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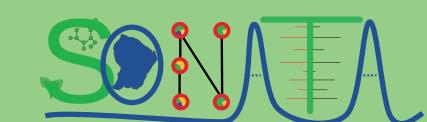
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Exploring the anti-mycobacterial potential of French Guiana's flora through a metabolomic study

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Context

Within the plant diversity of French Guiana's, many species are used by local populations for medicinal purposes. These plants, and their associated traditional uses, have been the subject of numerous census studies, and have attracted the attention of scientists for their therapeutic potential, particularly in the treatment of infectious diseases.

Bacterial infections such as tuberculosis (TB), caused by *Mycobacterium tuberculosis*, are a major public health problem in French Guiana. With a high incidence rate and the emergence of drug-resistant strains, there is an urgent need to explore therapeutic alternatives [1-2]. Local traditional knowledge could be a source of discovery for new anti-TB agents.

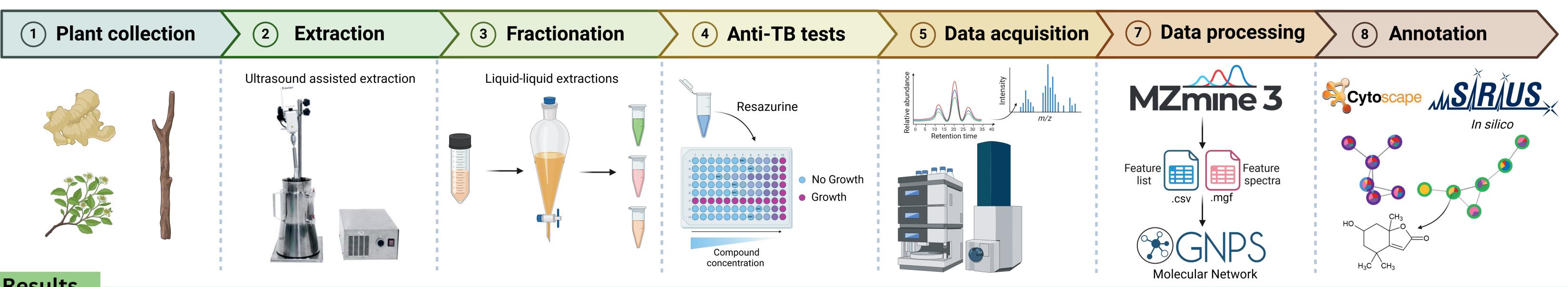
Objectives

In drug discovery from natural products, it can be challenging to describe the metabolites responsible for the biological activity. The use of the Mass Spectrometry based Molecular Network approach can facilitate structural dereplication and accelerate the annotation of structural entities in complex samples.

This study aims to facilitate the characterization of bioactive metabolites responsible for the anti-TB activity present in **9 plant species** from French Guiana.

Based on the literature, preliminary results indicate interesting activities for the plant species selected [3]. Some are also used in traditional medicine to treat tuberculosis symptoms (fever, cough, asthenia).





Results

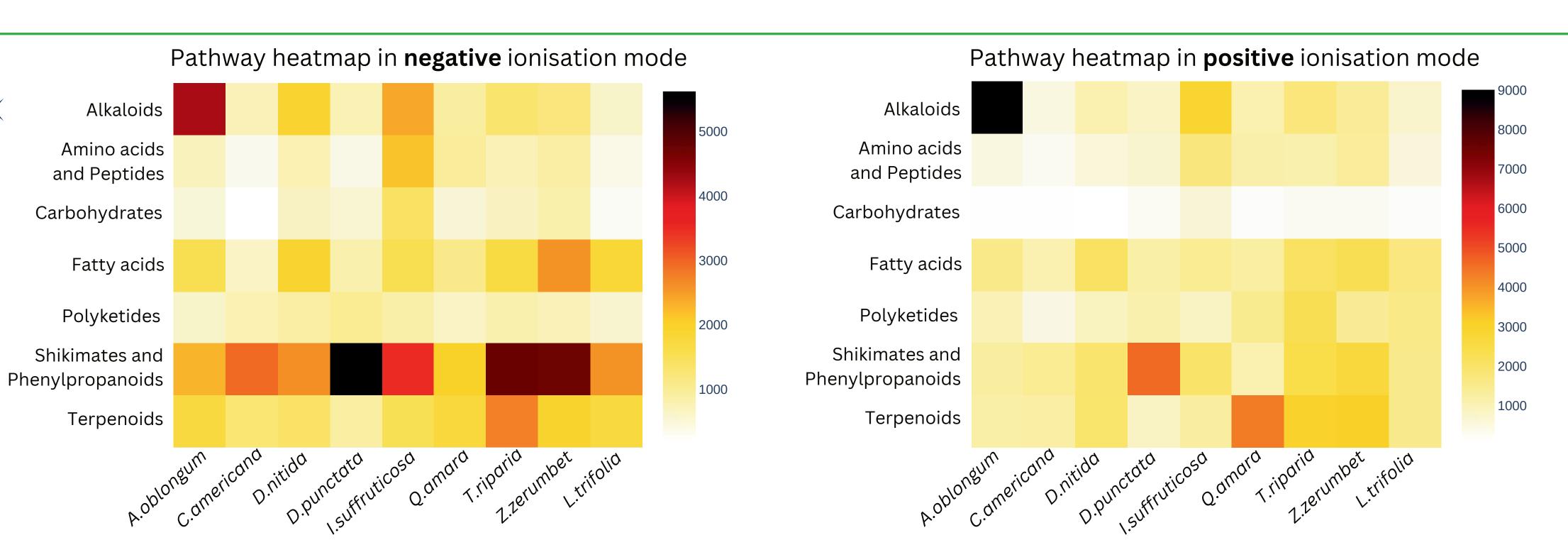
Chemodiversity

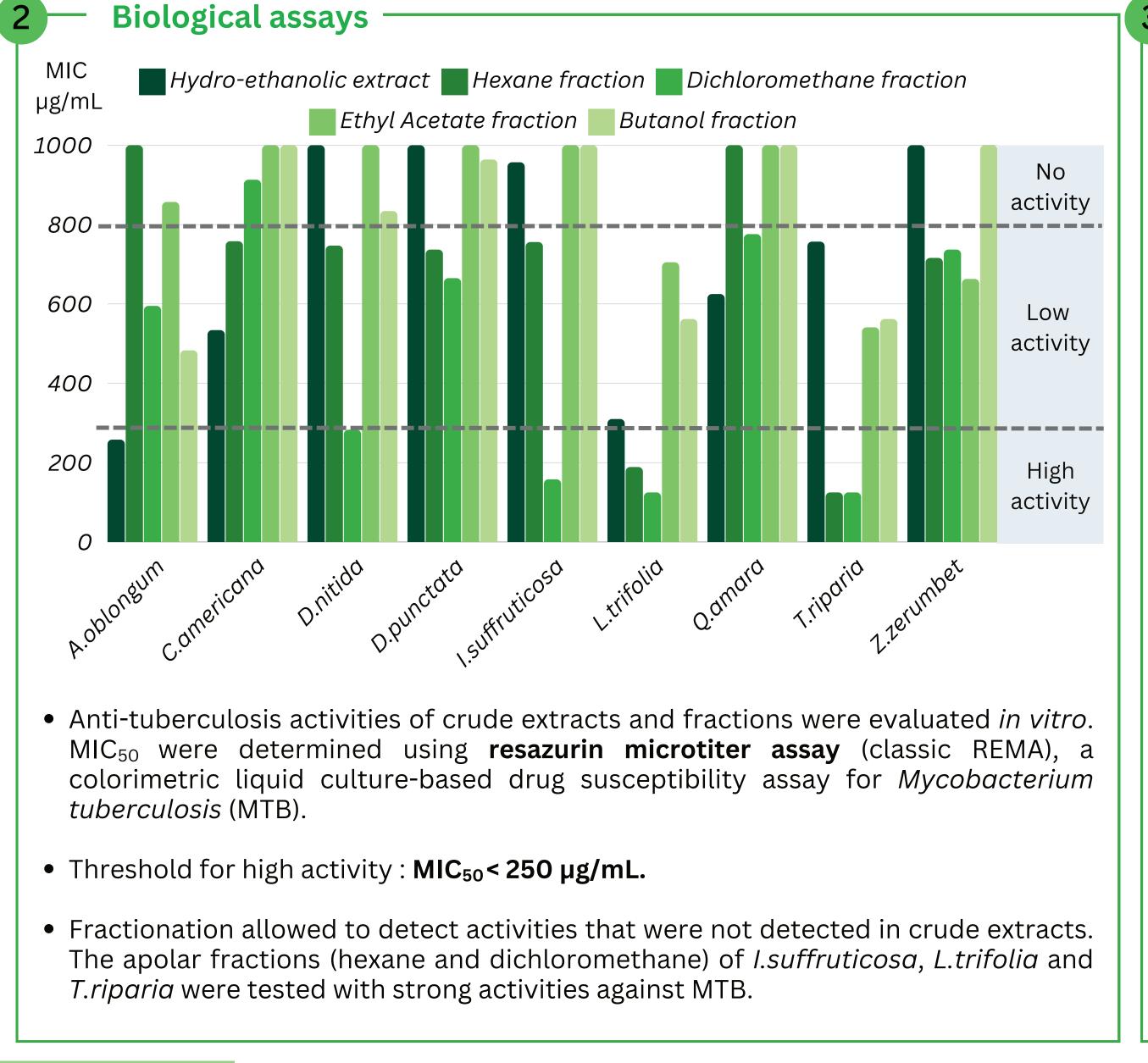
• Heatmaps were constructed based on the *in silico* prediction by CANOPUS module of the SIRIUS 5 software, and according to

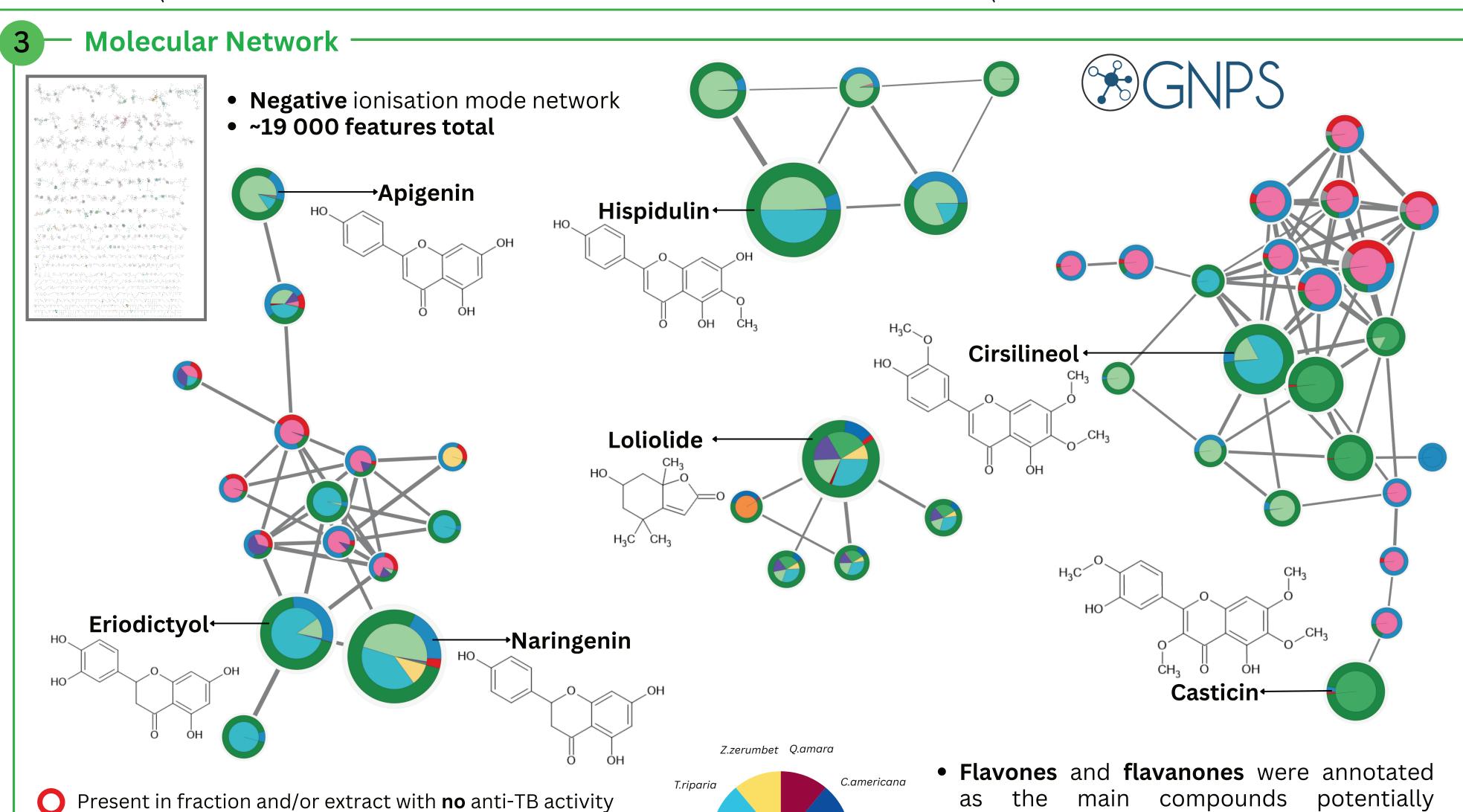
- The studied species contain mainly metabolites of the shikimates and phenylpropanonoids pathway: a large number of flavonoids and isoflavonoids were annotated.
- T.riparia and Q.amara are rich in terpenoids

the chemical pathways from NPClassifier.

• A.oblongum contains mainly alkaloids, specially of the tryptophan type.







Conclusion

The present experimental and analytical workflow allow us to prioritise data of biological interest, and annotate metabolites potentially responsible for the anti-TB properties in complex samples.

Present in fraction and/or extract with **low** anti-TB activity

Present in fraction and/or extract with **high** anti-TB activity

Relative abundance of the metabolite in the studied species

- Multiple flavonoids, of the **flavone** class, were annotated as potential new antitubercular compounds.
- Targeted purification of the proposed metabolites, coupled with NMR analysis will allow to confirm the proposed structures.
- Anti-TB assays will also be performed on isolated compounds to evaluate their activity.

References

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I.suffruticosa



responsible for the activity.

• Loliolide is a **beta-apocarotenoid**, already

tested with anti-TB properties in-vitro [4].



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