

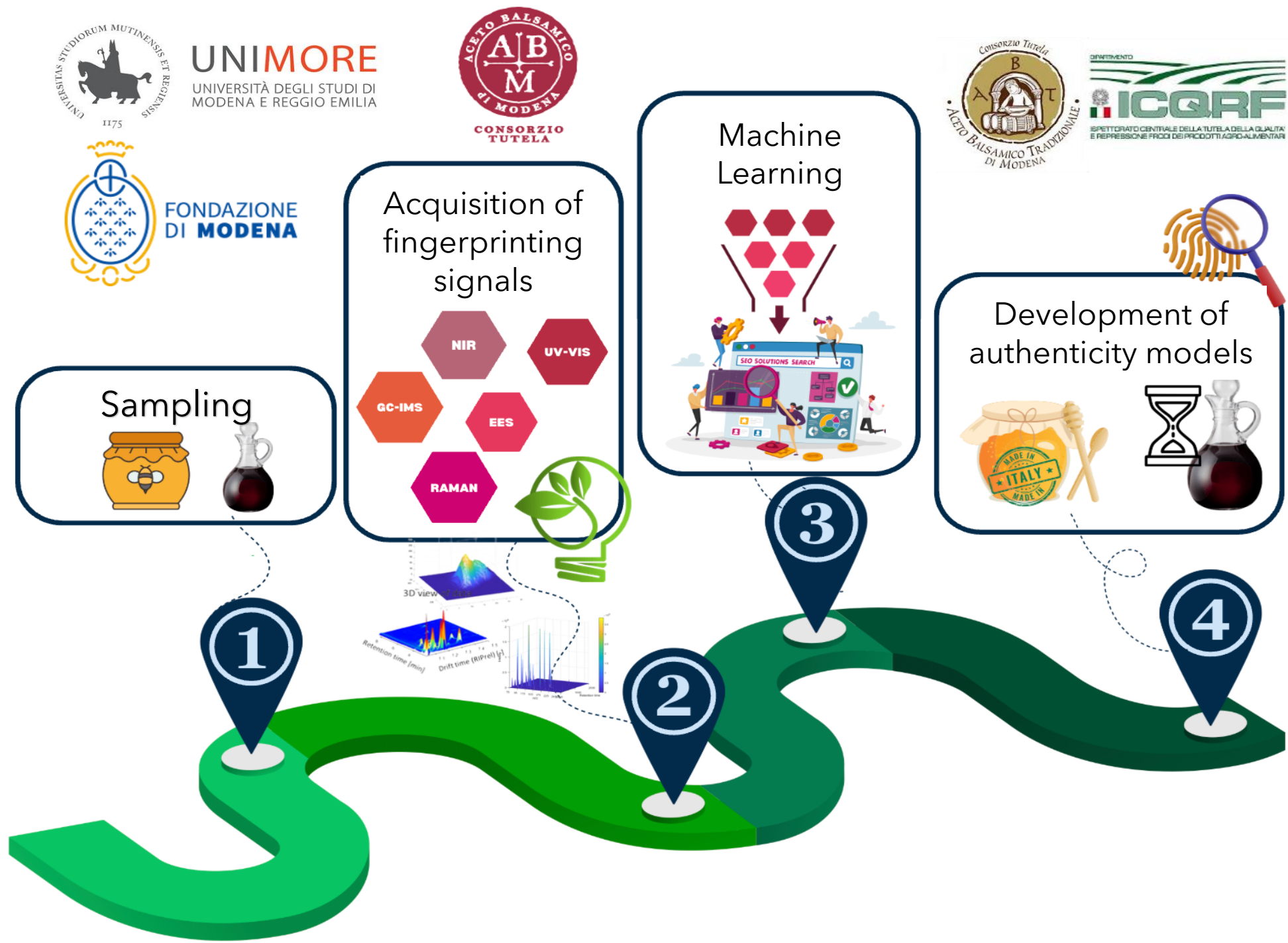
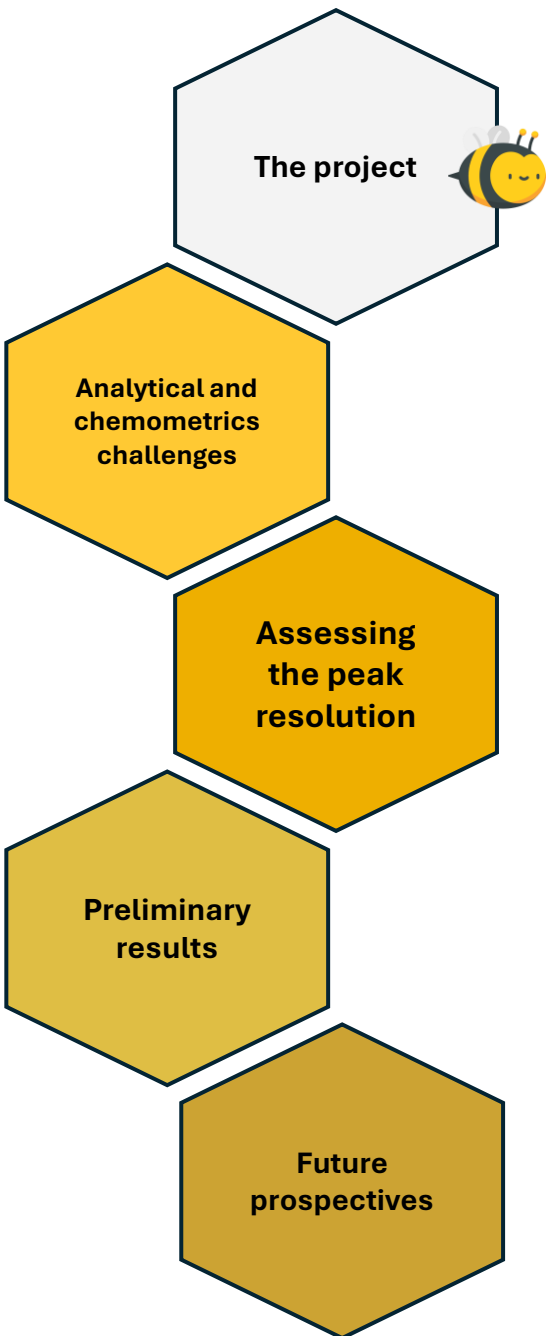
# Development of GC-IMS-based fingerprinting methods to support honey authenticity

**Samuele Pellacani**, Daniele Tanzilli, Marina Cocchi, Raniero Marabottini, Angela Napolitano, Rebecca Barbieri, Giorgia Gazzoli, Francesca Martone, Lorenzo Strani, Noemi Coppola, Caterina Durante

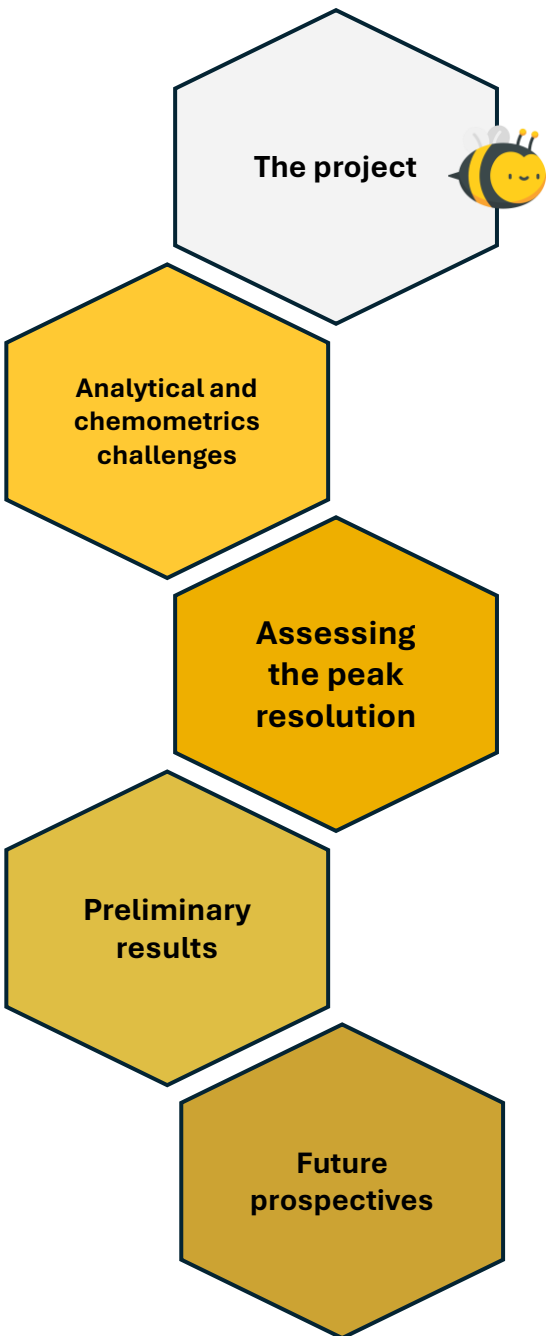


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# Sampling



**Citrus**  
17.9%

**Acacia**  
28.4%

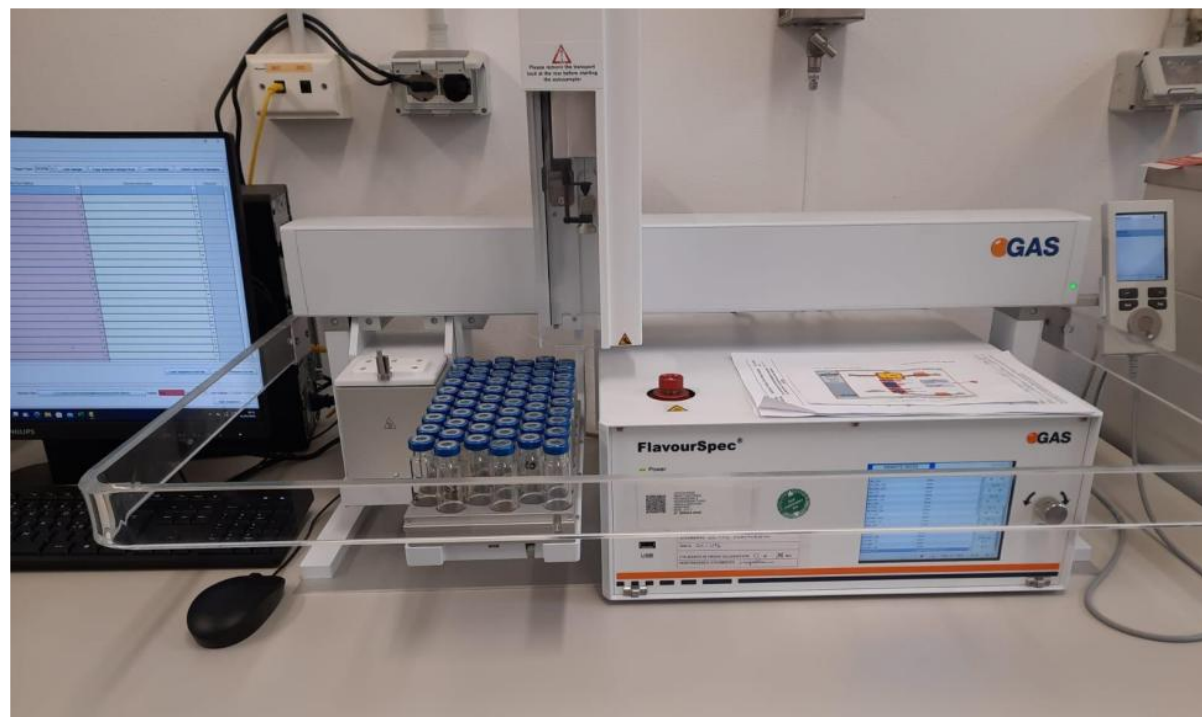
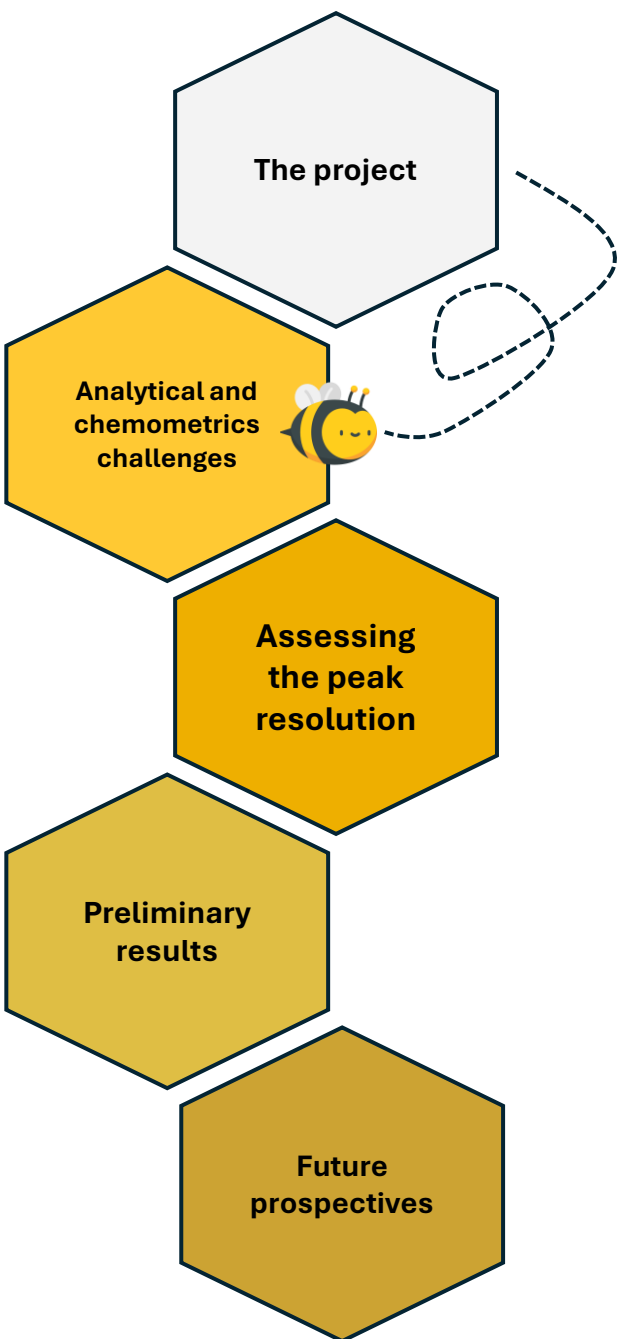
**Wildflowers**  
53.7%

**Not-italian**  
53.5%

**Italian**  
46.5%

Italian	31
Not-Italian	36

# Gas-Chromatography – Ion Mobility Spectrometry (GC-IMS)



GC-IMS FlavourSpec © (G.A.S.)



The project

Analytical and  
chemometrics  
challenges



Assessing  
the peak  
resolution

Preliminary  
results

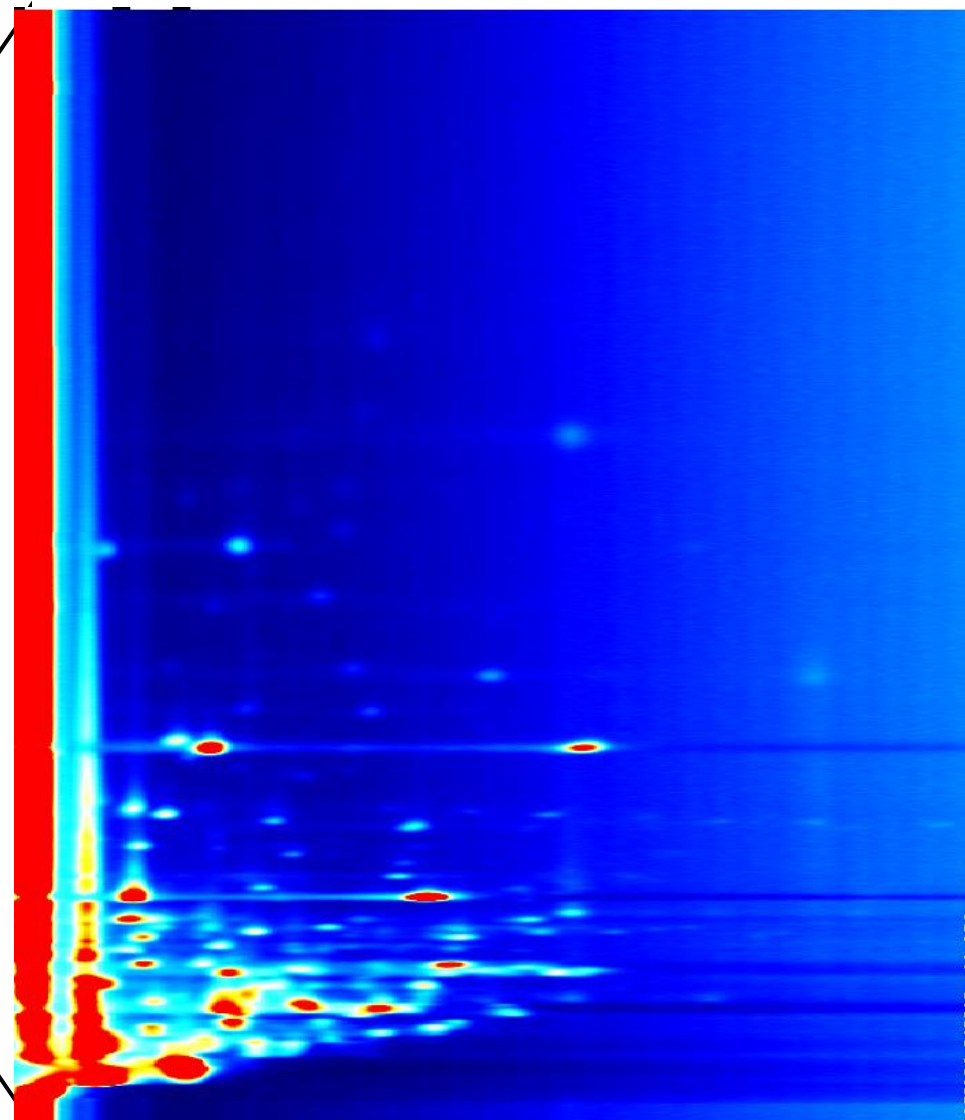
Future  
prospectives

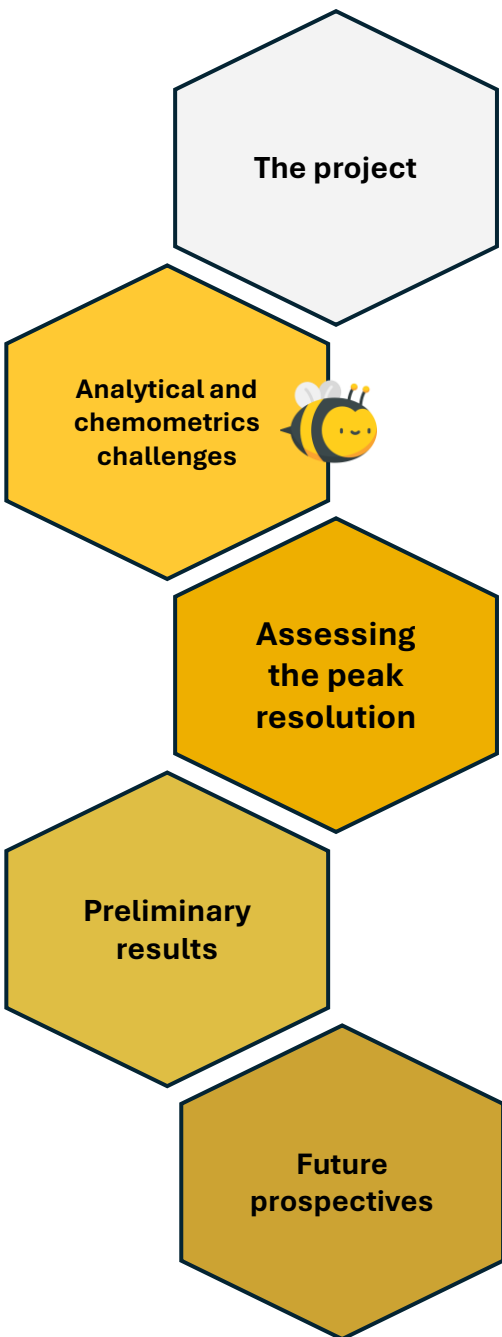


Retention time (sec)

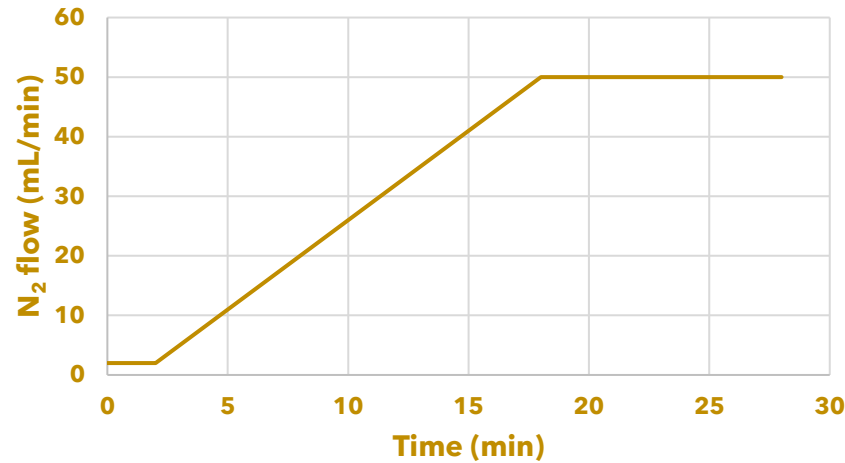
1,00  
RIP: 4.21ms

Drift time (ms)





## The instruments works at constant temperature



**Memory effect<sup>1</sup>** 



**Two blanks between samples**

**Long sequence  
(67 samples + blanks)**



**Control samples, one  
every 10 samples to look  
for instrumental drift**

<sup>1</sup>Jurado-Campos et al., "Usage considerations for headspace-gas chromatography-ion mobility spectrometry as a suitable technique for qualitative analysis in a routine lab" *Journal of Chromatography A*, **2021**, 1640, 461937.

# *iCoshift* alignment

The project

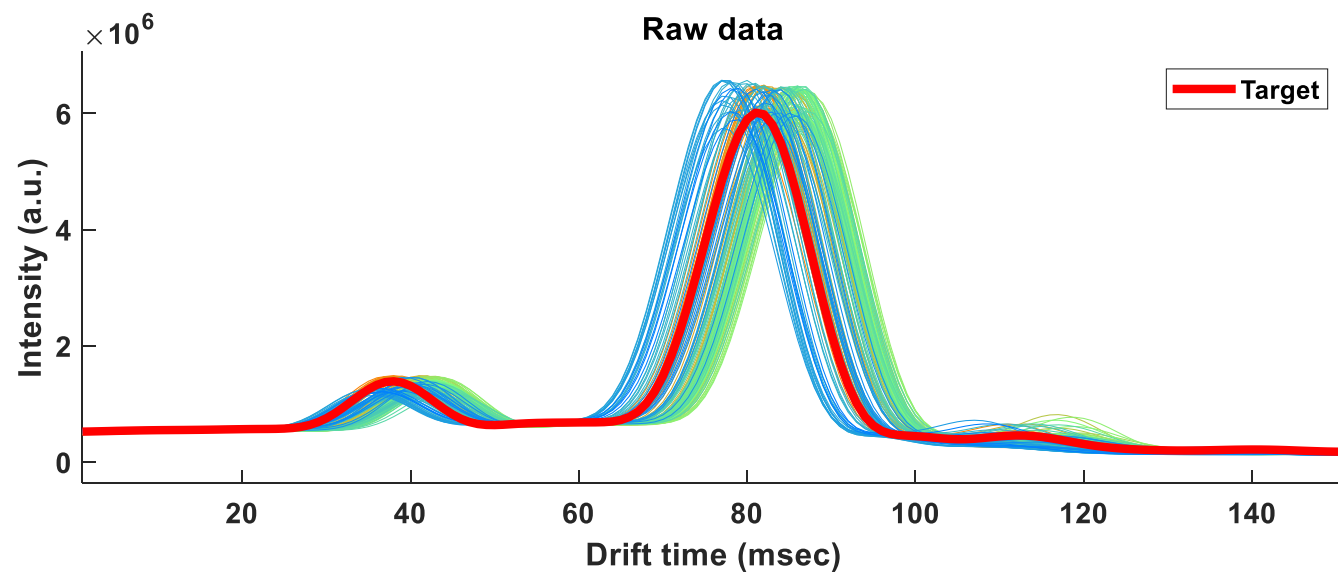
Analytical and  
chemometrics  
challenges



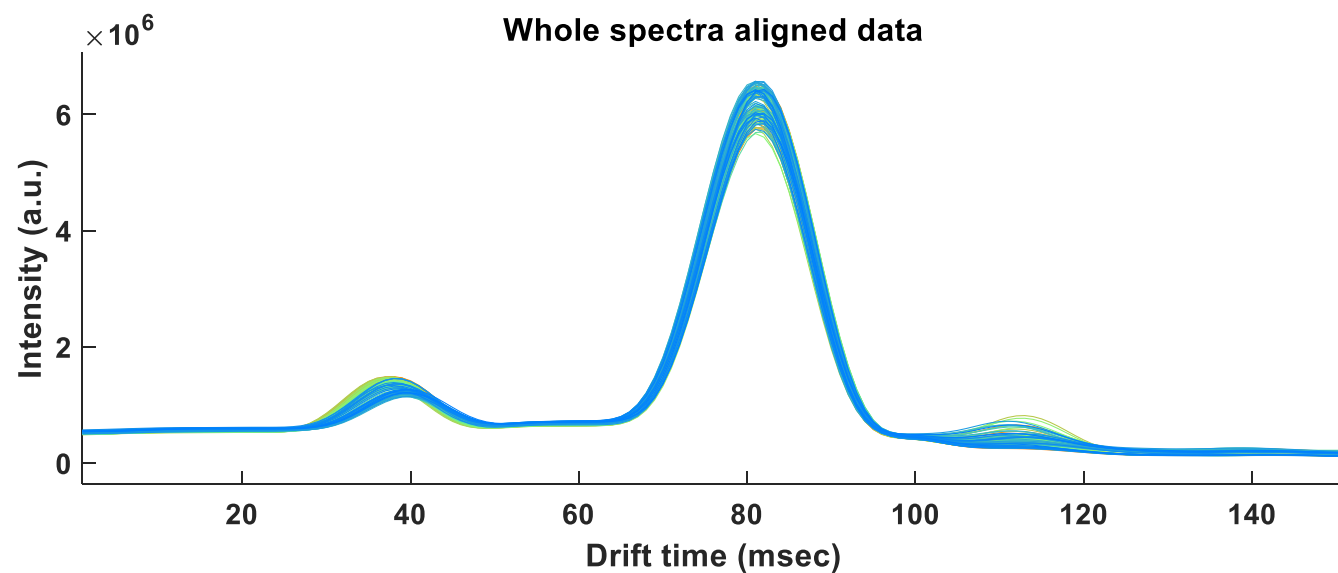
Assessing  
the peak  
resolution

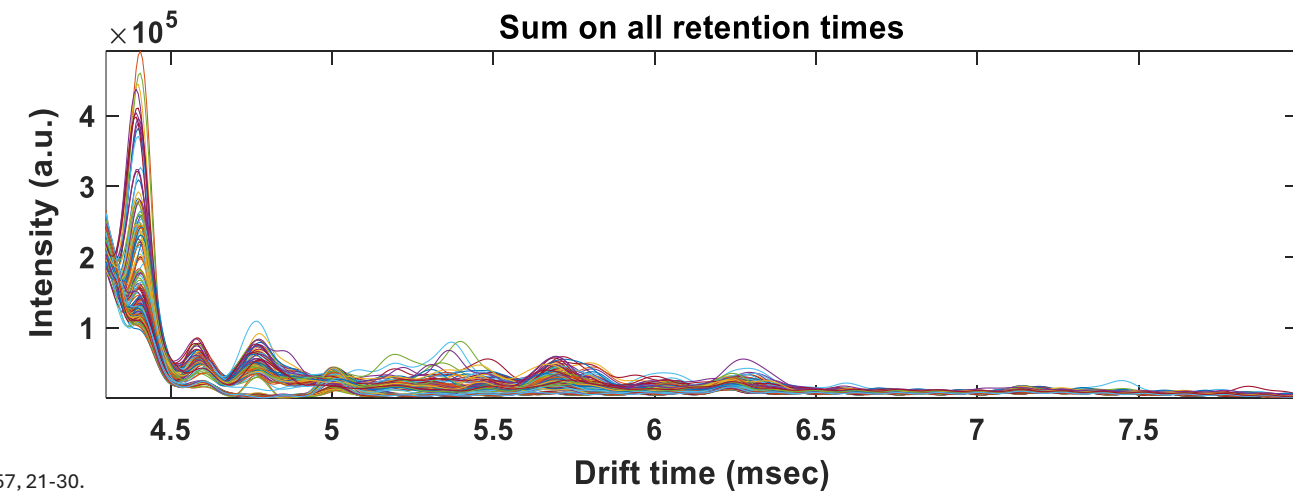
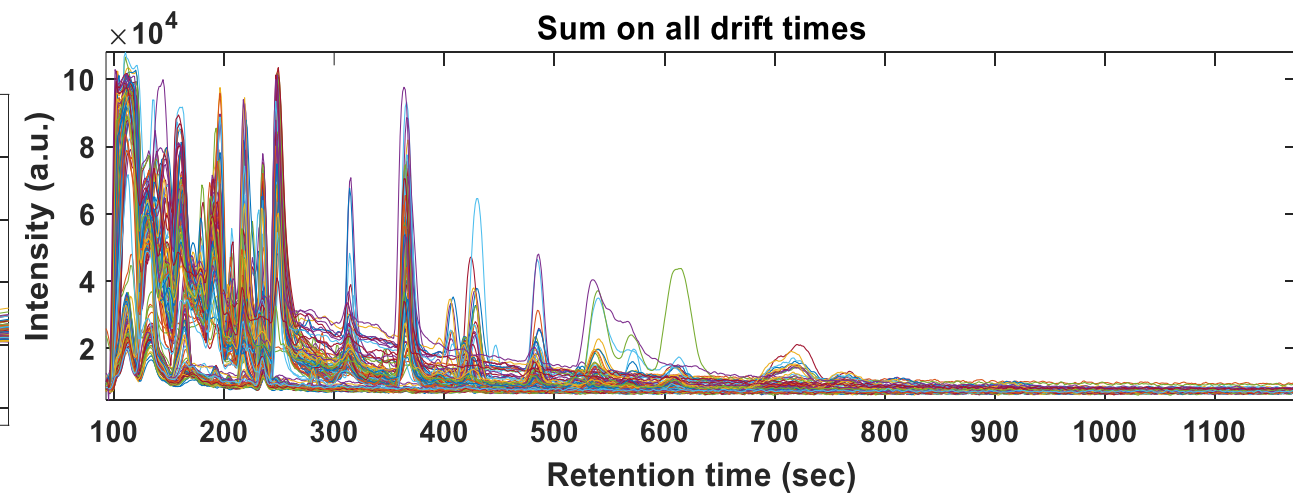
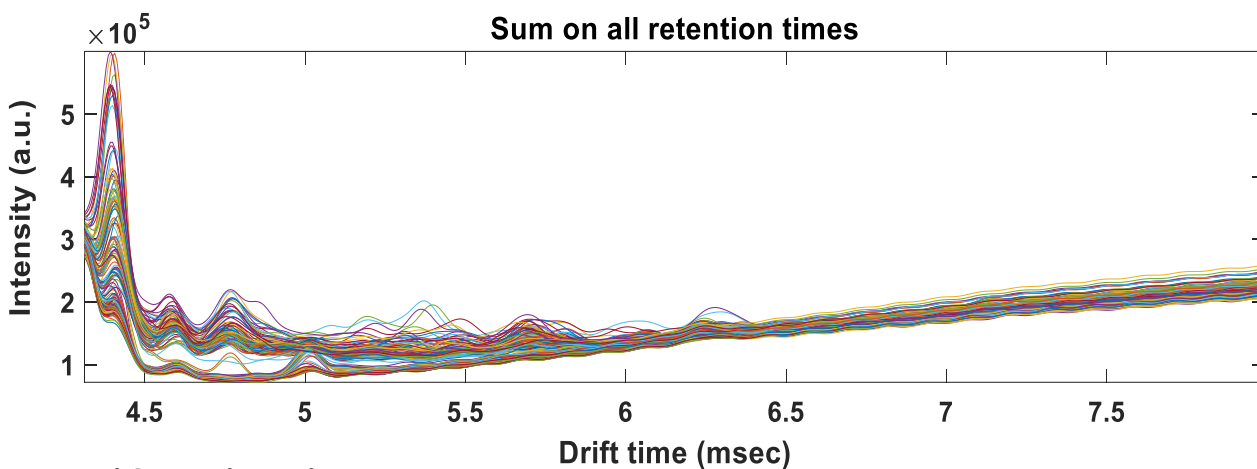
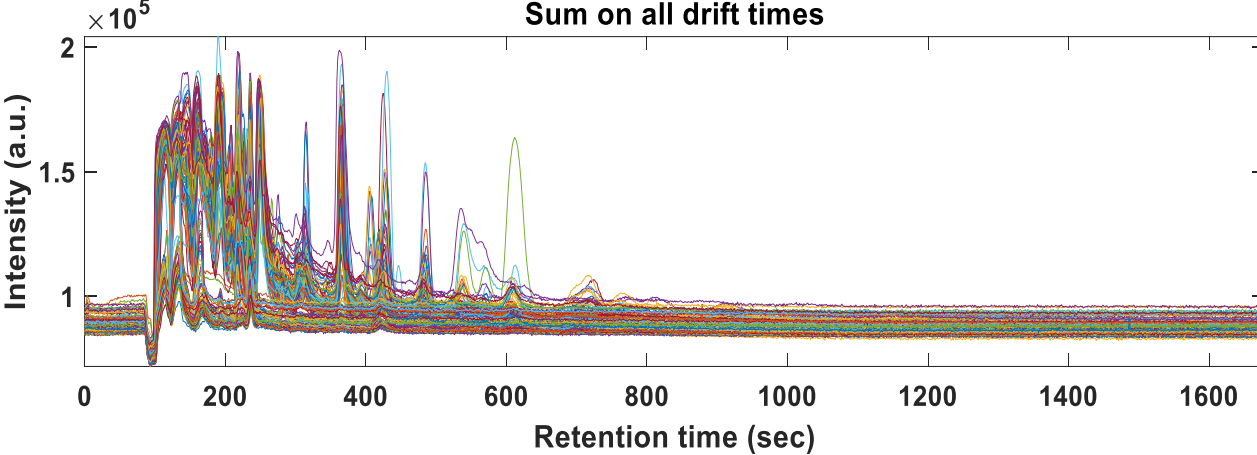
Preliminary  
results

Future  
prospectives

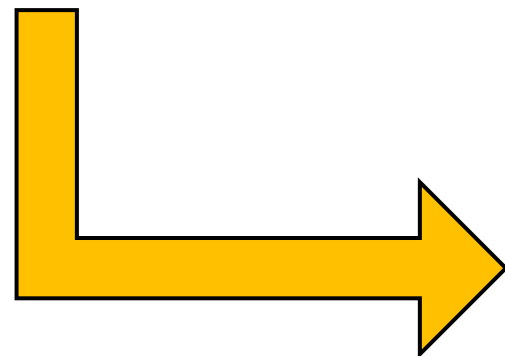


Alignment  
applied to the  
rest of the  
spectra





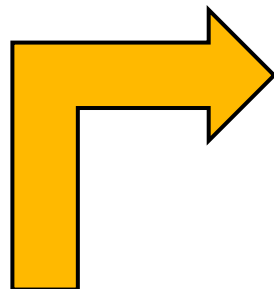
- ***iCoshift* alignment**
- **ROI selection**
- **Smoothing on retention times (SavGol, width = 11, order = 2)**
- **Smoothing and baseline correction on drift times (asymmetric least squares, default parameters<sup>2</sup>)**





The project

## Definition of 38 intervals



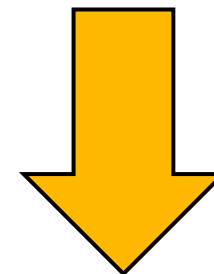
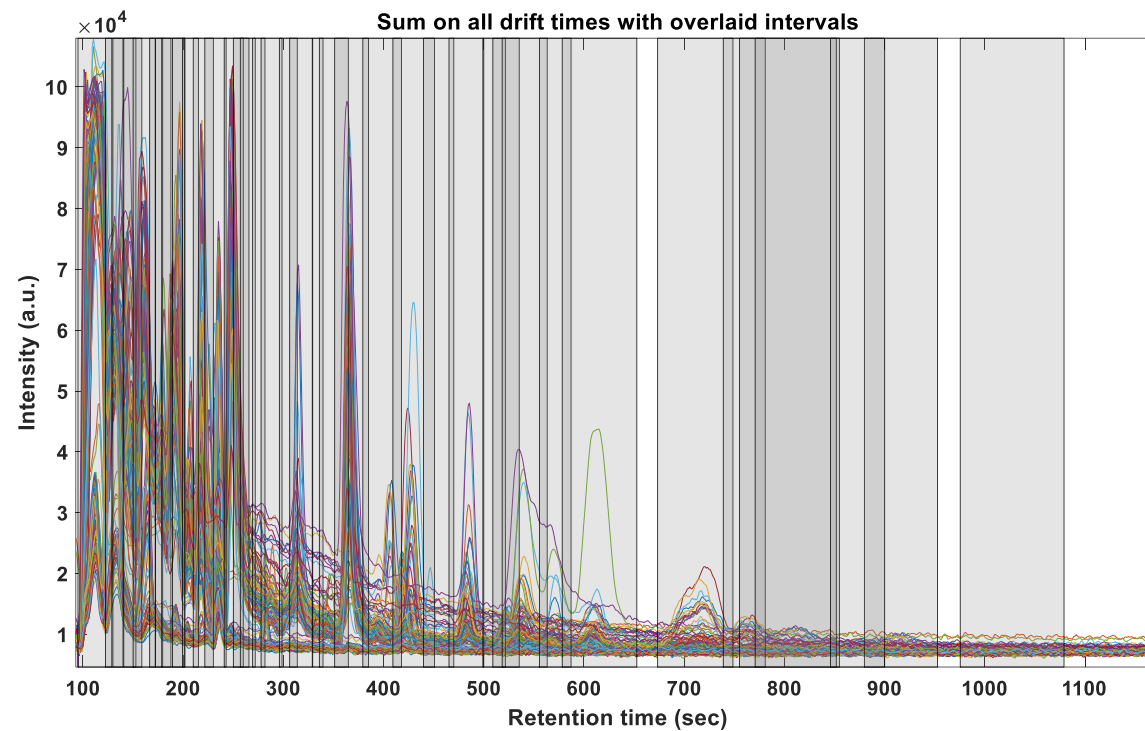
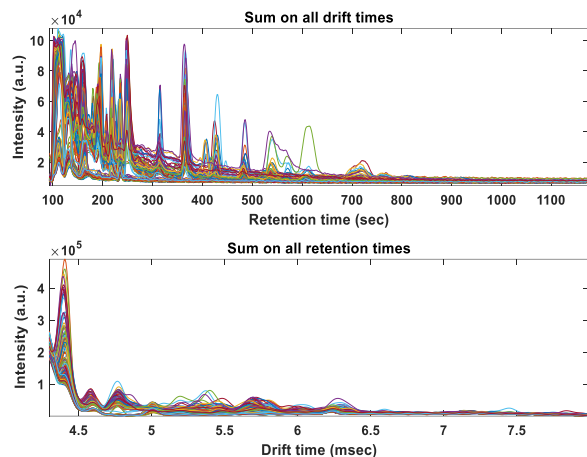
Analytical and  
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challenges



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resolution

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prospectives



**Peak deconvolution:**  
**MCR-ALS<sup>3</sup> (nonnegativity and unimodality constraints)**  
**PARAFAC2 (nonnegativity constraint)**

The project

Analytical and  
chemometrics  
challenges

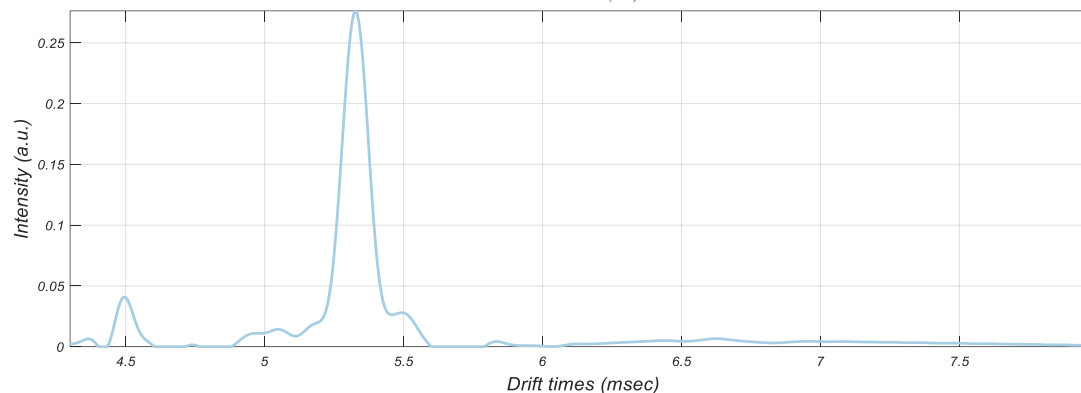
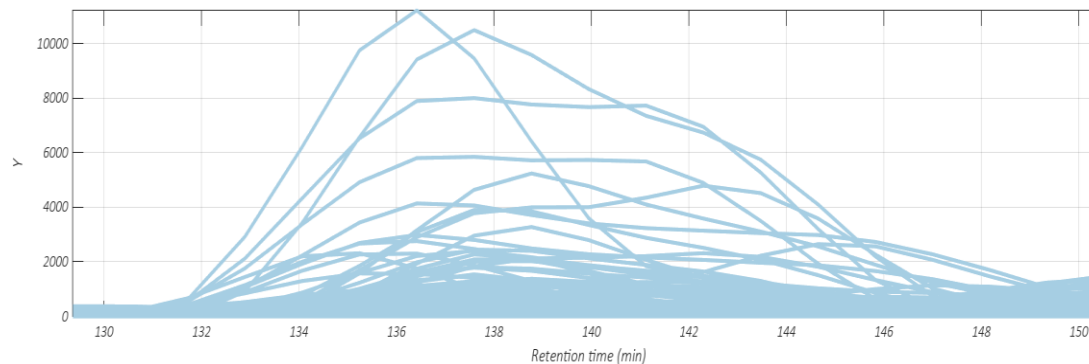
Assessing  
the peak  
resolution



Preliminary  
results

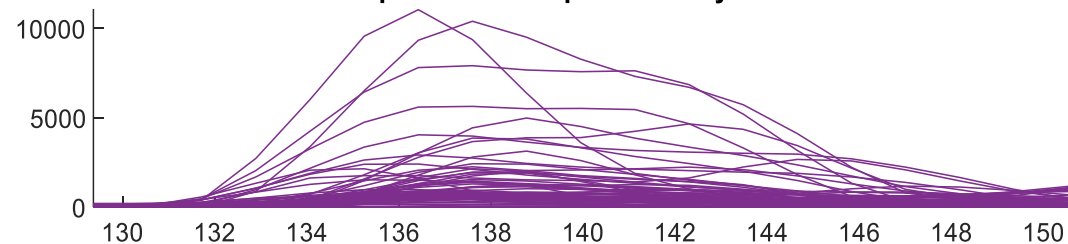
Future  
prospectives

## PARAFAC2 deconvolution

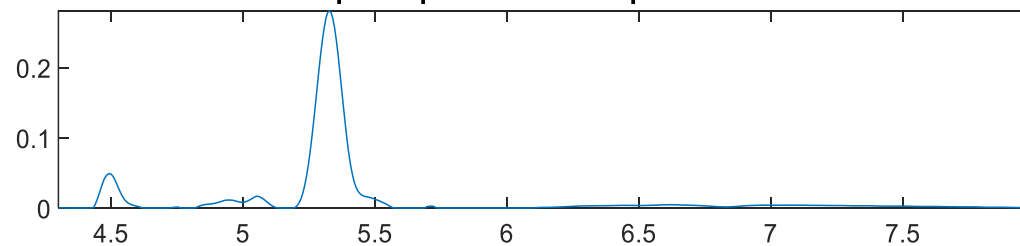


## MCR-ALS deconvolution

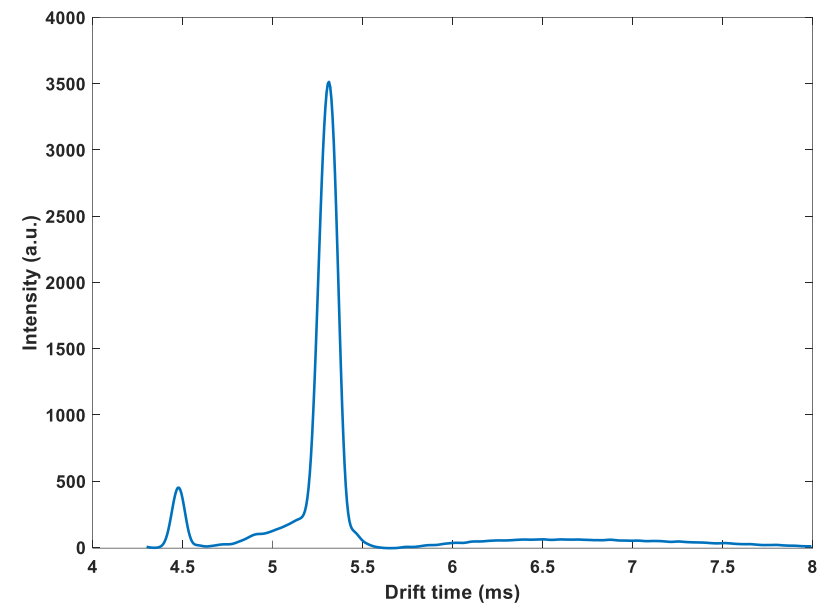
elution profile of component: 4 by time in sec

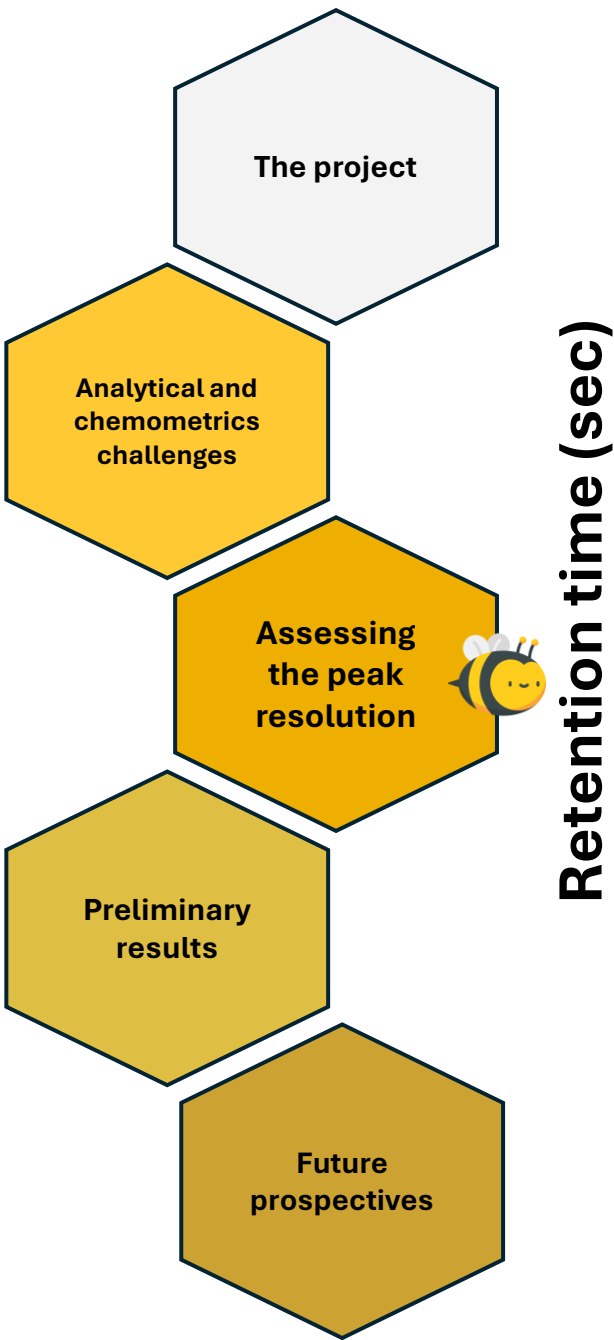


pure spectrum of component: 4

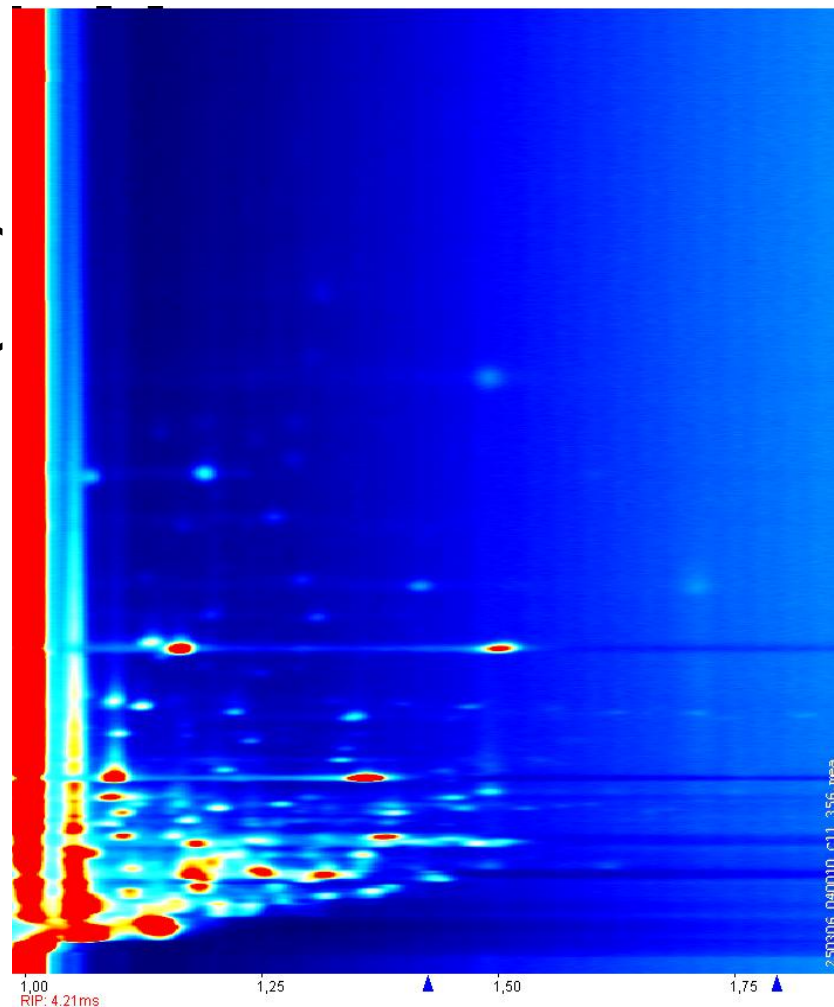


## Standard: 1-butanone





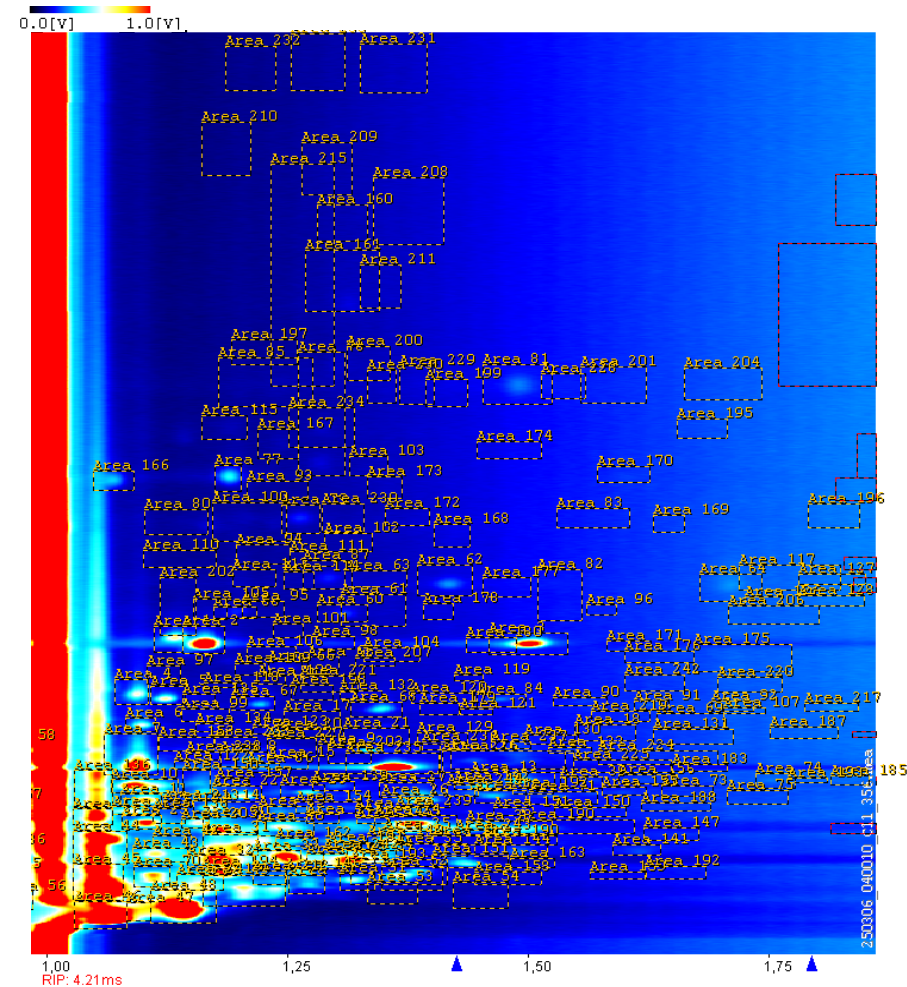
Retention time (sec)



Drift time (ms)

## Vocal software

Retention time (sec)

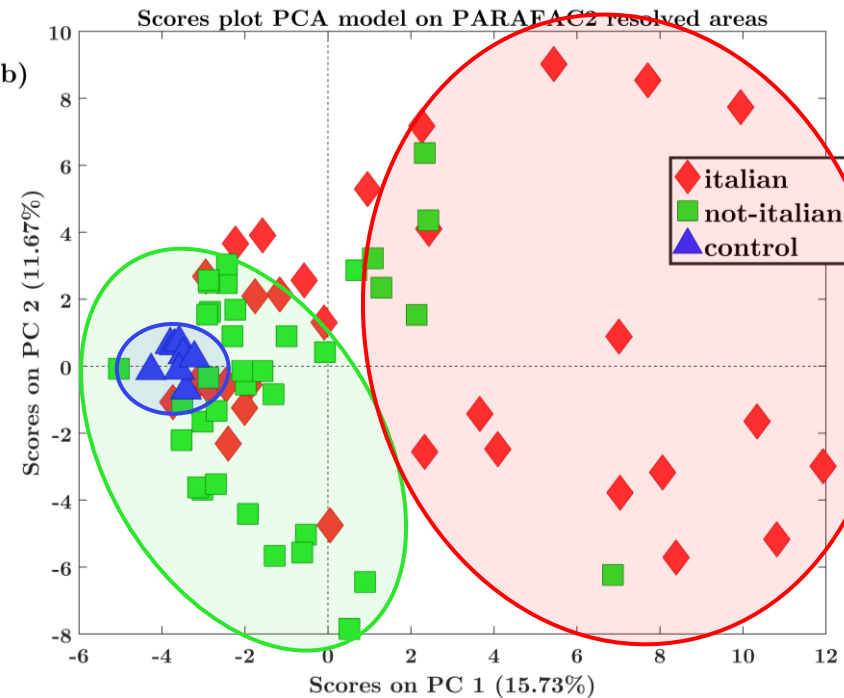


Drift time (ms)

244 intervals!!

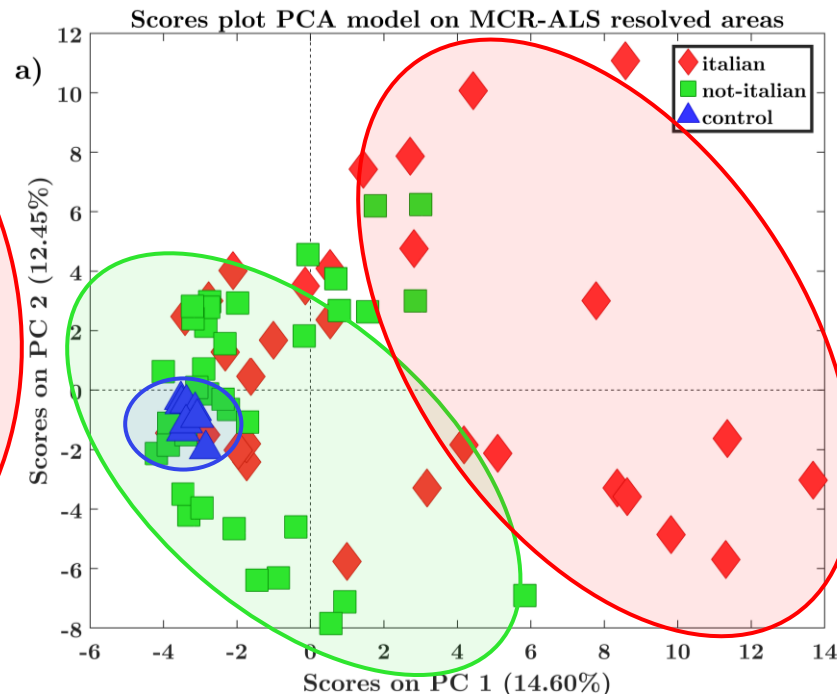
# Principal Component Analysis (PCA)

## PARAFAC2 resolved areas



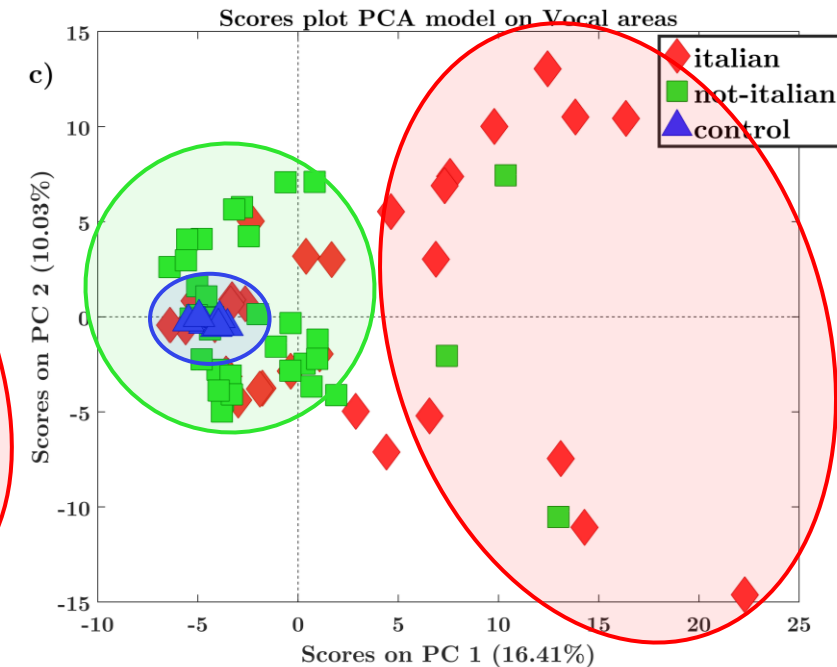
Area matrix (75 x 108)

## MCR resolved areas



Area matrix (75 x 114)

## Vocal areas



Area matrix (75 x 244)

The project

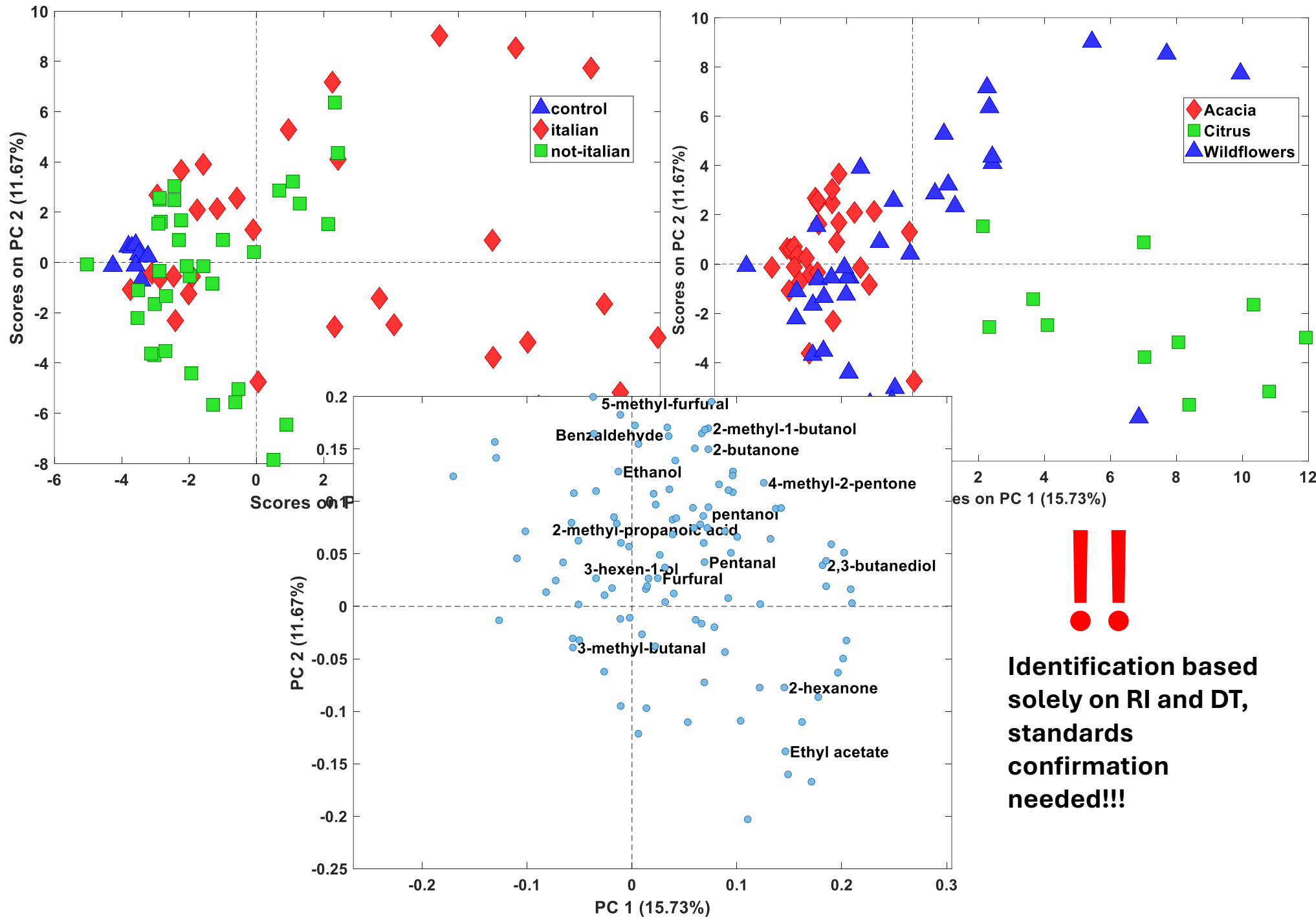
Analytical and chemometrics challenges

Assessing the peak resolution

Preliminary results

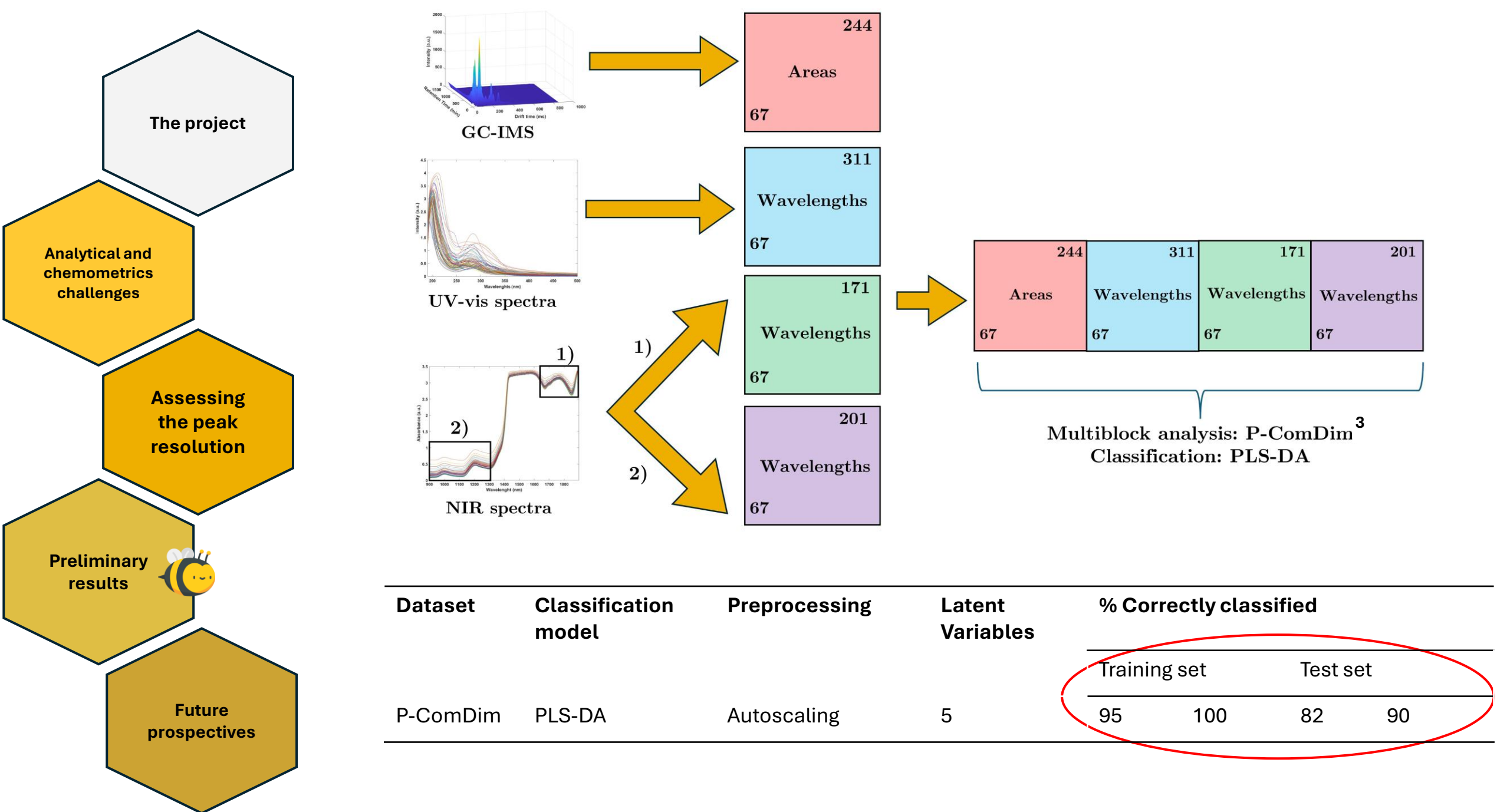


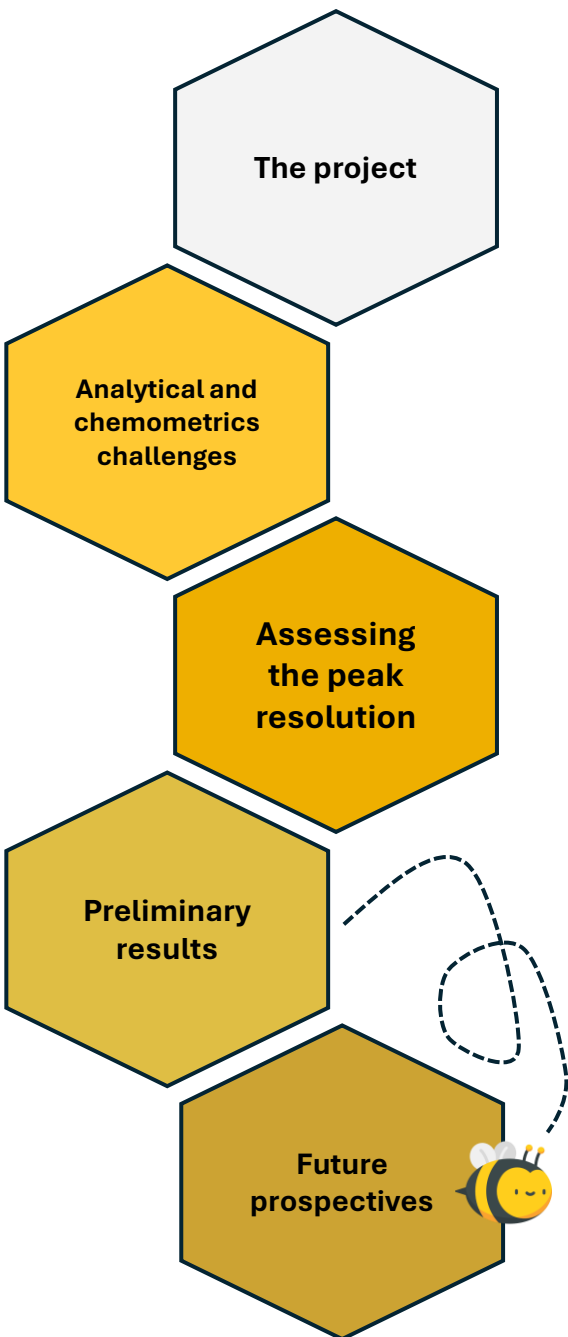
Future prospects



Identification based  
solely on RI and DT,  
standards  
confirmation  
needed!!!







- MCR-ALS and PARAFAC2 proved to be effective for the decomposition and resolution of GC-IMS data
- After careful preprocessing, GC-IMS data can be analysed using the same chemometric techniques as GC-MS, demonstrating that the same level of automation can be achieved
- GC-IMS partially separates Italian and Not-Italian samples and it is significantly influenced by the botanical origin
- GC-IMS combined with spectroscopic techniques allows for a good classification of Italian honey
- More acacia, wildflowers and citrus honey samples will be analysed for properly develop and validate a classification model



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