

Artigo

Botanical Characterization of Brazilian honeys by Nuclear Magnetic Resonance Spectroscopy and Chemometrics

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Rev. Virtual Quim., 2016, 8 (1), 277-287. Data de publicação na Web: 8 de janeiro de 2016<http://rvq.sbq.org.br>**Caracterização Botânica de Méis Brasileiros por Espectroscopia de Ressonância Magnética Nuclear e Quimiometria**

Resumo: O mel é um dos alimentos mais estudados devido às suas propriedades nutricionais e medicinais. RMN pode ser usado na análise de alimento para discriminação da origem e identificação de biomarcadores utilizando uma abordagem metabolômica. Neste estudo, nós apresentamos um exemplo desta estratégia para tipificar amostras de mel de diferentes origens botânicas. Onde 53 extratos de méis ricos em polifenóis foram preparados utilizando resina de adsorção XAD-4 e os seus espectros de RMN de ^1H associados à análise multivariada foram utilizados para detectar possíveis biomarcadores de origem floral. A análise de componentes principais foi aplicada com sucesso na discriminação entre amostras de mel de eucalipto, laranjeira e cambará.

Palavras-chave: Classificação de mel; RMN; Análise por Componentes Principais (PCA).

Abstract

Honey is one of the most studied foods due to its nutritional and medicinal properties. NMR can be used in food analysis for origin discrimination and biomarker identification using a metabolomic approach. In this study, we present an example of this strategy to discriminate honey samples from different botanical origins. The NMR spectra coupled with multivariate statistical analysis of 53 polyphenol-rich extracts prepared using adsorption resin XAD-4 of selected honey samples were analyzed to detect possible biomarkers of their floral origin. Principal component analysis was successfully applied for the discrimination among eucalyptus, orange and cambara honey samples.

Keywords: Honey classification; NMR; principal component analysis (PCA).

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Botanical Characterization of Brazilian Honeys by Nuclear Magnetic Resonance Spectroscopy and Chemometrics

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1. Introduction

Honey is a complex mixture of around 200 compounds, but carbohydrates make up most of the content. Minor substances such as enzymes, aminoacids, minerals, organic acids, vitamins, waxes and pollen grains can also be found.¹⁻³

Thus, honey is basically a supersaturated fructose and glucose solution, with about 17% of water in its final composition. That is due to the action of water evaporation promoted by the bees, which keep fanning the nectar to expose it to

air and therefore eliminate the excess humidity. Also, they use their proboscis (long hairy tongue) to absorb part of the water and insert some enzymes. Besides being a homeostasis process, water removal increases the concentration of fructose and glucose, making honey a more stable food and therefore less susceptible to microbiological contamination.⁴

Previous researches showed that honey has functional properties in humans that may promote health improvement, but which greatly depend on the floral origin.⁵ The control of commercial honeys, therefore, requires the correct determination of the

floral origins as part of the overall evaluation of the quality of honey.

Honey floral origin is generally determined by pollen analysis (melissopalynology), as the type of pollen may reflect the nectar source. However, melissopalynology has some limitations: specialized professionals with a great knowledge of morphology are necessary to ensure trustworthy results. Furthermore, some pollen grains, such as those from *Citrus* species and members of the *Lamiaceae* family, are sub-represented in the pollinic spectra of honey, while other species such as *Eucalyptus*, *Castanea* and *Myosotis* are over-represented. Thus, a minimum of 10% of *Citrus* sp. pollen content is enough to consider the honey as unifloral, while a minimum of 70% of *Eucalyptus* pollen is necessary to classify it as unifloral.⁶ The difficulties on the determination of pollen content in some types of honey may afford wrong results if not confirmed by other techniques such as the determination of floral markers by high resolution liquid chromatography or gas chromatography.⁷ Many other studies were carried out to develop other methodologies and other analytical procedures, generally associated to chemometric analysis to help evaluate the honey's floral origin.⁸

Among many other analytical methods used for food characterization, Nuclear Magnetic Resonance (NMR) has been applied to food with the aim of characterization as well as detection of adulteration and has earned general acceptance as an efficient method^{9,10} due to its non-destructive feature, along with high reproducibility and sensitivity. This new screening method using ¹H NMR has been combined with chemometrics. The approach consists in applying principal component analysis (PCA) to reduce the data dimensionality.

Lately, an increasing interest in the development of analytical methods that may complement pollen analysis on the determination of honey floral origin has been observed in the literature. The analysis of phenolic compounds by high pressure liquid chromatography (HPLC) has been shown to

be an efficient method in the study of floral and geographic origins of honey.¹¹⁻¹⁴ In the early years, researchers have employed aminoacid analysis¹⁵, volatile compounds,^{16,17} phenolic compounds,¹¹ aromatic acids and their esters¹⁸ and carotenoids¹⁹ for the determination of floral and geographic origins of European honeys, by using gas chromatography coupled to mass spectrometry.

2. Experimental

2.1. Reagents

Methanol used for the analyses was spectrophotometric grade (VETEC), and water was previously passed through a Millipore filter (Millipore Direct-Q UV with pump). Reference compounds benzoic acid, *m*-coumaric acid, *m*-methoxycinnamic acid, naringerin, kaempferol, caffeic acid, myricetin, gallic acid, ferulic acid, quercetin, *p*-coumaric acid, protocatechuic acid, syringic acid were all purchased from Sigma-Aldrich.

2.2. Sample Collection

Fifty-three honey samples [seven cambara (*Gochnatia* spp, Asteraceae), sixteen multifloral, thirteen orange (*Citrus* sp) and seventeen *Eucalyptus* spp. (Myrtaceae)] were collected from different regions of the states of Rio de Janeiro and São Paulo, Brazil. Samples were collected from 2004 to 2010 and kept stored at 4 °C until analysis

2.3. Sample Preparation

The procedure for the fractionation of honey was carried out as previously described.²⁰⁻²² Honey samples (50 g) were thoroughly mixed with 250 mL of distilled water and the solution pH was adjusted to 2

with concentrated HCl followed by filtration through cotton wool to remove solid particles. The filtrate was mixed with 75 g of Amberlite XAD-2 resin, pore size 9 nm, and particle size 0.3-1.2 mm (Supelco, Bellefonte, PA, USA) and magnetically stirred for 10 min. The mixture was then packed into a glass column (25 x 2.0 cm). The column was washed with 250 mL of acidified water (pH 2 with HCl) and subsequently rinsed with 300 mL of neutral distilled water to remove all sugars and other polar compounds in honey. The phenolic compounds were eluted from the sorbent with 500 mL of methanol. The methanol extracts were concentrated under vacuum at 40 °C in a rotary evaporator. The residue was dissolved in 10 mL of distilled water and extracted five times with 10 mL of ethyl acetate. The ethyl acetate extracts were combined and the solvent was removed under vacuum. The dried residue was then redissolved in 1 mL of methanol (HPLC grade) and filtered through a 0.45 µm pore size membrane filter (Sartorius Stedim Biotech, Germany). Three replicate extractions were performed for each sample, and the standard deviation did not exceed 5%. The yields of extracts were expressed as a mean of three extractions and ranged from: *Citrus* (10 to 32 mg), *Cambara* (24 to 64 mg) and *Eucalyptus* (17 to 22 mg) per 50 g of honey.

2.4. ¹H NMR Spectroscopy

¹H NMR spectra were acquired for all 53 samples on a Bruker Avance Ultra Shield 500 spectrometer (Bruker Biospin GmbH, Rheinstetten, Germany) operating at 500.13 MHz using a broadband detection 5 mm probe. All the spectra were obtained at 298 K, using a 30° pulse, an acquisition delay of 2 s and 16 transients and a constant receiver gain. The spectra were processed using ACD/1D NMR Processor (Advanced Chemistry Development, Inc, Canada). Phasing, baseline correction and integration were done automatically by group treatment of all 53 spectra. The whole spectra from 0.00 to

12.00 ppm were integrated using ACD's Intelligent Bucketing, resulting in 221 integrals for each sample.

2.5. Sample Fortification

Fortification was performed by successive addition of 1.0 mg of 13 pure phenolic and flavonoid standards in each honey extract previously identified by HPLC, as shown in **Figure 1**. ¹H NMR spectra of the corresponding solutions were acquired, under the same conditions described above, in order to assign the corresponding signals of the standards in the different honey samples.

2.6. Multivariate Analysis

Principal component analysis (PCA) is an important multivariate tool for exploratory data analysis.²³ It aims at dimension reduction, i.e. using fewer variables to represent a data set. This complexity reduction is achieved through an orthogonal linear transformation of the original coordinate system so that the greatest variance by any projection of the data comes to lie on the first coordinate (called the first principal component), the second greatest variance on the second coordinate and so forth.

The new variables (loadings) are linear combinations of the original variables and have no physical meaning. However, they provide a simpler way to visualise and understand the original data set.

In NMR, in multivariate analysis studies, the variables (the X matrix) are the integrals of the whole or part of the spectrum for each sample. Each sample is a point in an *n* dimensional space of integrals. When PCA is performed, each sample can be represented in a lower dimensional space so that unseen relationships and groups can be visualized.

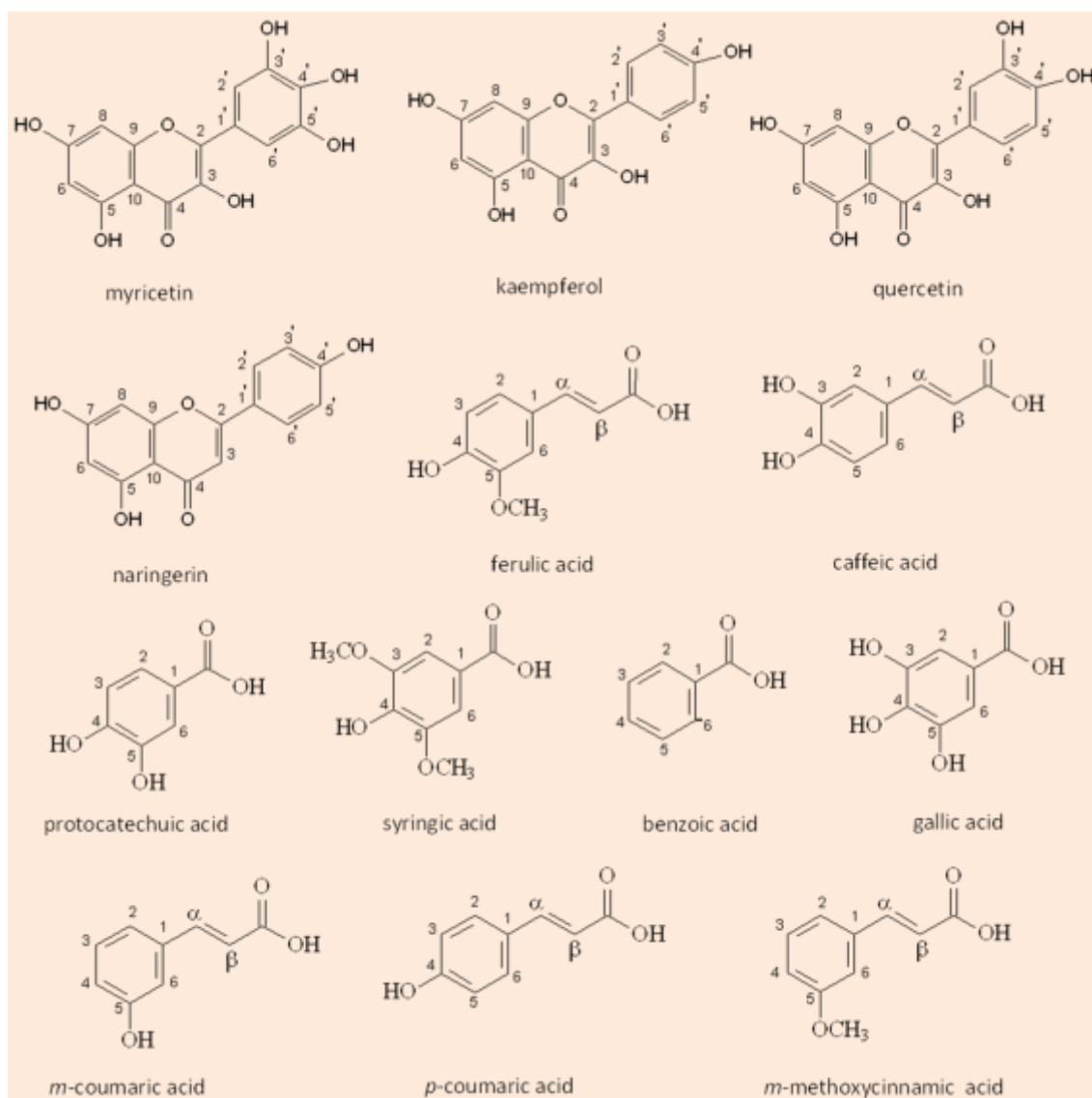


Figure 1. Structures of phenolic compounds identified in honey extracts after fortification

2.7. HPLC analysis of the phenolic profile of honey

Polyphenol identification analyses were carried out using an HPLC apparatus (LC-20AT, Shimadzu, Japan) equipped with a diode array detector (DAD, SPD-M20A) coupled to a LCStation ChemStation data-processing station. The column used was a C18 LiChroCART (250 mm x 4 mm; 5 μ m; Merck, Darmstadt, Germany), operating at 35 $^{\circ}$ C. The mobile phase consisted of solvent A (water and acetic acid, 99:1) and solvent B (water, acetonitrile and acetic acid, 59:40:1).

The gradient program was as follows: from 25 to 100% B in 30 min, and then isocratic by 10 min. The injection volume was 20 μ L, and the flow rate was 1 mL min⁻¹. Chromatograms for the phenolic acids were monitored at 270 nm and for flavonoids at 360 nm. The identification of phenolic compounds was based on a comparison of chromatographic data (retention times and UV spectra) with authentic markers.

3. Results and discussion

The ^1H NMR spectra of the honey extracts exhibited characteristic signals for phenolic compounds, as shown in **Figure 2** (fortification), which were confirmed by

HPLC-DAD (**Figure 3**). Due to the overlapping of signals, their attribution in the NMR spectra was only possible for major phenolic compounds after fortification. The detailed attribution of the phenolic reference compounds (**Figure 1**) is summarized in **Table 1**.

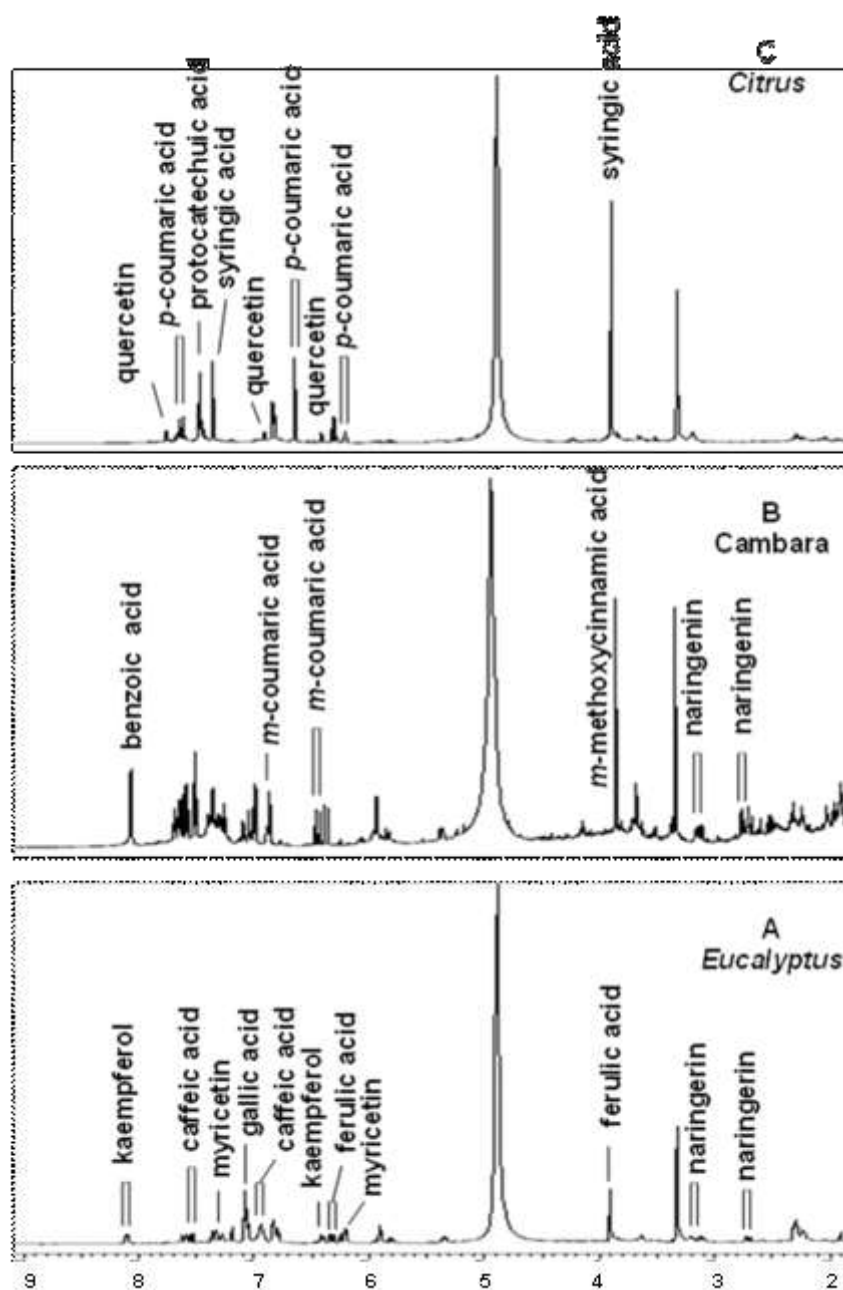


Figure 2. Representative 500 MHz ^1H NMR spectra (δ 2.0-9.0) of (A) *Eucalyptus*, (B) Cambara and (O) Orange honey extracts along with the assignment of resonances following spiking of 13 standard polyphenols. (For resolution reasons the spectra presented were processed using Gaussian multiplication of the FID (LB = -2, GB = 0.1 prior to Fourier transform)

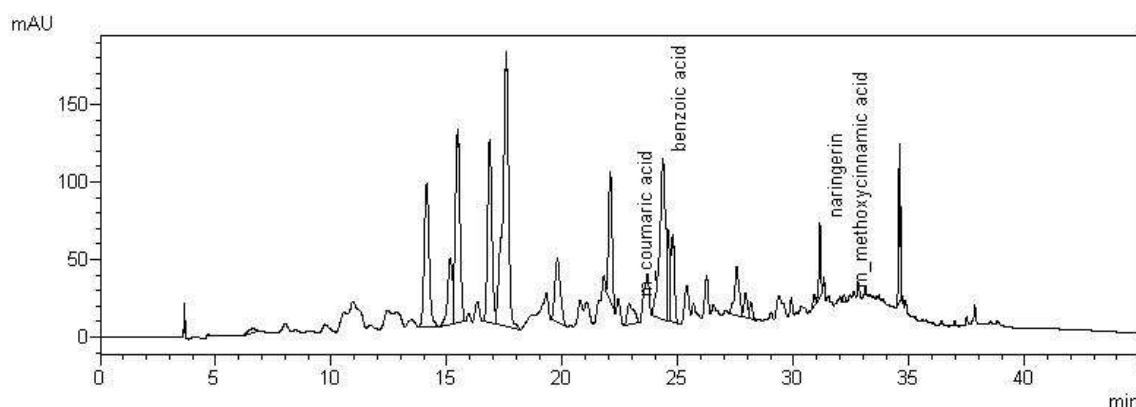


Figure 3. Chromatogram of the honey extract of Cambara at $\lambda= 280\text{nm}$

Table 1. ^1H NMR chemical shifts of phenolic components detected in honey

N°	Compound	$\delta^1\text{H}$ (multiplicity, assignment)
1	benzoic acid	8.04 (dd, H1, H6)
2	<i>m</i> -coumaric acid	6.41 (dd, H- β), 7.86 (dd, H- α)
3	<i>m</i> -methoxycinnamic acid	3.84 (s, OMe3)
4	naringerin	2.7 (dd, H3), 3.09 (dd, H3)
5	kaempferol	8.08 (d, H2', H6'), 6.41 (d, H8)
6	caffeic acid	7.53 (d, H- α), 6.93 (dd, H 6)
7	myricetin	7.28 (s, H-2'), 6.21 (d, H-6)
8	gallic acid	7.08 (s, H2, H6)
9	ferulic acid	6.47 (d, H- β), 3.90 (s, OMe5)
10	quercetin	6.89 (d, H5'), 6.40 (d, H8), 6.19 (d, H6)
11	<i>p</i> -coumaric acid	7.60 (d, H- α), 6.28 (d, H- β), 6.80 (d, H3, H5)
12	protocatechuic acid	7.46 (dd, H1, H6)
13	syringic acid	7.32 (s, H2, H6), 3.87 (s, OMe3, OMe5)

s, singlet; d, doublet; dd, doublet of doublet

The ^1H NMR spectra were segmented, integrated and normalized, and the data were sub-projected to PCA. Two PCA's were performed, one using multifloral and orange honey samples (**Figure 4**) and the other using

cambara, orange and *eucalyptus* monofloral honey samples (**Figure 5**). Both analyses afforded a very clear discrimination among the honey floral origins, thus becoming a good alternative to melissopalynology.

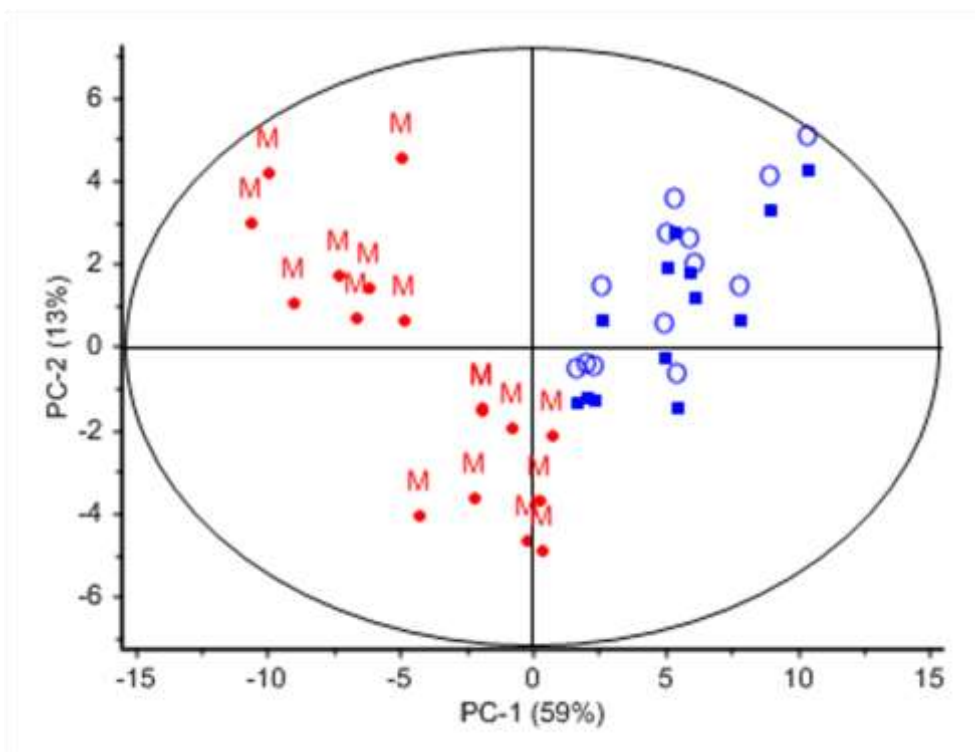


Figure 4. PCA scores plots derived from ¹H NMR spectra of orange and multiflora honey extracts: (O) Orange and (M) Multiflora

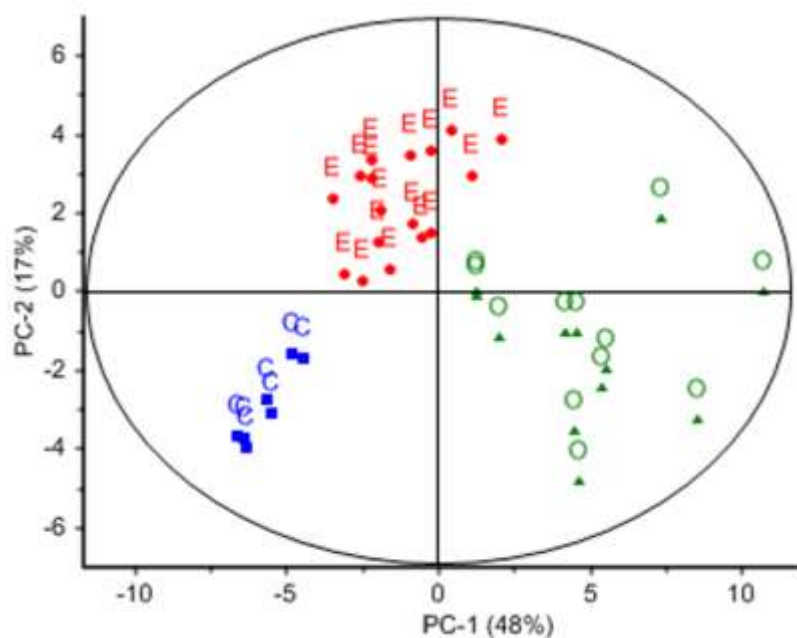


Figure 5. PCA scores plots derived from ¹H NMR spectra of orange (O), cambara (C) and eucalyptus (E) honey extracts

Biomarkers characteristic of some flora responsible for the clear discrimination have been described in the literature, being observed in the PCA.²⁴ Some of these

biomarkers were found in a great part of the samples studied in this work. Figure 2 (fortification) shows the major phenolic compounds found in each floral origin, which were used in the fortification in order to make it possible to identify them in the samples, due to their low concentration in the honey extracts.

Many studies have been carried out to seek for alternatives to sensorial and melissopalynological honey typification.²⁵ Among them, ¹H NMR associated to chemometrics has been highlighted. There is a huge diversity in honey constituents, which depends intimately on its botanical origin, though seasonal and environmental factors may also interfere on its composition, including its phenolic profile. Many authors have suggested the possible correlations between floral origin and honey's phenolic profile.²⁶ Therefore, flavonoids, phenolic acids, aminoacids and some aromatic and heterocyclic aldehydes are being evaluated in order to help distinguish honey floral origin.

The presence of some individual components or group of compounds is being evaluated as possible biomarkers that may be used in honey typification. The flavonoids myricetin, tricetin, luteolin, quercetin, and kaempferol have been suggested as floral markers for *Eucalyptus* honeys, as well as gallic acid and some other benzoic acid derivatives²⁷, while hesperitin, quercetin, crisin, caffeic acid, *p*-coumaric acid and ferulic acid have been described as possible *Citrus* honey markers.²⁷

4. Conclusions

In this study, phenolic compounds such as myricetin, kaempferol and gallic acid were identified in *Eucalyptus* honey samples, as well as naringenin, caffeic and ferulic acids. Quercetin and *p*-coumaric acid could also be identified in citrus samples, which have already been described as possible markers²⁷, as well as syringic and protocatechuic acids. Finally, *p*-methoxycinnamic acid and *m*-

coumaric acid were identified in all the Cambara honey samples, as well as naringenin and benzoic acid. Principal component analysis was successfully applied for the discrimination among *eucalyptus*, orange and cambara honey samples. Finally, the association of PCA and NMR techniques might be an important tool to evaluate honeys with poor pollen content and even with excess pollen content, which can mask the nectar predominance of other species and interfere on the melissopalynological analysis.

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