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Metabolomic approach using LC-MS/MS analysis and molecular networking to follow up bioactive constituents of *Calophyllum inophyllum* nuts during drying process





SPECTROPOLE

Centre d'analyses chimiques

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Mont Tekao Cascade de Vaipo

Nuku Hiva collection sites

Strategy



Hatiheu, S08°49.679' W140°04.89

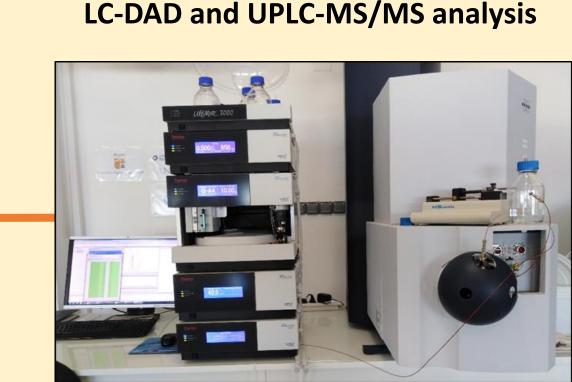


(A) Harvesting between July and September 2020 in the Marquesas Islands(B) Opening the fruits(C) Drying of the nuts from 0 to 10 weeks

DRYING PROCESS



Laboratory scale extraction:
Grinding of 15 ± 2 g of dry nuts,
maceration with EtOAc (ethyl
acetate), sonication and drying



 \simeq 135 resinoid extracts analyzed in French Polynesia (LC-DAD, Luna C18 (250 x 4.6 mm, 3 µm)) and Marseille (UPLC-QqToF-MS/MS, Luna C18 (150 x 2.1 mm, 1.6 µm))

PRE-PROCESSING ANALYSIS

Workflow4metabolomics

P-value: 0.032, NMC: 76.2%

R packages: mixOmics (*plsda* function); ade4 (graphical representation); RVAideMemoire (discriminant compounds)

MOLECULAR NETWORKS

GNPS Cytoscape

Calophyllic action Tamanolide

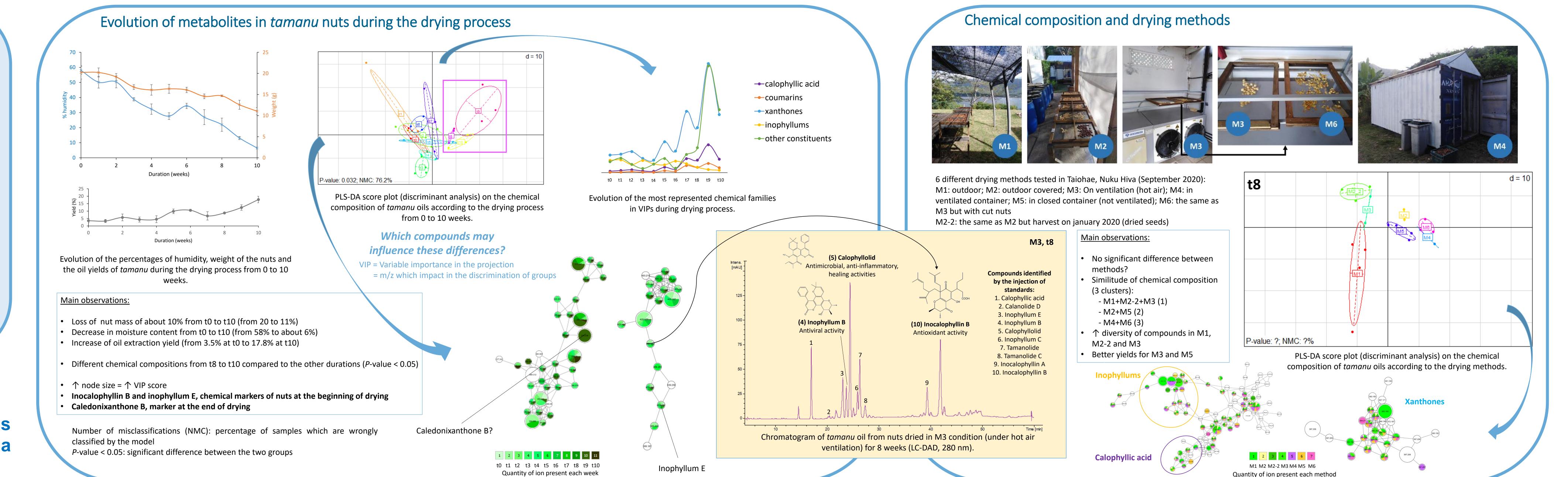
Context

Tamanu oil, obtained from the nuts of Calophyllum inophyllum L. (Calophyllaceae), was traditionally used to cure various skin problems and ailments in French Polynesia [1]. Nuts and containing oil are also used for skin care [2]. They were reported to treat different kinds of skin affections and used as natural cosmetic ingredient [3]. Since the drying of nuts is an important stage for oil preparation, the objective of our study was to develop an analytical method to evaluate oil quality (chemical composition and oil yields) during nuts drying process following different drying parameters (abiotic factors or duration). In this project, we propose to use a metabolomic approach combining LC-MS/MS analysis and molecular networking to identify markers inducing chemical composition variability during drying process of tamanu nuts and previously unidentified constituents.

Objectives:

(1) Evaluate quality of oil by oil yields and chemical composition during drying process (2) Assess the drying process efficiency by a metabolomics approach

Results



Alignment on W4M

110 Samples

1176 ions

Conclusion and perspectives

Using a metabolomic approach combining LC-MS/MS analysis and molecular networking, the obtained data revealed differences in the metabolites chemical classes occurring in resulting oils (neoflavonoids, coumarins, xanthones and triterpenes) and provided the identification of precedently unreported constituents. **Inophyllums seemed to be more abundant in the beginning of drying process compared to calophyllic acid** after nine weeks. For the first time, our study provides new findings regarding the occurrence and evolution of the metabolites in *tamanu* nuts during the drying process. The developed method provided a powerful analytical tool aiming a better identification of bioactive components formed in *tamanu* nuts during drying process and will be helpful to control high quality oil for a natural active cosmetic ingredient. An efficient metabolomic approach was thus implemented to identify markers inducing variability in chemical composition during drying process of *tamanu* nuts and previously unidentified constituents. Thus, this set-up analytical method successfully applied in *C. inophyllum* could be used to the study of metabolites of a wide range of plant metabolites content.

Acknowledgements

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References

[1] Pétard P., Ed. Revue et augmentée HaerePo No Tahiti, 1986, 224-225. [2] Léguillier et al., PlosOne, 2015, 10(9):e0138602, doi:10.1371/journal.pone.0138602. [3] Raharivelomanana et al., OCL, 2018, 25(5), doi:10.1051/ocl/2018048.



