

Biochemical characterization of the genetic resources of wild coffee trees collection using near infrared spectroscopy

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Introduction

This study aims to enrich the CIRAD infrared spectral database (1) for coffee beans (>2200 references, mainly C. arabica and C. canephora samples) and improve the predictive models for the major seed constituents (2) using the chemical diversity of wild coffee species present at the Coffea Biological Resources Centre (3) (maintained by IRD and CIRAD in Reunion Island).

Materials/Methods

462 coffee samples from 32 species were dried ground and analyzed for their NIR spectrum. Based on spectra diversity, 90 representative samples were selected using Mahalanobis distances (fig.1) and analyzed for their caffeine, trigonelline, fat and chlorogenic acids (CGA) contents. New models were developed.

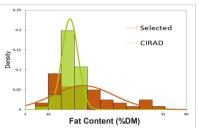


Figure 2: Fat content distribution for CIRAD database and new selected samples

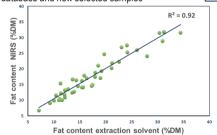


Figure 3: Fat content predicted vs laboratory values

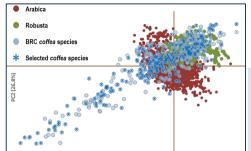






Figure 1: Spectral repartition on PC1 and PC2

Results/Discussion

The 90 samples originated from 22 coffee species.

The caffeine content ranged from 0 to 3.3%, trigonelline from 0.25 to 1.52%, CGA from 0,22% to 10,53% and fat from 7.12% to 34,45% (DMb). The predicted models present standard errors of 0.08%, 0.07%, 0.58% and 0.48% for respectively caffeine, trigonelline, CGA and fat. These performances are close to original models with an increase of the content ranges (e.g. Fat, fig.2 & 3).

Conclusion/Perspectives

This study demonstrated the efficiency of the use of genetic diversity present in the Coffea genus to enhance the robustness of the database. The resulting calibrations cover a larger range of values. Mahalanobis distances permitted an efficient improvement of the representativeness with a limited number of samples. New calibrations will be applied to the whole database (400 samples) in order to describe biochemical diversity in wild coffee species.

References:

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