

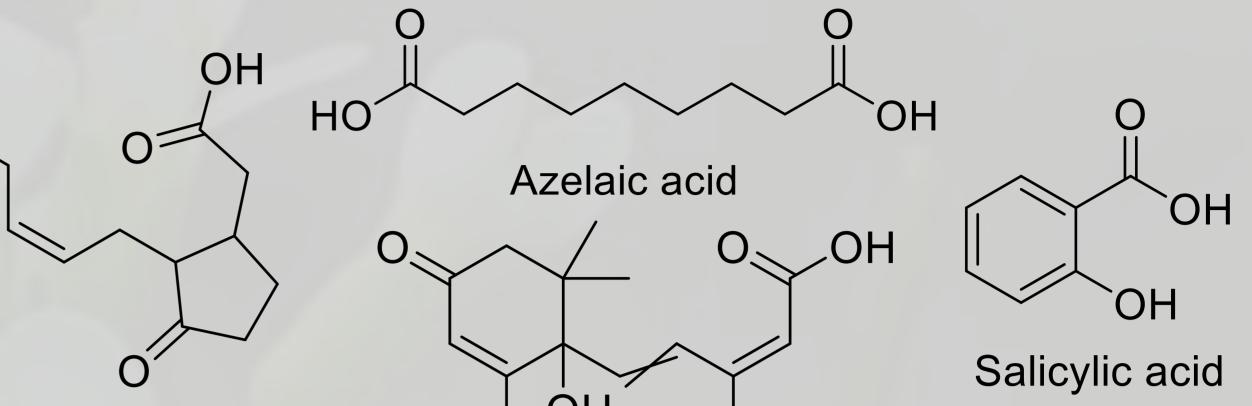
Study of the role of complex lipids in plant responses to stress



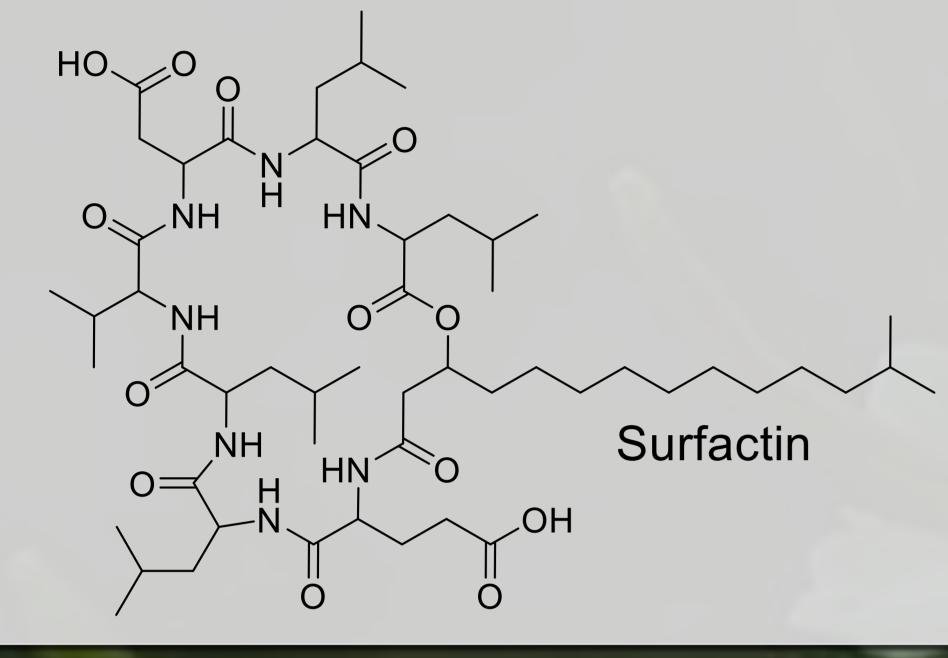
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Context

Plants are subjected to various biotic and abiotic stresses during their development, including wounding, pathogen attack, insects, viruses, temperature variations, drought, and many others. Among the plant responses to these stresses, the production of phytohormones, reactive and cytosolic calcium species notably induced. are oxygen Phytohormones in particular play a regulatory role in plant



development, reproduction and death.



Jasmonic acid

OH

Abscisic acid

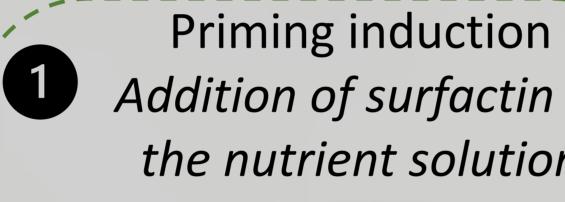
Nevertheless, plants are able to resist certain external attacks more effectively through interactions with non-pathogenic micro-organisms such as plant growth-promoting rhizobacteria (PGPR) i.e, the *Bacillus* genus. This is called microbial priming, which is part of the plant induced systemic resistance.

Among the secondary metabolites produced by these non-pathogenic microorganisms, Bacillus lipopeptides, and more precisely surfactins, are well known to be microbial elicitors that enable priming in plants.

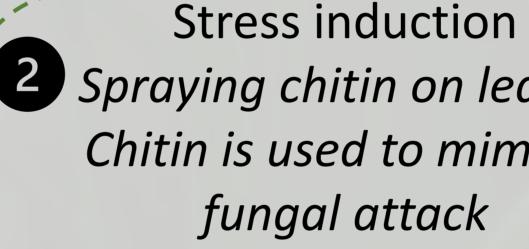
Methods

Cultivation of wild type 0 A. thaliana in hydroponics conditions

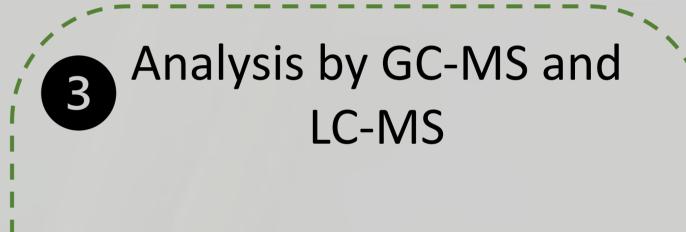




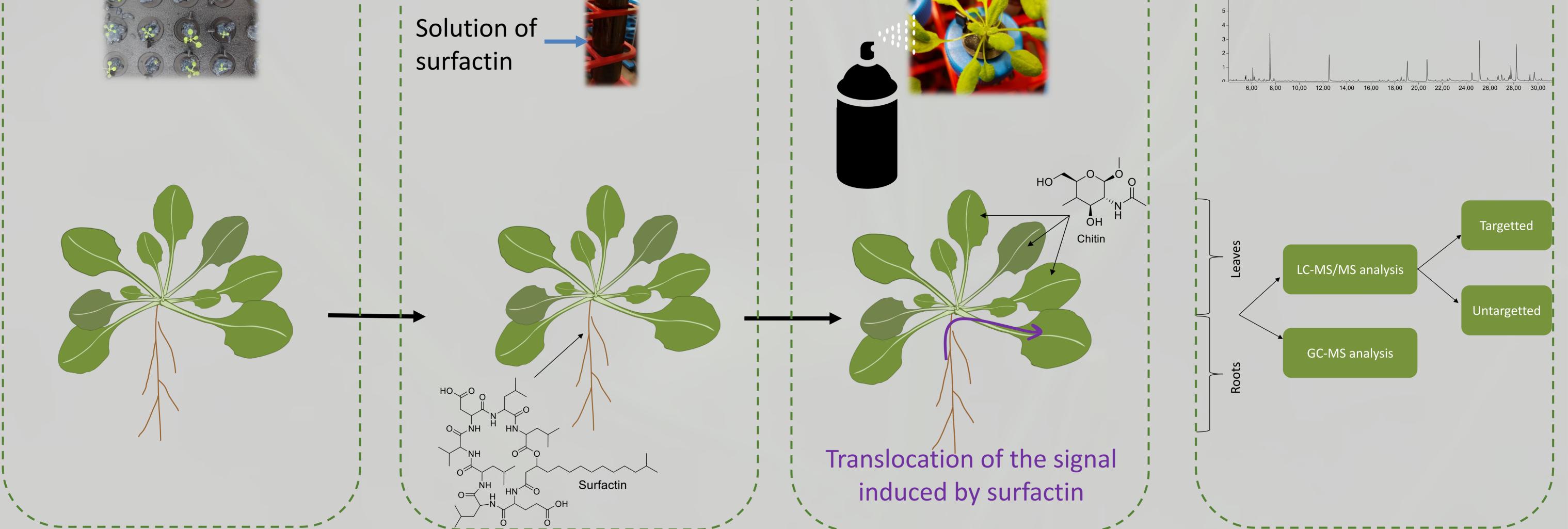
Addition of surfactin in the nutrient solution



2 Spraying chitin on leaves Chitin is used to mimic a fungal attack



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Conclusion and perspectives

The present study aims to understand how surfactin modifies the metabolome of root tissue and to identify and characterize the molecules that are translocated from roots to leaves. In addition, the immune response set up in leaves of plants that have been primed and then attacked by a pathogen will be explored. Two LC-MS/MS approaches will be combined to answer these aims: a targeted analysis of phytohormones and a non-targeted analysis. In addition to liquid chromatography analyses, particular attention will also be paid to volatile organic compounds emitted by the plant, and that will be analyzed by GC-MS. These two techniques will enable to characterize changes in the metabolome of leaves and roots following surfactin priming in A. thaliana.

References

Zhao B., & al. J. Agric. Food Chem (2021) Prsic J., & al. Front. Plant Sci. (2020) Jeworutzki E., & al. The Plant Journal (2010) Mousavi S., & al. Nature (2013)

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