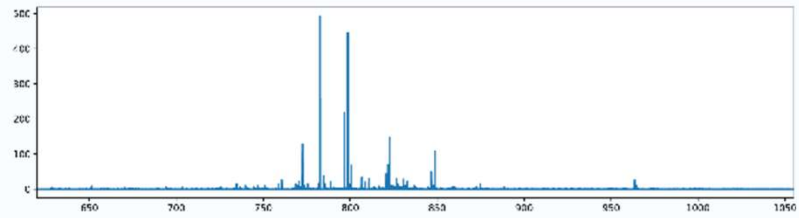
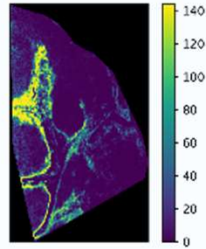


PCA first applied in 2007!

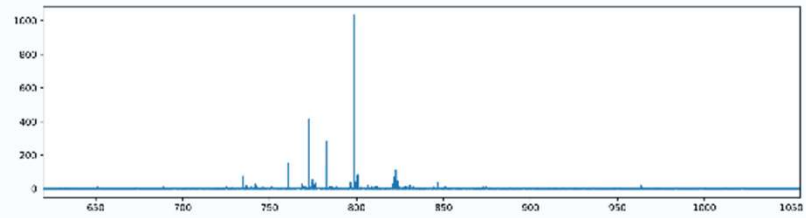
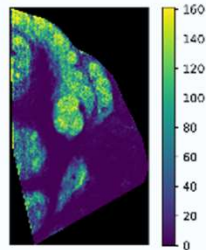
NNC 1



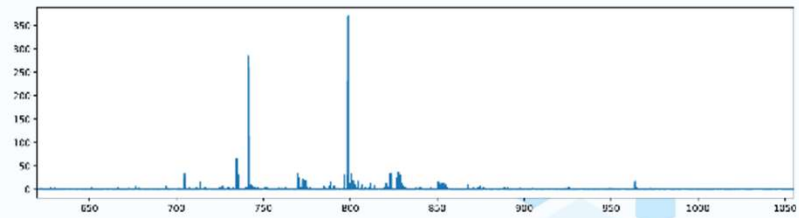
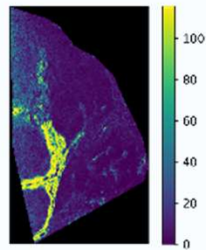
NNC 2



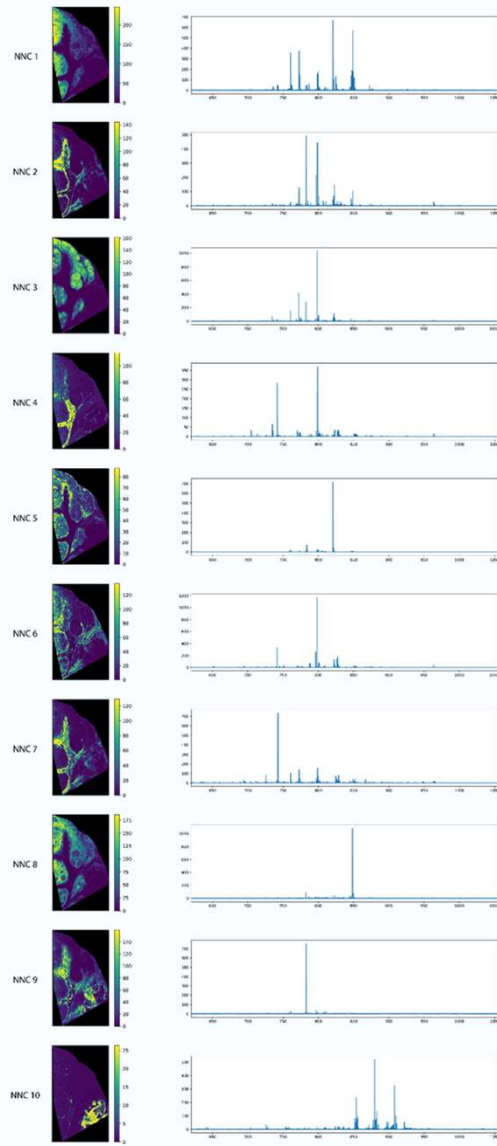
NNC 3



NNC 4

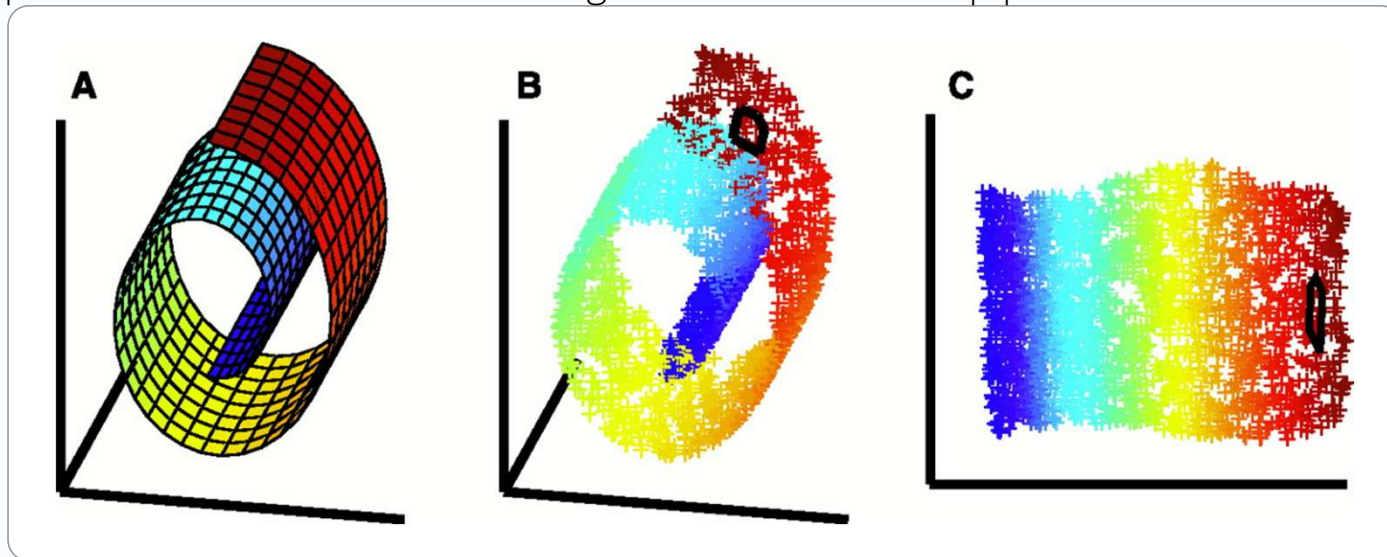


Still bread
and
butter!



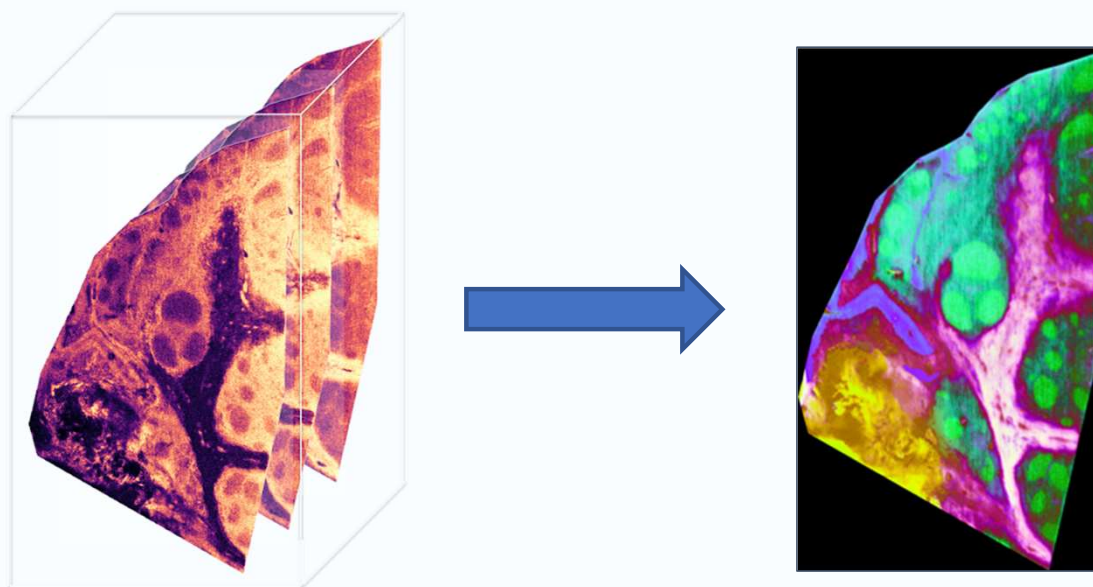
Manifold learning: concept

Capture local structure rather than global structure – keep points that look alike close



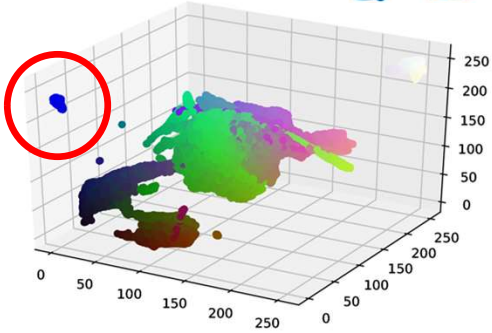
Example Locally Linear Embedding, Roweis et al., Science 2000

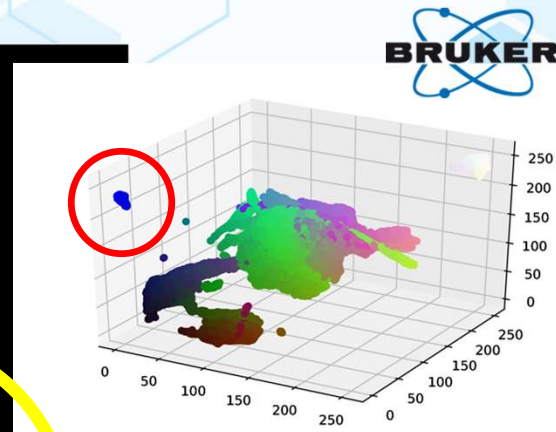
UMAP dimensionality reduction



Evaluation of Distance Metrics and Spatial Autocorrelation in Uniform Manifold Approximation and Projection Applied to Mass Spectrometry Imaging Data

T. Smets, N. Verbeeck, M. Claesen, A. Asperger, G. Griffioen, T. Tousseyn, W. Waelput, E. Waelkens, B. De Moor - 2019, Analytical Chemistry





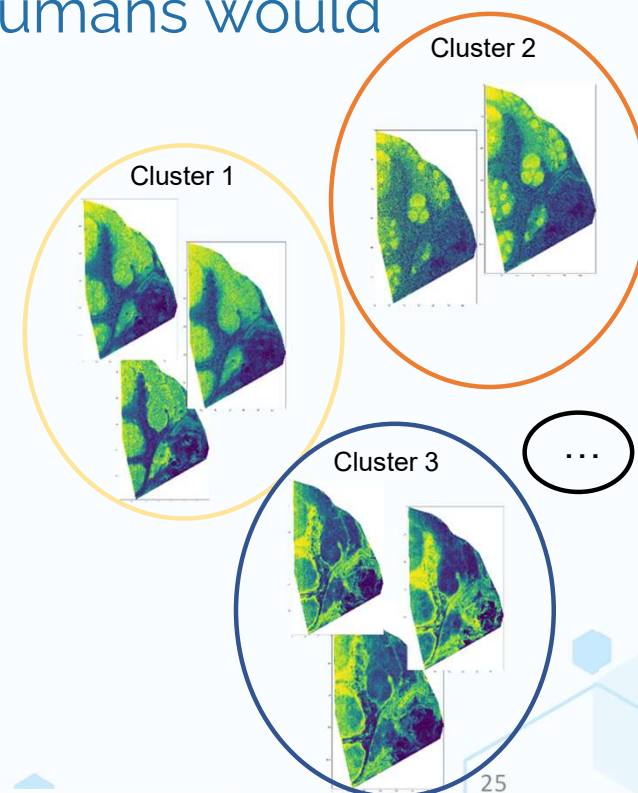
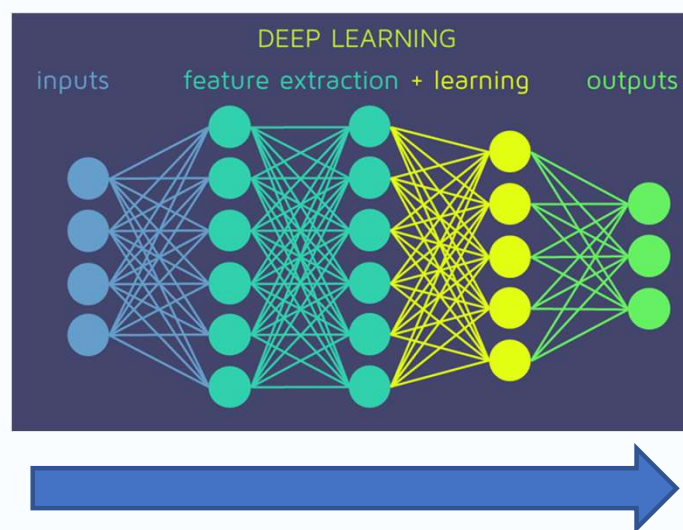
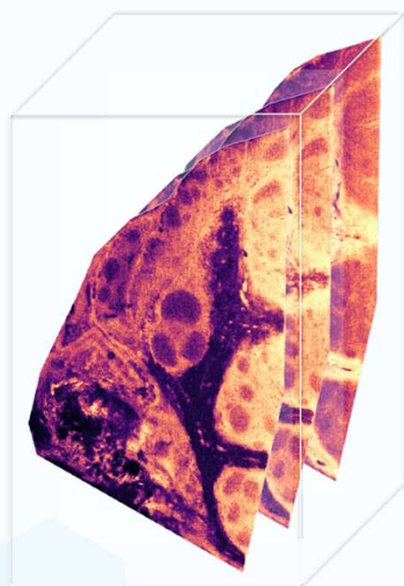
H&E image available in "Site-to-Site Reproducibility and Spatial Resolution in MALDI-MSI of Peptides from Formalin-Fixed Paraffin-Embedded Samples", Ly et al. <https://onlinelibrary.wiley.com/doi/10.1002/prca.201800029> (used with permission)



Wanqiu Zhang
Deep learning & AI

Clustering ion images

- Find similar images and group together as humans would

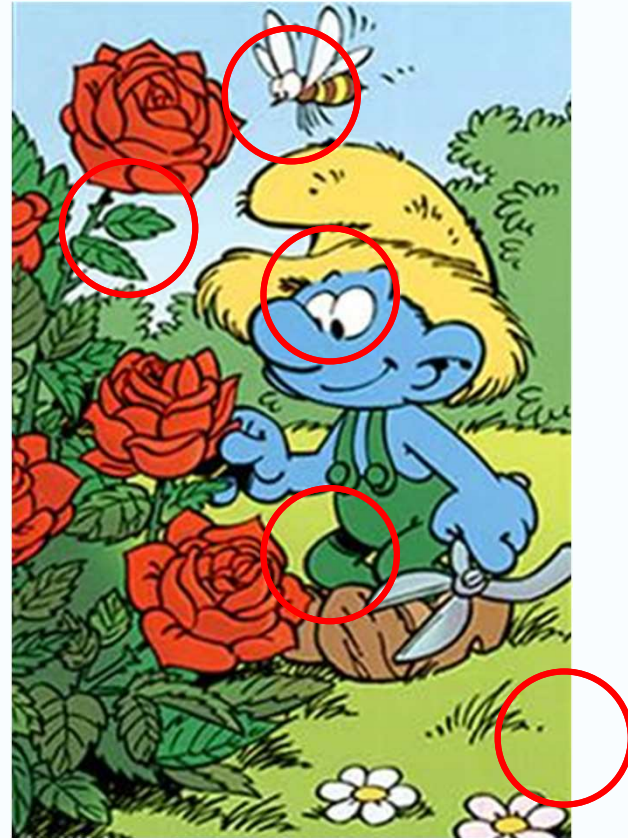
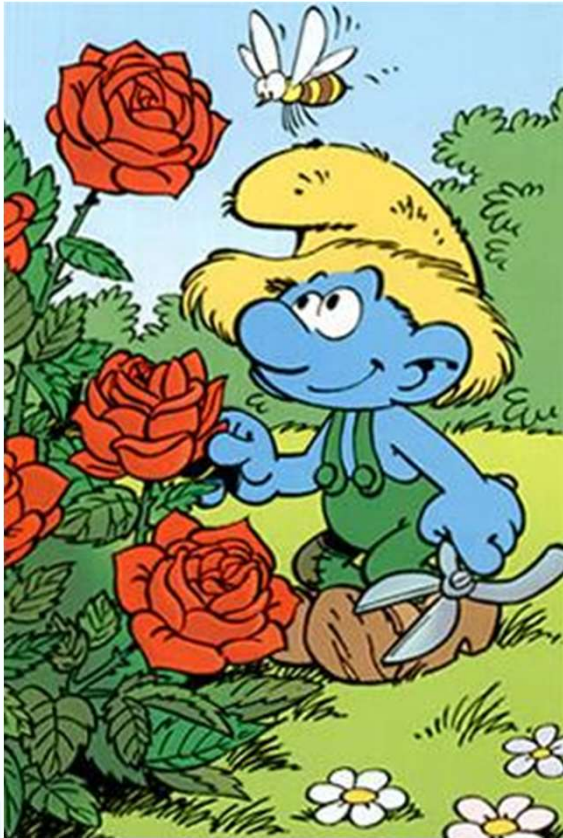


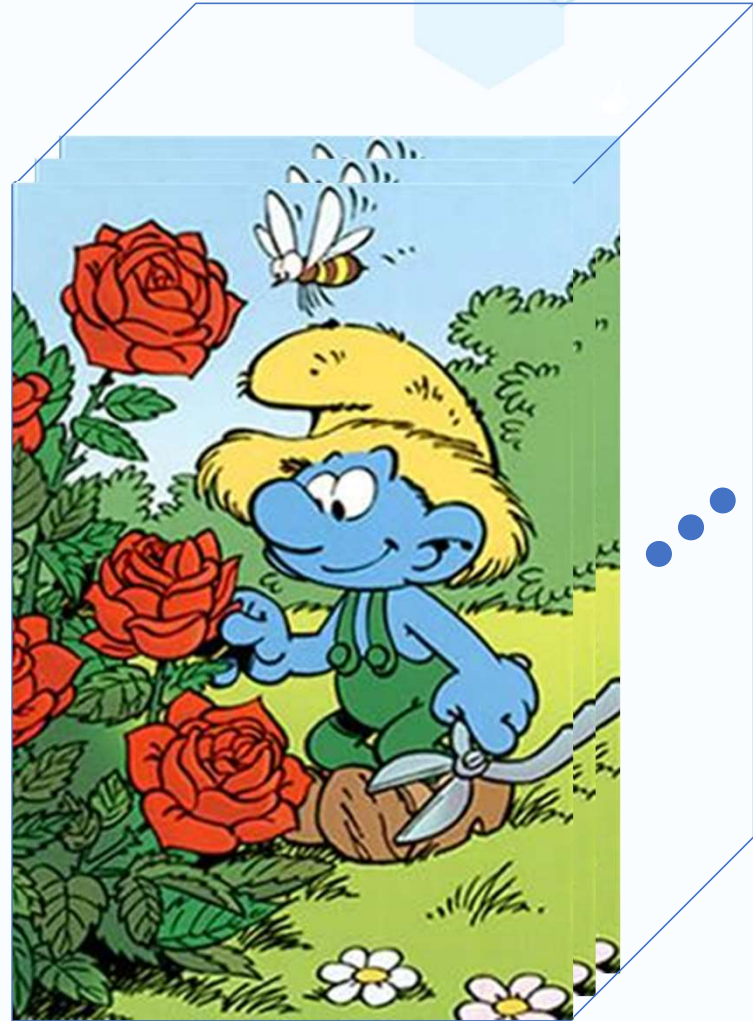
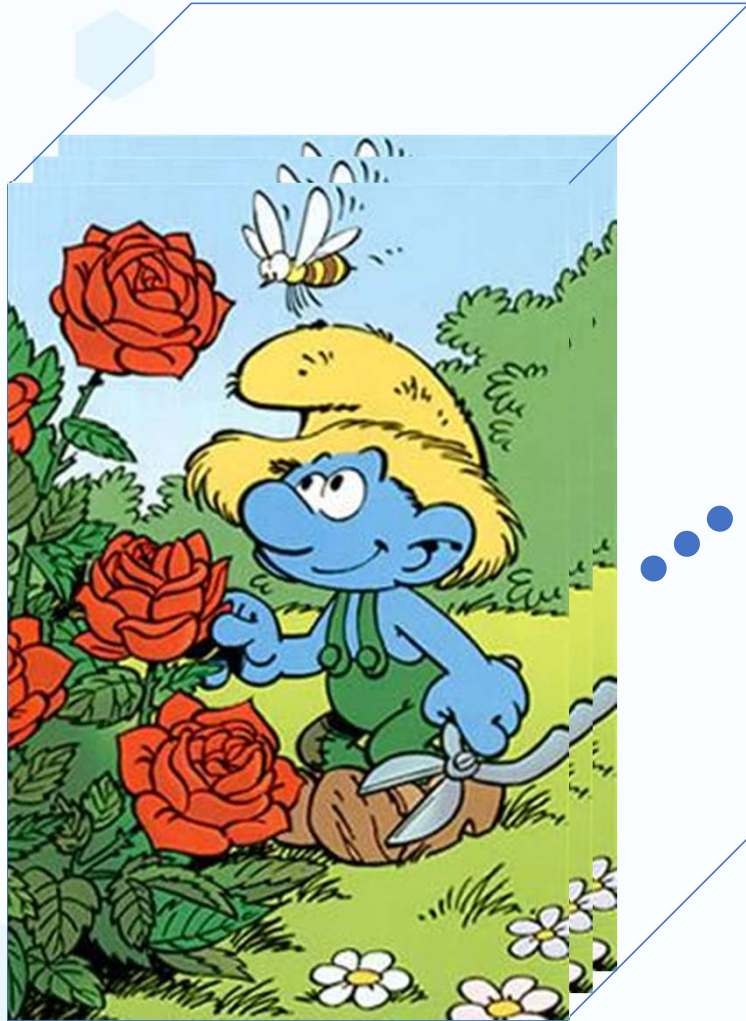
Comparing multiple MSI experiments

Spot the 5 differences

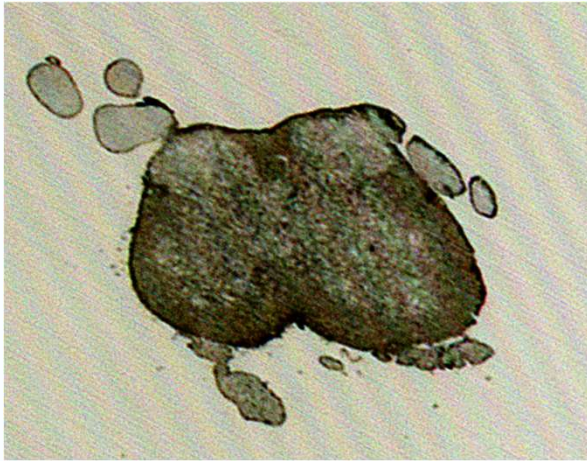


Spot the 5 differences





Multiple tissues



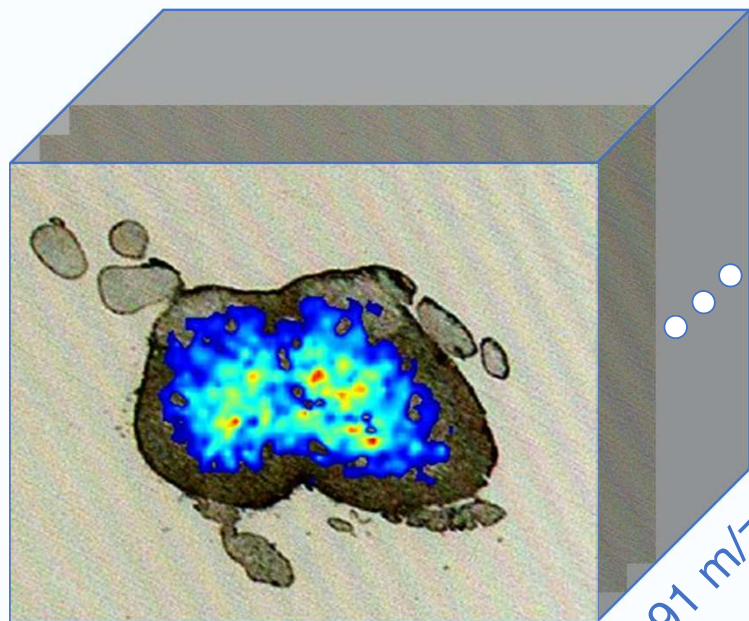
ALS
mouse model



Healthy

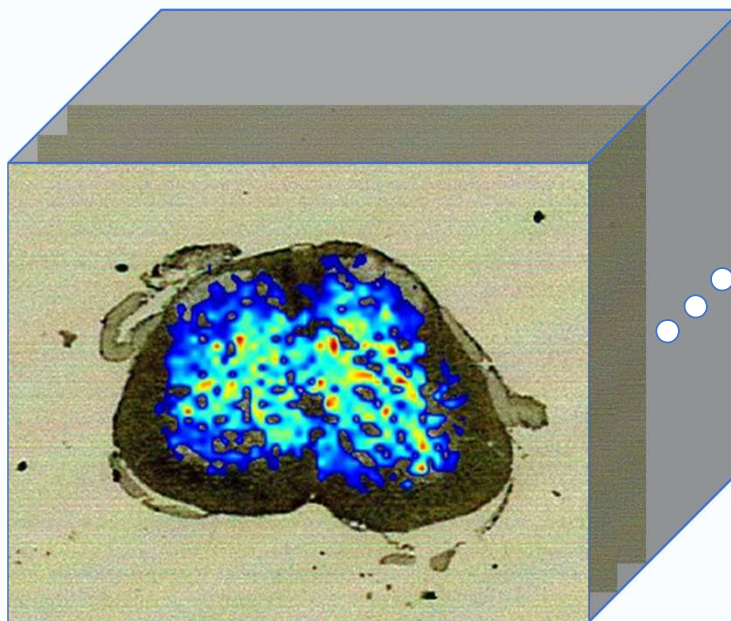
W. Robberecht, L. Vandebosch, N. Verbeeck,
R. Van de Plas, E. Waelkens, B. De Moor

Multiple tissues



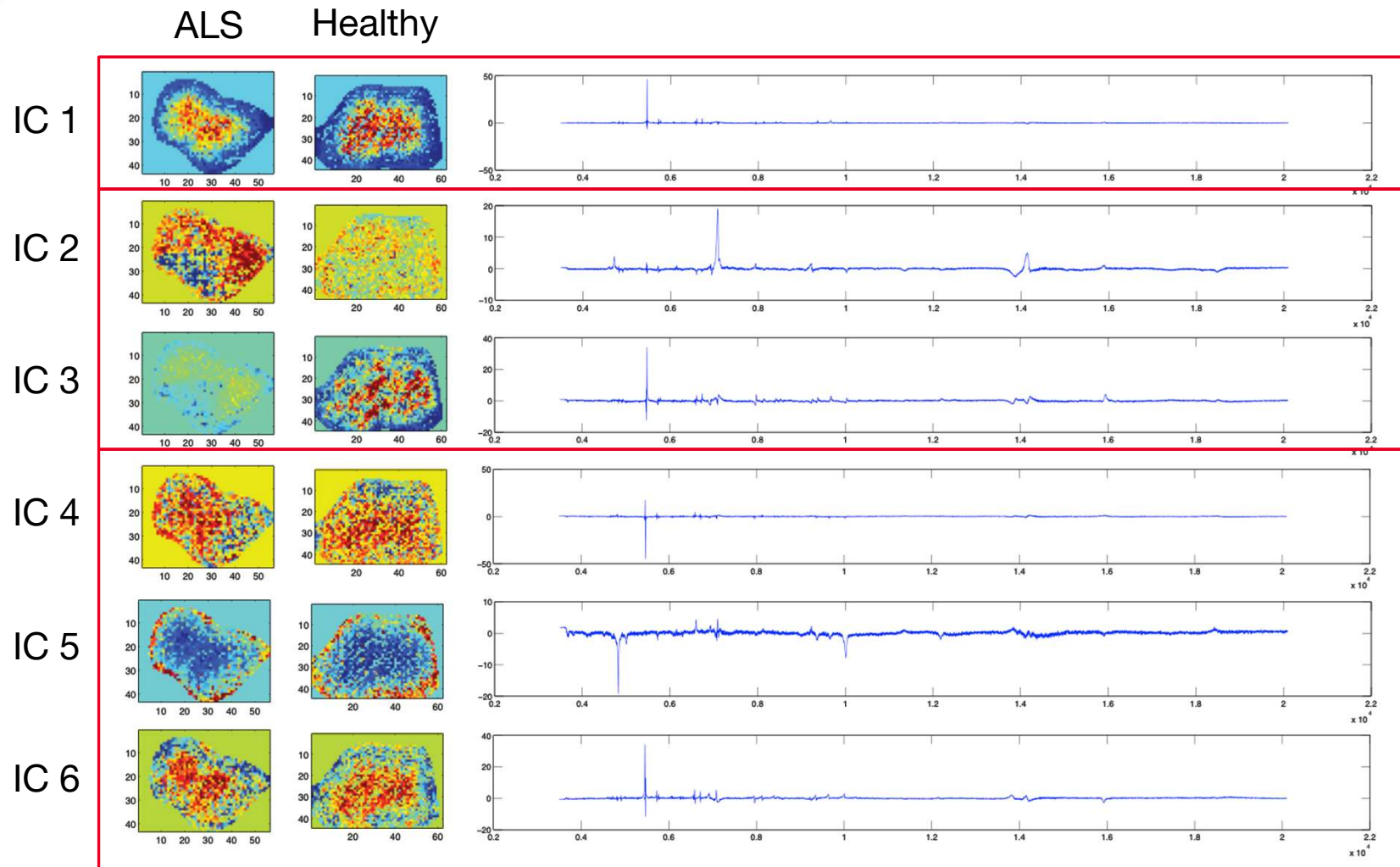
15691 m/z-bins

ALS

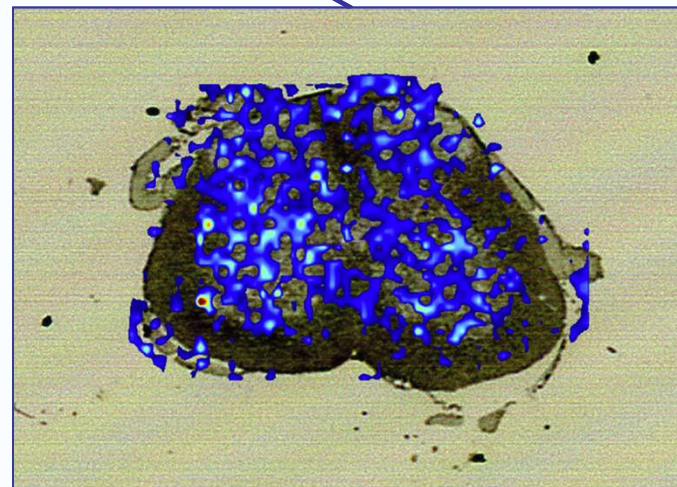
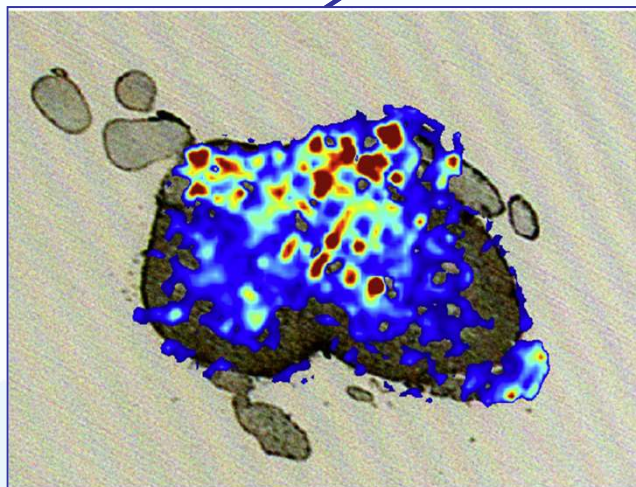
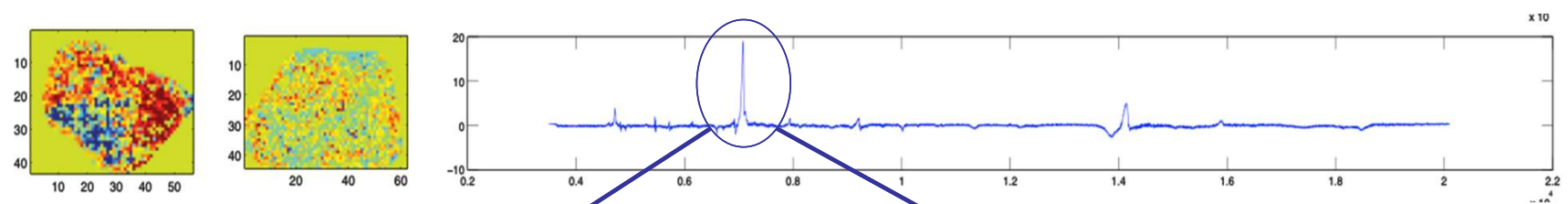


Healthy

GICA Results



Ion images





US visit opportunity

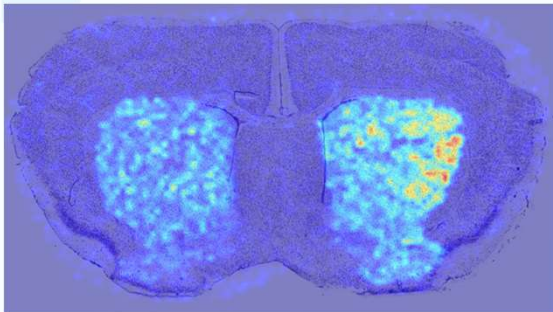


US visit opportunity

Go for it!

bdm

Protein



Bruker MALDI-TOF/TOF in
linear mode

Matrix: SA sublimated

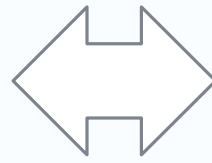
Mass range: m/z 3000 to
22000

Nr. m/z bins: 31232

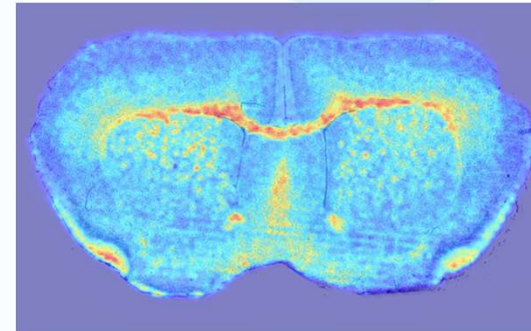
Spatial res.: 100 μ m

Spatial grid: 53 x 98

Nr. pixels: 4371



Lipid



Bruker MALDI-TOF/TOF in
linear mode

Matrix: DAN sublimated

Mass range: m/z 400 to
1000

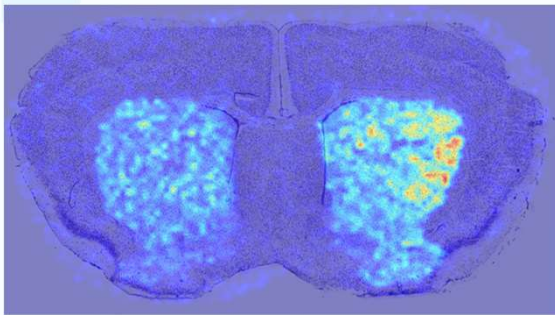
Nr. m/z bins: 26624

Spatial res.: 80 μ m

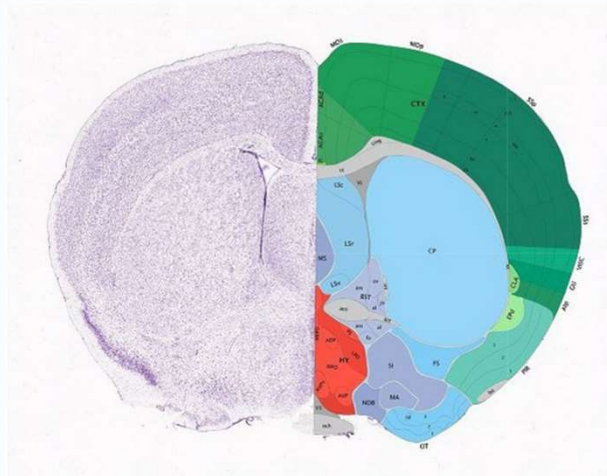
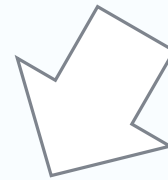
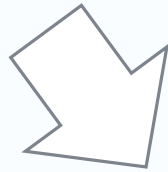
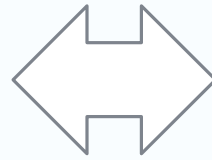
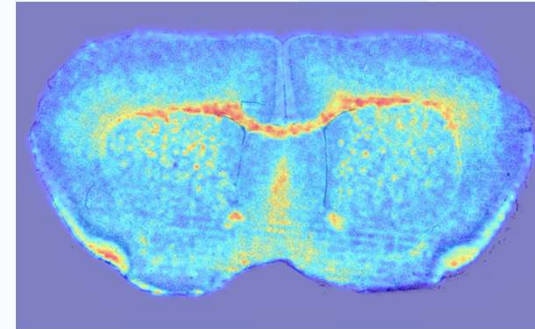
Spatial grid: 127 x 77

Nr. pixels: 7626

Protein



Lipid



atlas.brain-map.org/atlas#atlas=1&plate=100960305&structure=485&x=5712&y=3696&zoom=-4&z=3

Search for structures here

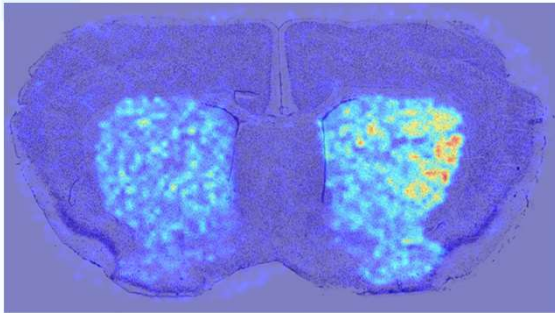
- CTXsp Cortical subplate
- CNU Cerebral nuclei
 - STR Striatum
 - STRd Striatum dorsal region**
 - STRv Striatum ventral region
 - LSX Lateral septal complex
 - LS Lateral septal nucleus
 - LSc Lateral septal nucleus, cau
 - LSr Lateral septal nucleus, rost
 - LSv Lateral septal nucleus, ver
 - SF Septofimbrial nucleus
 - SH Septohippocampal nucleus
 - sAMY Striatum-like amygdalar nuclei
 - PAL Pallidum
 - PALd Pallidum, dorsal region
 - PALv Pallidum, ventral region
 - PALm Pallidum, medial region
 - MSC Medial septal complex
 - MS Medial septal nucleus
 - NDB Diagonal band nucleus
 - TRS Triangular nucleus of septum
 - PALc Pallidum, caudal region
- CB Cerebellum
- BS Brain stem

Mouse, P56 Coronal
 Acronym: STRd
 Name: Striatum dorsal region

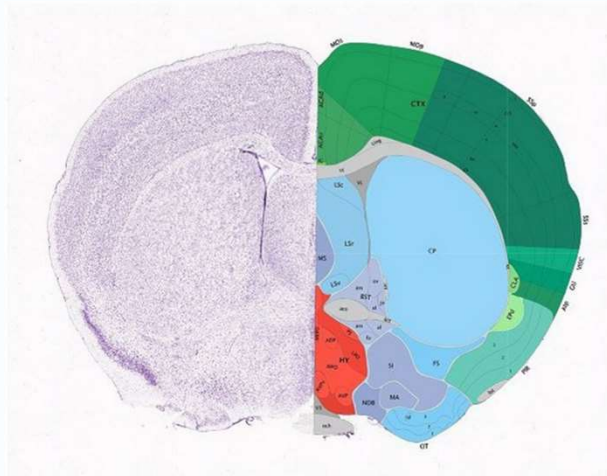
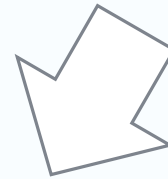
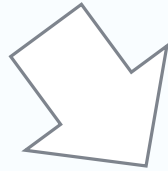
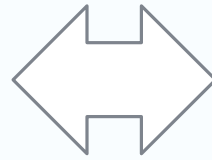
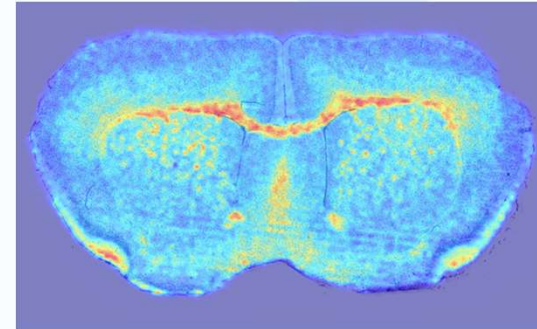
1675 microns

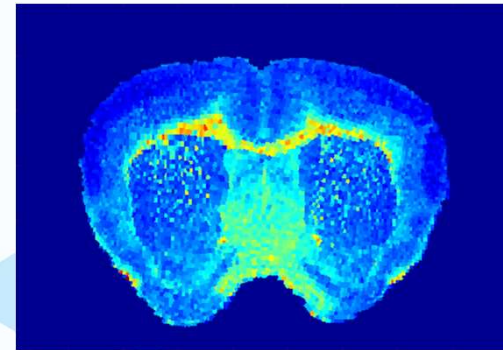
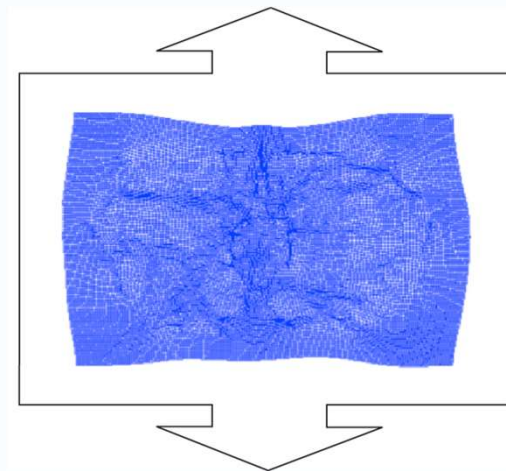
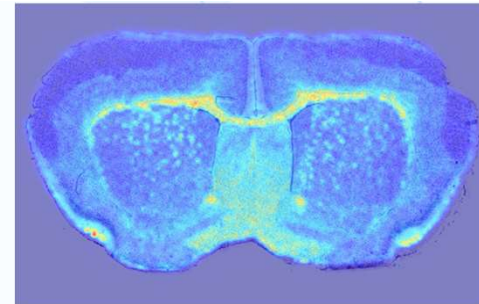
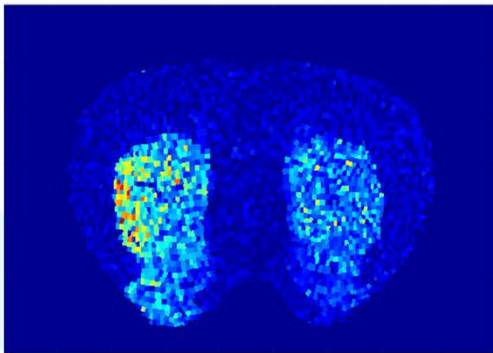
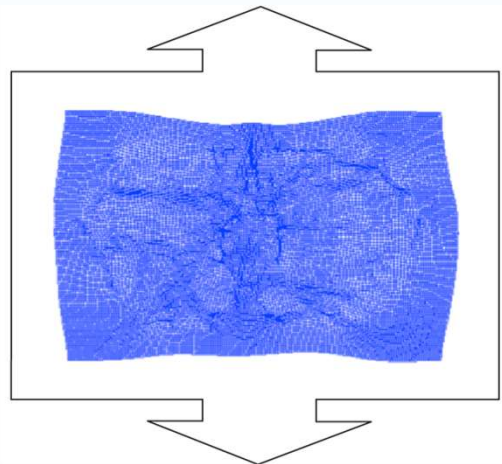
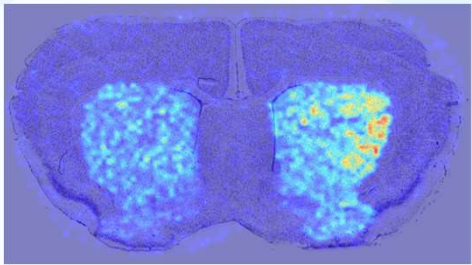
Mouse, P56 Coronal Atlas

Protein

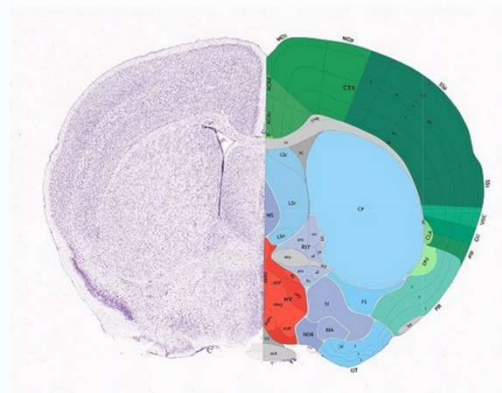


Lipid

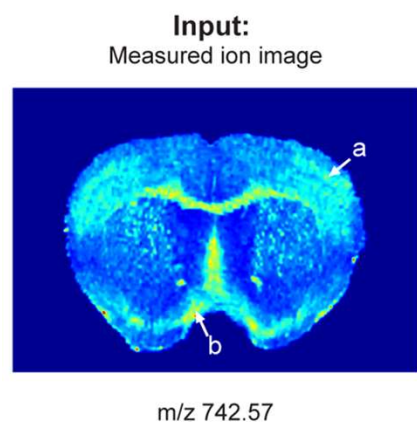




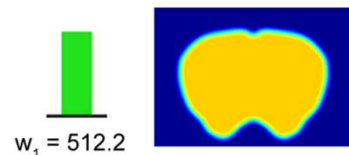
Nonrigid Registration



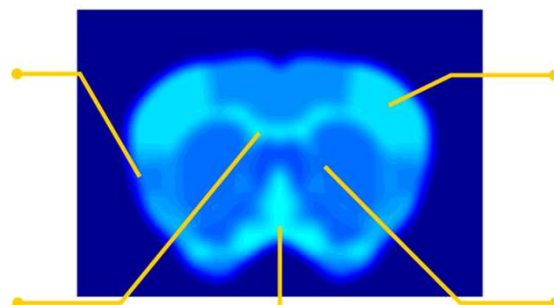
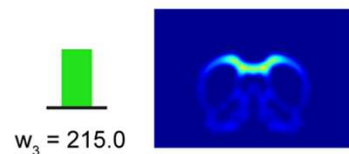
A.



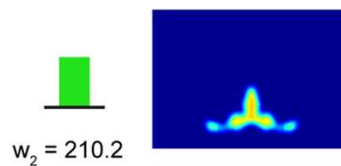
Basic cell groups and regions



Fiber tracts and ventricles



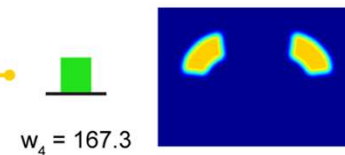
Pallidum



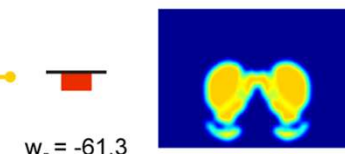
Output:

Approximation of m/z 742.57 and analysis of primary anatomical zones involved.

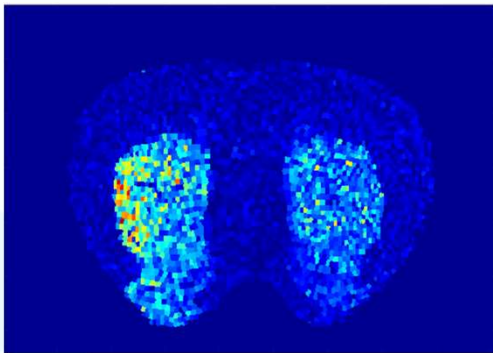
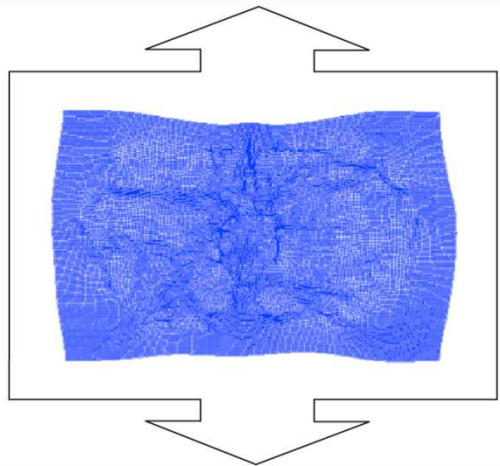
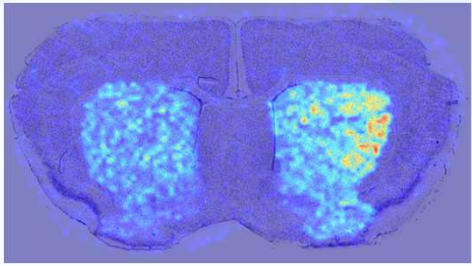
Somatosensory areas



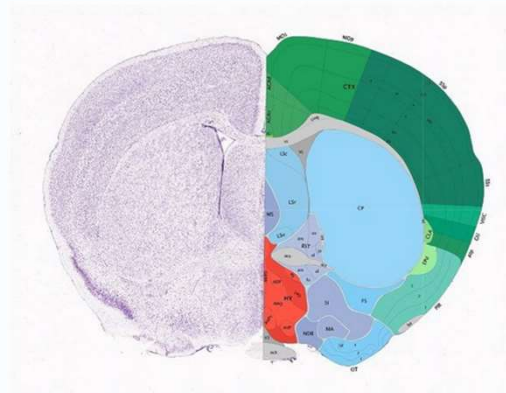
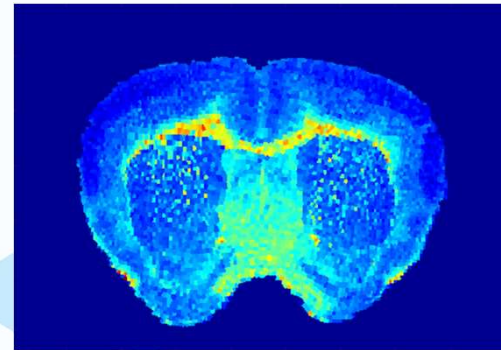
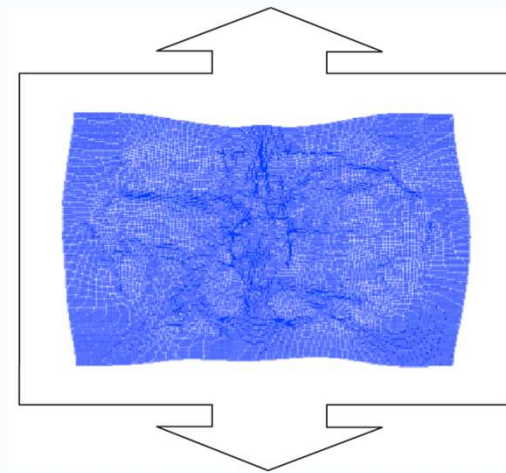
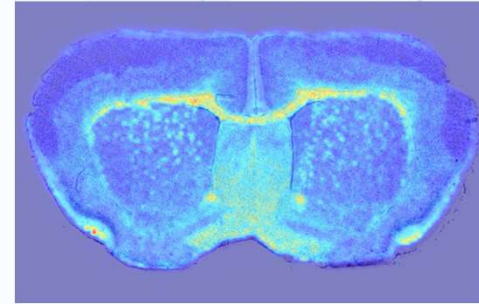
Striatum

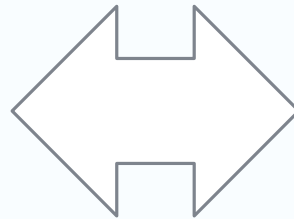
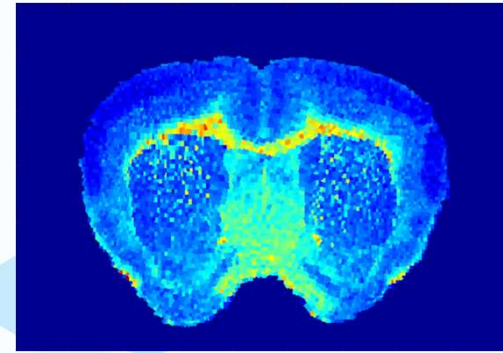
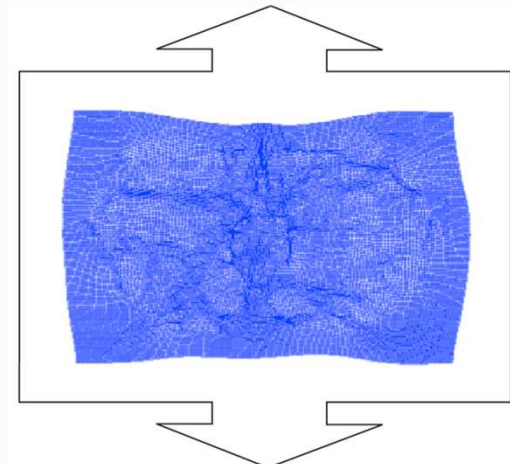
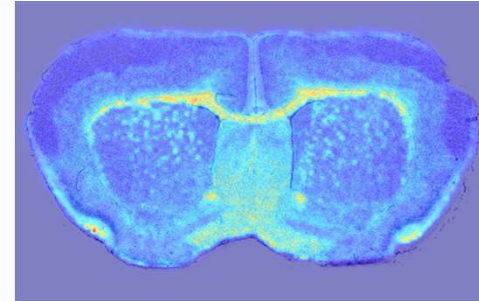
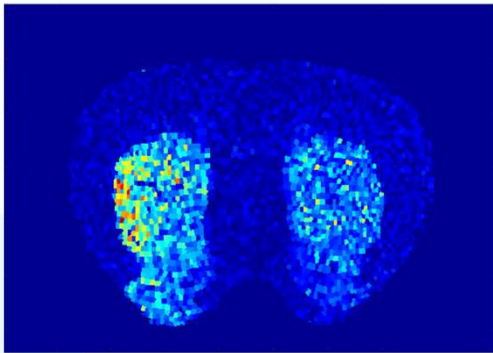
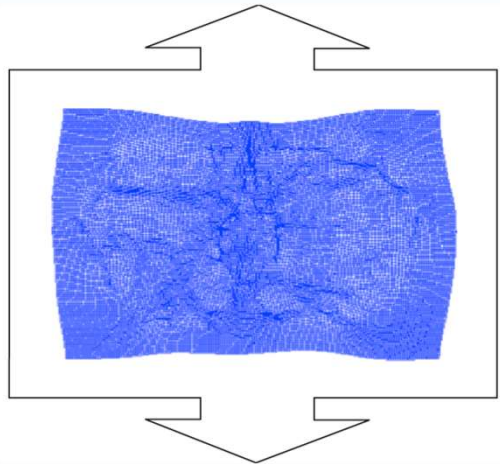
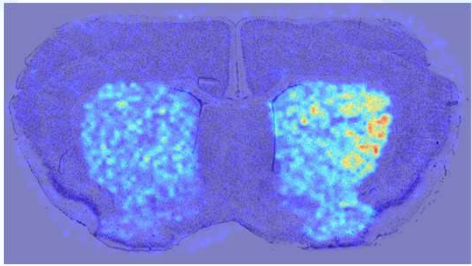


Verbeeck, N.; Yang, J.; De Moor, B.; Caprioli, R. M.; Waelkens, E.; Van de Plas, R., *Anal. Chem.* 2014.



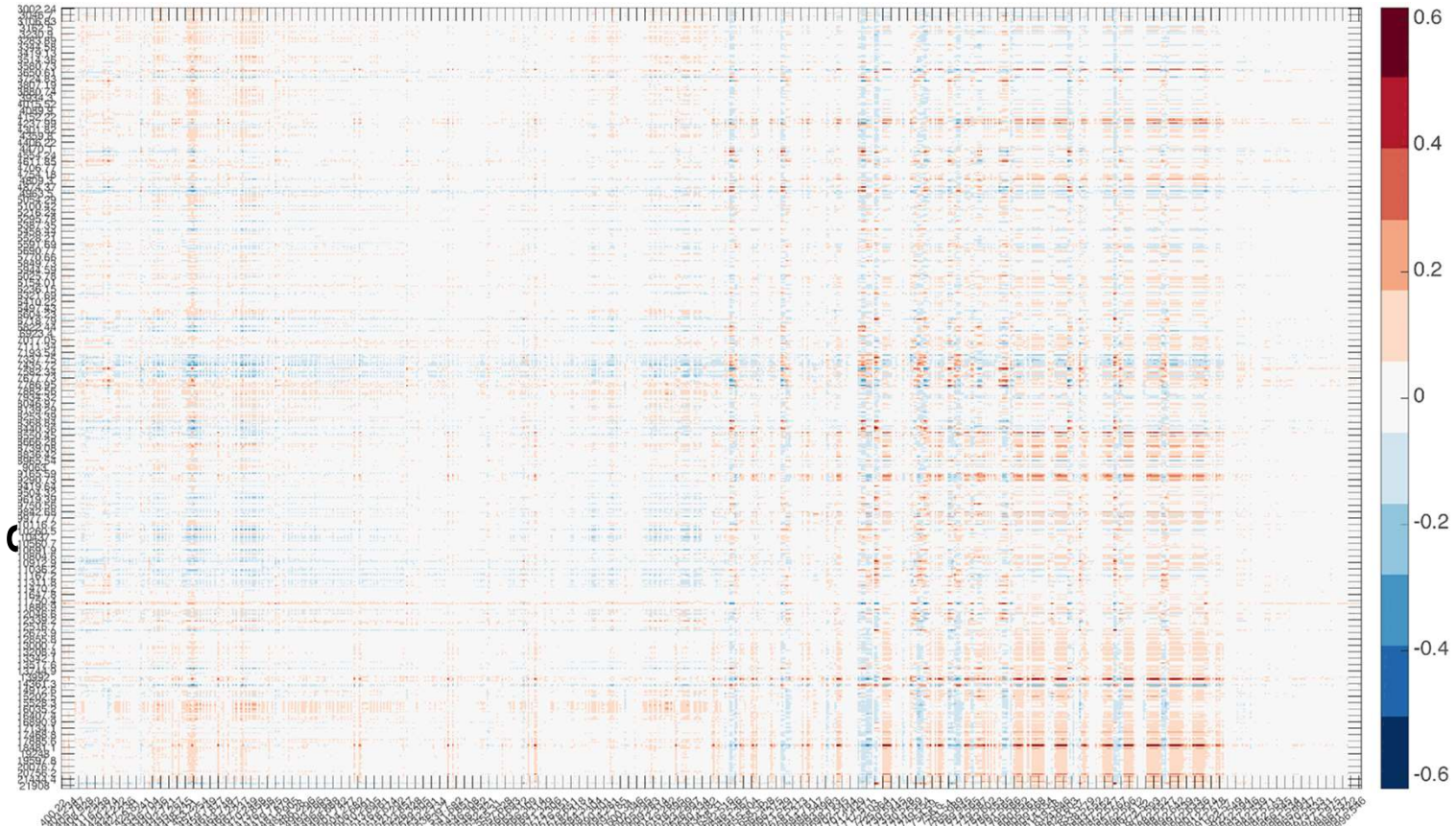
Nonrigid Registration





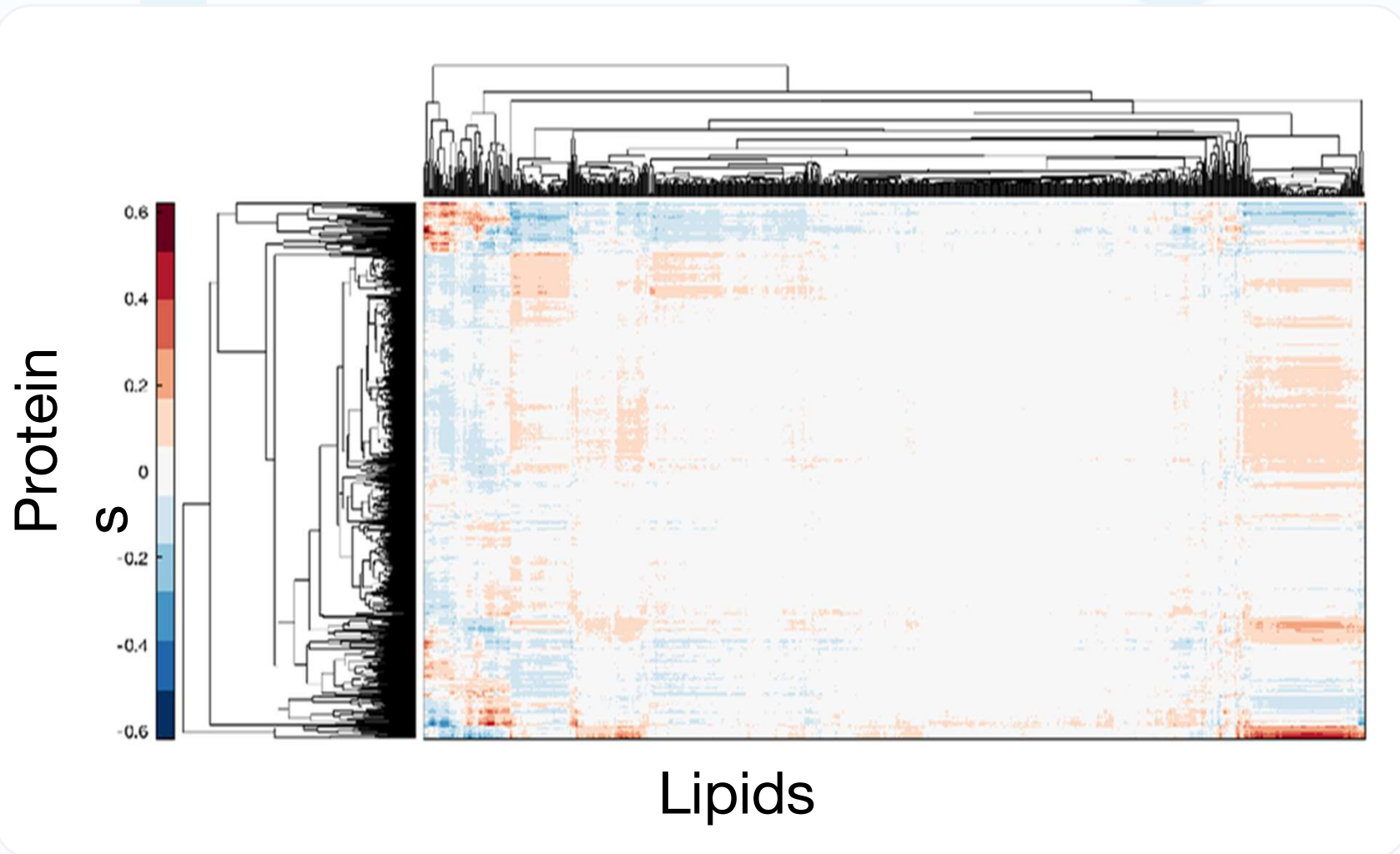
Correlation table

Protein

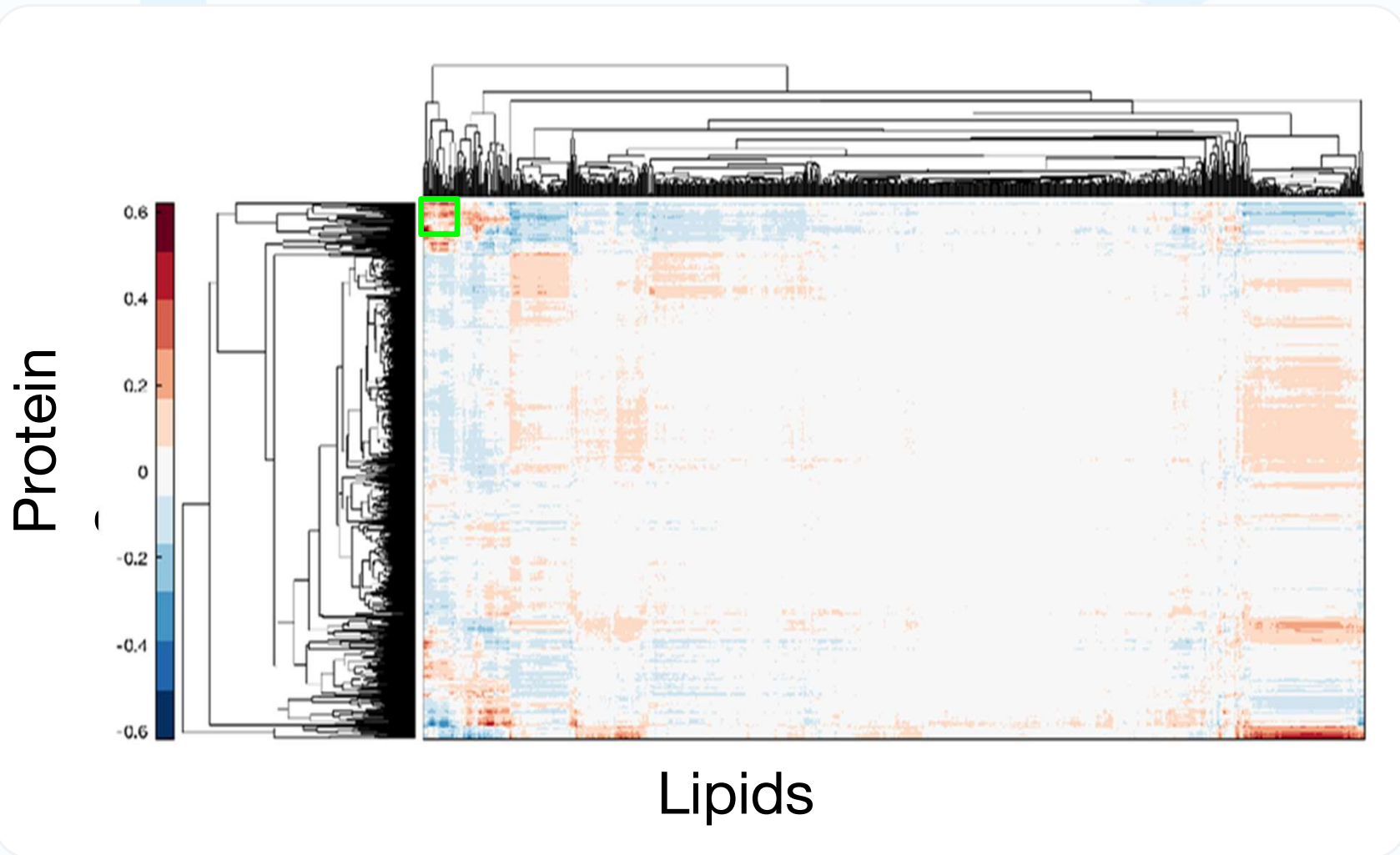


Lipids

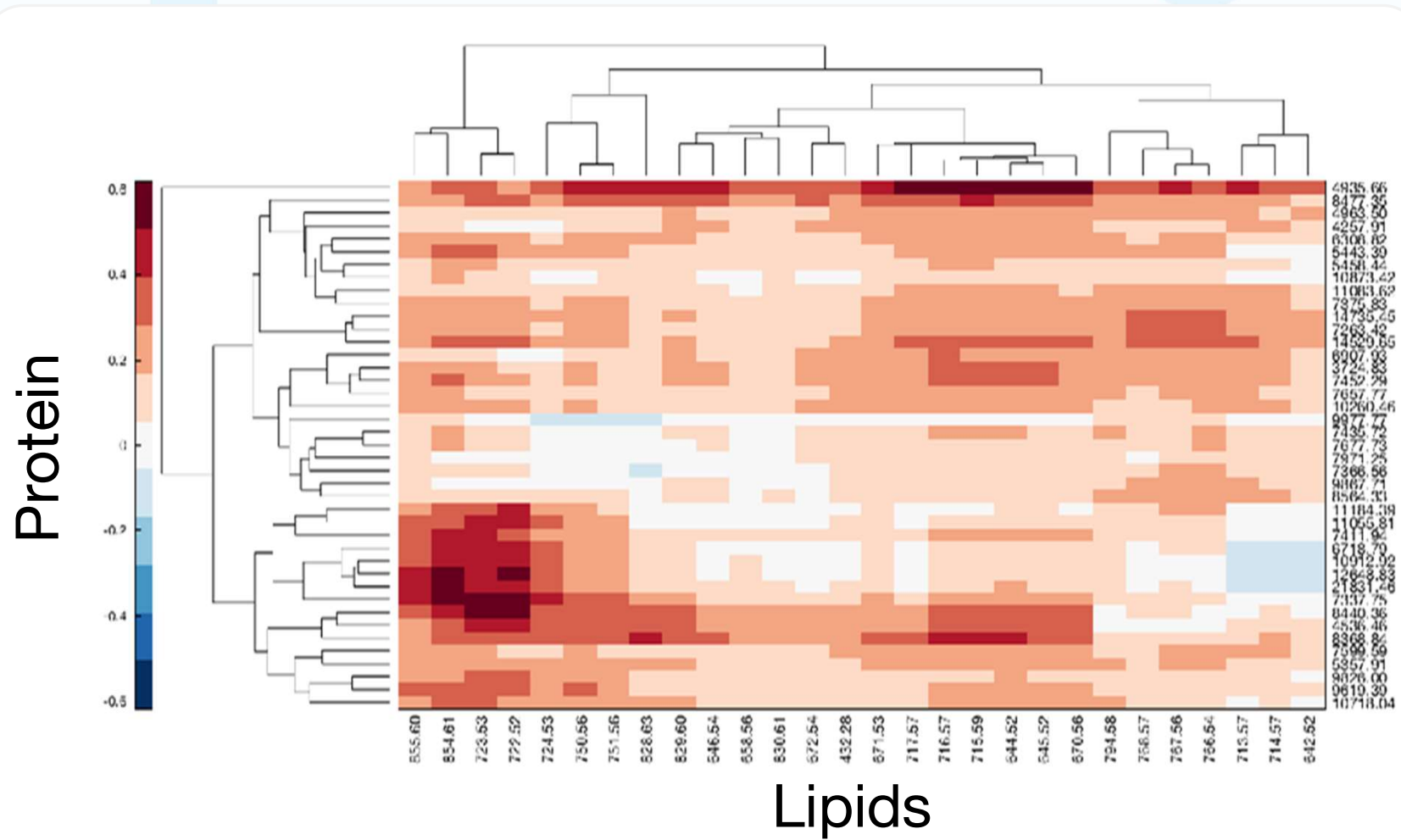
Bi-clustered correlation table



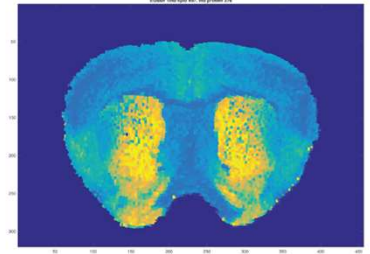
Bi-clustered correlation table



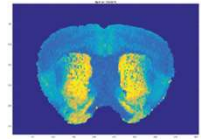
Bi-clustered correlation table (zoom-in)



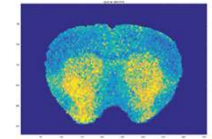
Lipids



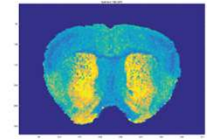
Cluster image



m/z 722.52

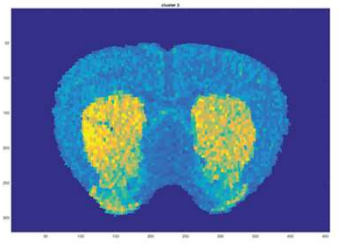


m/z 723.52

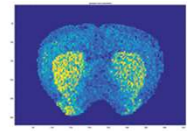


m/z 854.61

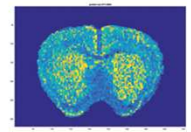
Proteins



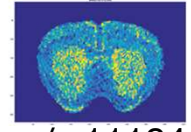
Cluster image



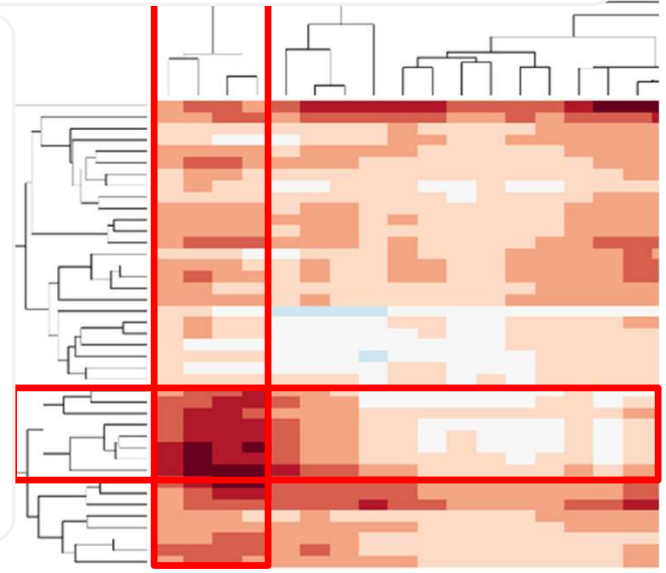
m/z 6718.8



m/z 11056



m/z 11184

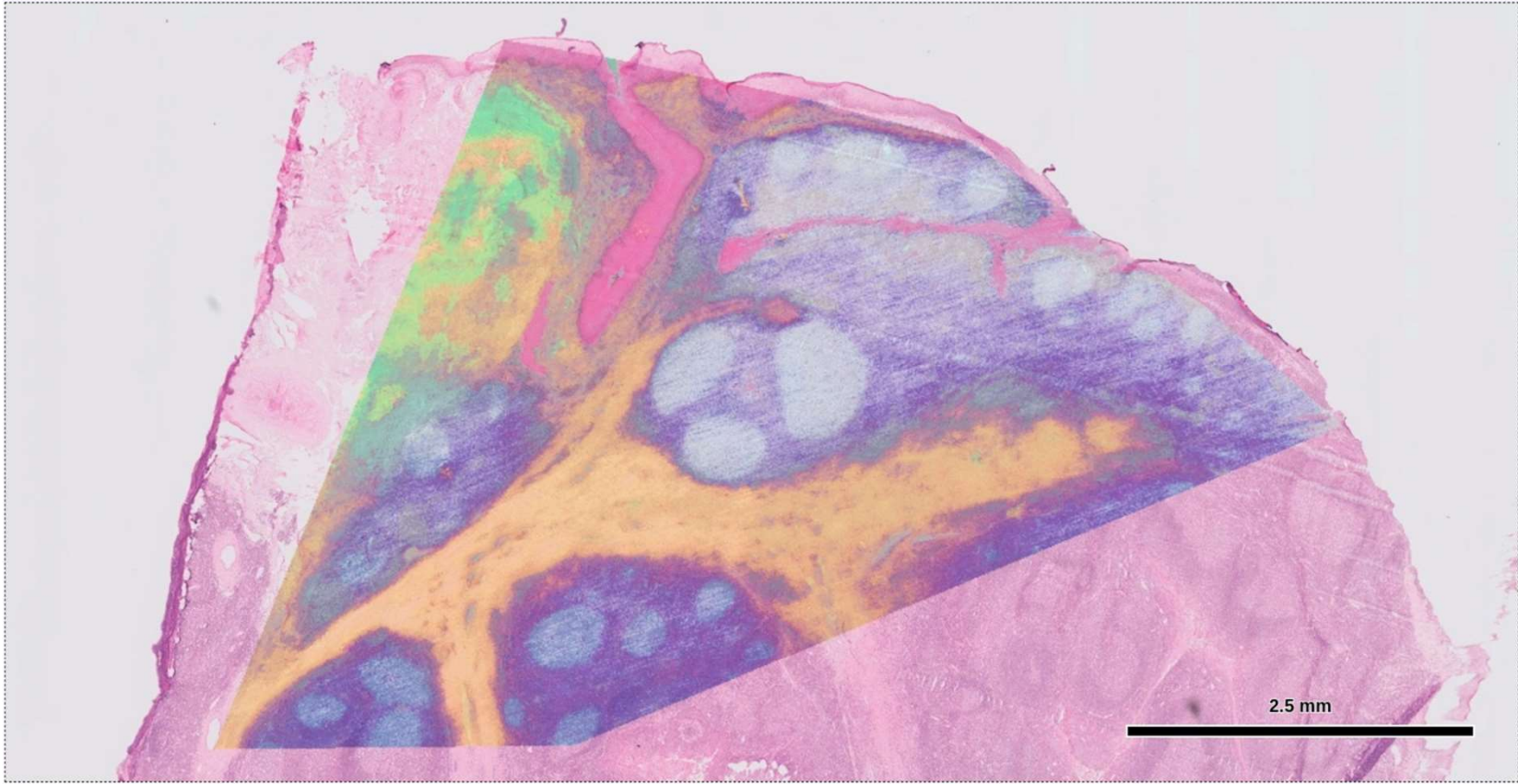




Aspect Analytics

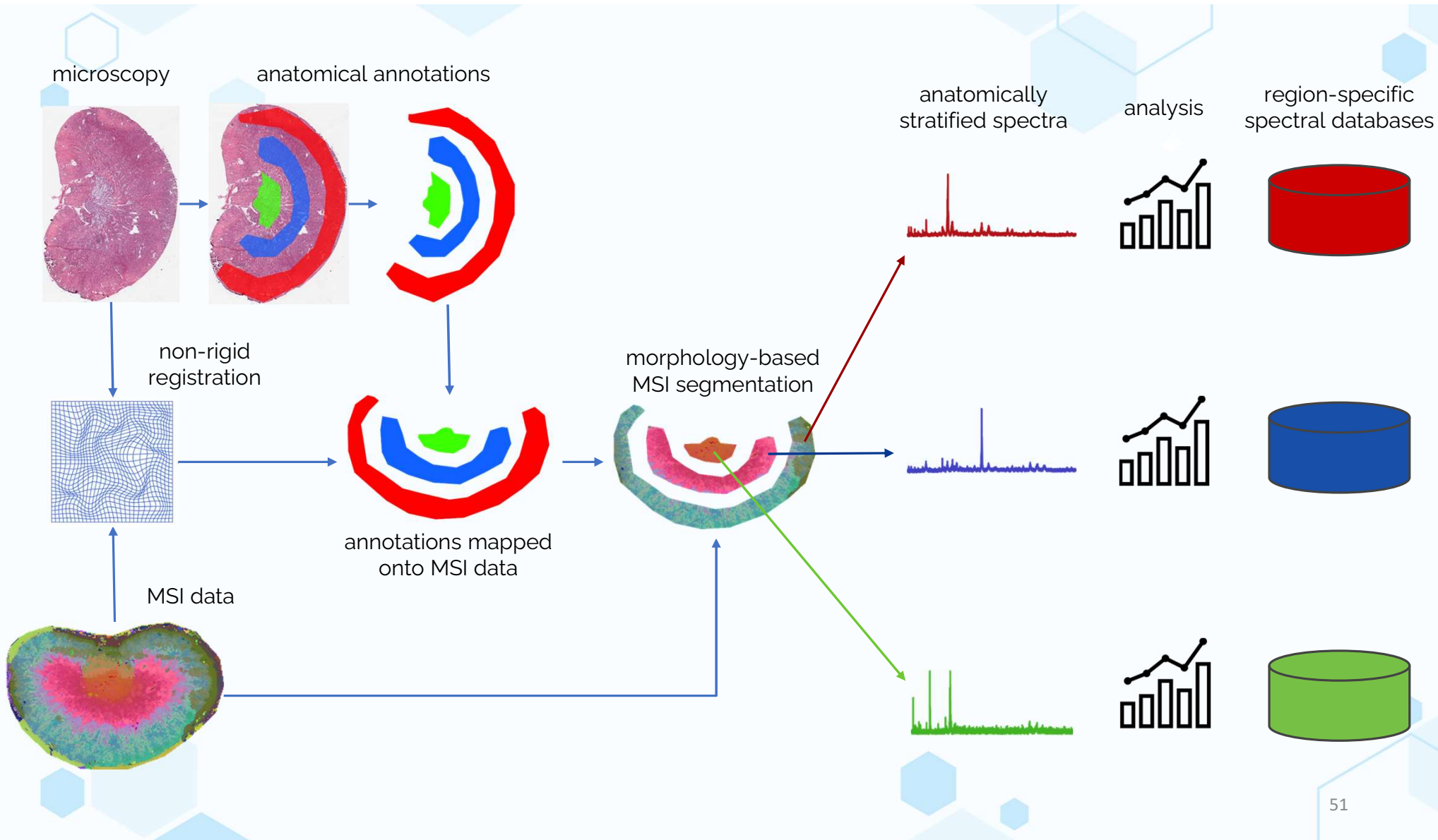


ROI selection Control point selection **Registration results** ROI to MSI Classifier training



MSI alpha
50 %





Thank you for your attention!