

#### Development of a multimodal approach to study plant-microorganisms interactions using mass spectrometry imaging, Raman imaging and non-targeted metabolomics

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## Introduction

450 million years ago, an essential symbiotic relationship called arbuscular mycorrhizal symbiosis (AMS) was formed between plants and glomeromycota fungi<sup>1</sup>, enabling the first land plants to colonize the soil<sup>2</sup>.

It begins with a molecular dialogue between partners located in the soil, leading to the development of specialized fungal structures in the roots, called **arbuscules**, facilitating the exchange of nutrients<sup>3</sup>.

Understanding this metabolic dialogue is a challenge, but innovative techniques such as LC-HRMS and MALDI\* mass spectrometry imaging enable a complementary localization and quantification of metabolites in AMS. The integration of stimulated Raman microscopy enhances our understanding at the cellular level in a non-invasive way.

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# **Untargeted Metabolomics**

### Data acquisition

by U-HPLC-HRMS (Thermo Scientific) **UHPLC : Vanguish with CAD and DAD detectors** 

MS : Q-Exactive Plus ESI +/-

Biphasic extraction following MetaToul AgromiX protocol (MTBE, MeOH,  $H_2O$ ).

*Lipidomic* analysis (apolar compounds) **C18** analysis (semi-polar compounds) Hilic analysis (polar compounds)

The first results of this study concern *Marchantia paleacea* mycorrhized by Rhizophagus irregularis.



A : Marchantia paleacea

B : Thallus embedded in agarose7%

C : Transverse section (100 $\mu$ m thickness using vibratome) with arbuscular mycorrhizal fungi (purple area)/







Data acquisition with MALDI-MSI\* (timsTOF fleX - Bruker)

> 10 µm resolution Positive mode





Data acquisition with Raman stimulated microscopy (STELLARIS 8 CRS - Leica)

### Sample preparation:

• MALDI matrix : DHB • Deposition with the HTX TM3 **sprayer** (HTX technologies LLC)

> Hey ! Scan me and download the file to see 3D PCA of discriminative segmentation !

**RED** = **Non mycorrhized** area ions **GREEN** = **Mycorrhized** area ions **BLUE** = **Centre of** mycorrhize ions





 $\Rightarrow$  Identification and mapping of Raman signals characteristic of biomarkers identified by LC-HRMS (single cell and hyphae scale) with tunable OPO laser.

*Optimization of analysis conditions:* 

 $\Rightarrow$  Magnification  $\Rightarrow$  Number of accumulations  $\Rightarrow$  Signal averaging

(A) BrightField snapshot of a mycorrhized M. paleacea cell at a x40 magnification with confocal Raman microscope.

(B) BrightField snapshot of a *R. irregularis* hyphae at a x40 magnification with confocal Raman microscope.

Hi again ! Scan me to see real-time vibrations signals from laser excitation !



\*Matrix-Assisted Laser Desorption-Ionization

### $\Rightarrow$ <u>Targeting</u> of biomarkers identified by LC-HRMS

## Conclusions

These results provide a great example on how multimodal approaches can help visualizing biological interactions between different organisms in a more constitutive way:

- → Global LC-HRMS analysis helps unravelling the metabolic segregation occurring during AMS establishment.
- → MALDI imaging helps localizing the metabolites involved at the tissue scale with high sensibility level.
- -> Despite not being as resolutive as MALDI, the high resolution provided by Raman stimulated microscopy helps us to localize metabolites families. Allowing us to better quantify exchanges occurring during AMS formation.



(A) Raman spectrum of a non-mycorrhized cell of *M. paleacea* between 1400 and 1700cm<sup>-1</sup>. (I 35 % ; SRS gain 100 ; accumulation 2) (B) Raman spectrum from the hyphae of *R. irregularis* between 1400 and 1700cm<sup>-1</sup>. (I 35 % ; SRS gain 88,5 ; accumulation 1) (C) Raman spectrum from the inside of a mycorrhized cell of *M. paleacea* between 1400 and 1700cm<sup>-1</sup>. (I 30 % ; SRS gain 100 ; accumulation 1)

Wavenumber (cm<sup>-1</sup>

 $\rightarrow$  Two main peaks around 1445cm<sup>-1</sup> and 1660cm<sup>-1</sup> which could be respectively attributed to the signal emitted by saturated fatty acids and unsaturated fatty acids.

 $\rightarrow$  The difference in intensity between non-mycorrhized and mycorrhized cells could be explain by the presence of arbuscular mycorrhizal fungi.

## References

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