



## DEVELOPMENT OF ORIGINAL ANALYTICAL TOOLS FOR THE CHEMICAL PROFILING OF MICROALGAE

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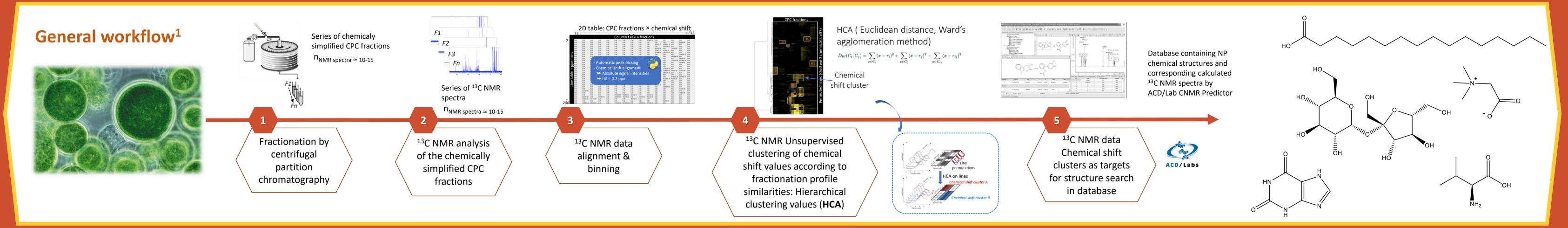


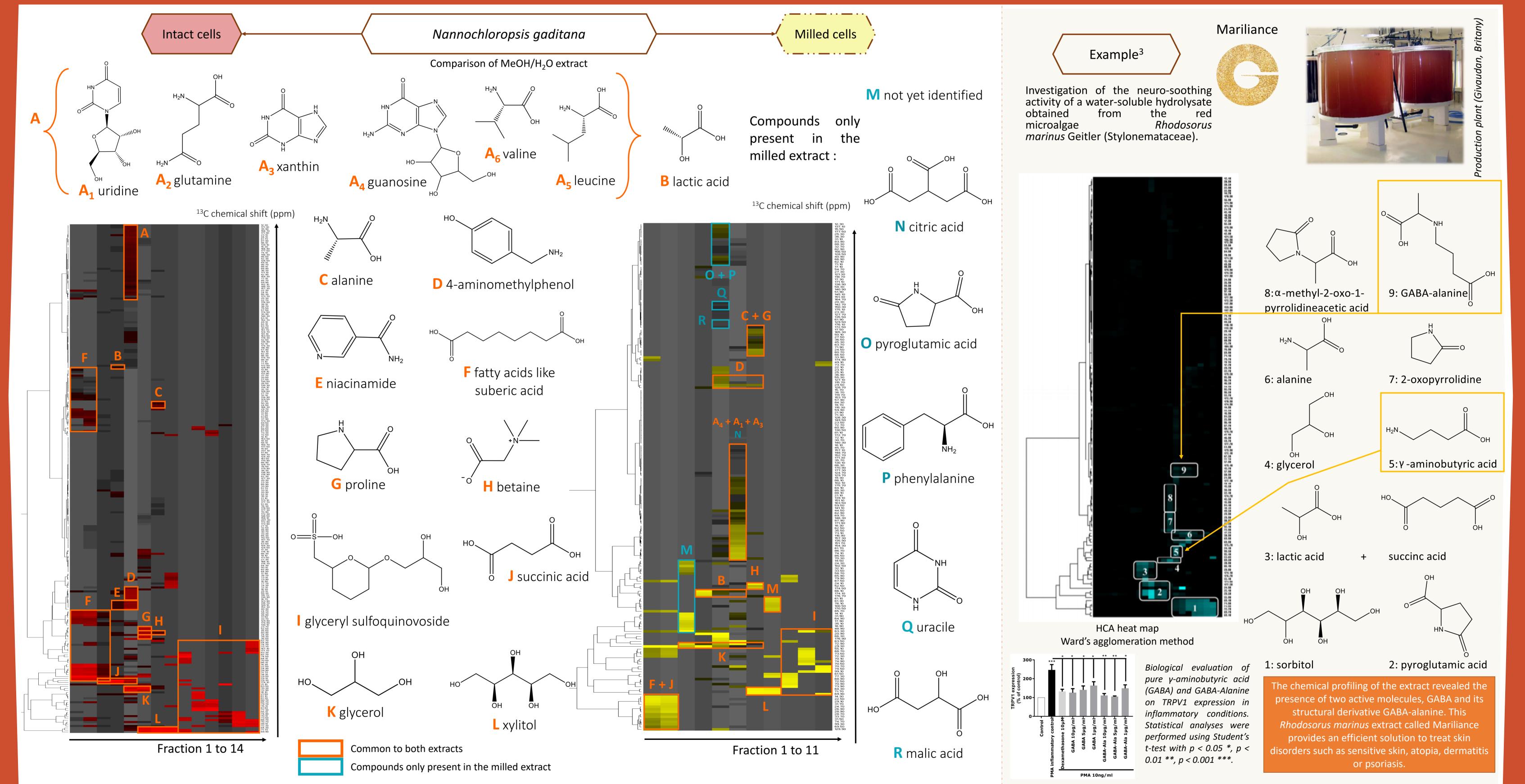
## Context

Research efforts are necessary to sustain the development of blue biotechnology as an emerging industrial field by diversifying industrial microalgae applications in high value added sectors such as materials, nutraceuticals, cosmetics or drugs. Access to these new products has to be done through the development of disruptive and safer processes with low environmental impacts as well as to adapt the micro-algae biorefinery according to chemodiversity potential, process routes, and market needs. Such an approach necessarily implies an increased knowledge of the chemical diversity of the "microalgae" raw material.

## Introduction

An original and efficient analytical tool was developed for the chemical profiling of microalgae extracts.<sup>1,2</sup> This dereplication strategy combines extract fractionation by **centrifugal partition** chromatography and <sup>13</sup>C NMR spectroscopy combined with unsupervised classification for the compound-by-compound grouping of experimental chemical shift values according to their chromatographic emergence profile. The grouping proceeds through the hierarchical clustering analysis performed on <sup>13</sup>C chemical shifts. The chemical shift clusters are then used as search targets in a database enriched with data from literature and predicted chemical shifts.





## **Conclusion & Perspectives**

The combination of an efficient multigram-scale fractionation method with <sup>13</sup>C NMR analysis and HCA for pattern recognition of <sup>13</sup>C signals across spectra of the fraction series enabled the direct identification of the main metabolites present in a crude natural extract (*i.e.* from microalgae) without compound-by-compound tedious, time and solvent consuming purification steps.<sup>4</sup> It also allows the study of the impact of the fractionation and production process on the chemical profile of the microalgae extracts.

<sup>1</sup>Hubert, J.; Nuzillard, J.-M.; Purson, S.; Hamzaoui, M.; Borie, N.; Reynaud, R.;

Renault, J.-H. Anal. Chem. 2014, 86 (6), 2955–2962.

<sup>2</sup>Bakiri, A.; Plainchont, B.; Emerenciano, V. de P.; Reynaud, R.; Hubert, J.; Renault, J.-H.; Nuzillard, J.-M. Mol. Inform. 2017, 36 (10), 1700027.

<sup>3</sup>Scandolera, A.; Hubert, J.; Humeau, A.; Lambert, C.; De Bizemont, A.; Winkel, C.; GoToS3 Kaouas, A.; Renault, J.-H.; Nuzillard, J.-M.; Reynaud, R. Mar. Drugs 2018, 16 (3), 96. **ALPO** <sup>4</sup>Wolfender, J.-L.; Nuzillard, J.-M.; van der Hooft, J. J. J.; Renault, J.-H.; Bertrand, S. Anal. Chem. 2019, 91 (1), 704–742.

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