

Phytochemical analysis and botanical origin of *Apis mellifera* bee pollen from the municipality of Canavieiras, Bahia State, Brazil

Análise fitoquímica e origem botânica de uma amostra de pólen apícola do município de Canavieiras, Estado da Bahia, Brasil

Giuseppina Negri^{1*} , Lidia Maria Ruv Carelli Barreto², Fábila Lugli Sper³, Claudemir de Carvalho³, Maria da Graça Ribeiro Campos⁴

¹ Universidade Federal de São Paulo (UNIFESP), Departamento de Medicina Preventiva, Centro Brasileiro de Informações sobre Drogas Psicotrópicas, São Paulo/SP - Brasil

² Universidade de Taubaté (UNITAU), Taubaté/SP - Brasil

³ Faculdade de Pindamonhangaba (FUNVIC), Pindamonhangaba/SP - Brasil

⁴ Universidade de Coimbra, Faculdade de Farmácia, Centro de Química de Coimbra /Polo III, Azinhaga de Santa Comba, Coimbra - Portugal

*Corresponding Author

Giuseppina Negri, Universidade Federal de São Paulo (UNIFESP), Centro Brasileiro de Informações sobre Drogas Psicotrópicas, Departamento de Medicina Preventiva, Rua Botucatu, 740, Vila Clementino, São Paulo/SP - Brasil, e-mail: gnegri@terra.com.br

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Abstract

An *Apis mellifera* bee pollen sample from Bahia state in Brazil was studied to evaluate its botanical origin and phytochemical composition. The bee pollen sample was collected in the municipality of Canavieiras, in an area with a very high predominance of *Cocos nucifera* L (Aracaceae), which was identified as the major taxon (99%), thus being the possible botanical origin of this pollen. The main constituents found in the non-polar extract analysed by GC-EI-MS were saturated fatty acids and long chain esters, together with phytosterols such as ergosta-5,24(28)-dien-3-ol, campesterol and sitosterol, detected in smaller quantities. Flavonoid glycosides, as well as hydroxycinnamic acid amide derivatives were detected in the polar extract analysed by HPLC-ESI-MS/MS. The presence of flavonoid glycosides, hydroxycinnamic acid amide derivatives, fatty acids and phytosterols have been reported in many bee pollen taxa. To the best of the authors' knowledge, this is the first study of the chemical composition of bee pollen from *C. nucifera*, which is cultivated for its coconut fruit.

Keywords: *Apis mellifera* bee pollen; *Cocos nucifera*; Fatty acids; Flavonoid glycosides; Phytosterols.

Resumo

Uma amostra de pólen apícola de abelhas *Apis mellifera*, coletada no Estado da Bahia, foi estudada com o objetivo de identificar a sua origem botânica e a sua composição química. A amostra de pólen apícola foi coletada no município de Canavieiras, em uma área onde ocorre a predominância de *Cocos nucifera* L (Aracaceae), a qual foi identificada como o maior táxon (99%) desta amostra, podendo ser considerada a origem botânica deste pólen. Os principais constituintes encontrados no extrato não polar, que foi analisado por CG-EM, foram ácidos graxos saturados e ésteres de cadeia longa, além de esteroides, tais como ergosta-5,24(28)-dieno-3-ol, campesterol e sitosterol, os quais foram detectados em menor quantidade. Flavonoides glicosídeos e amidas derivadas do ácido hidroxicinâmico foram encontrados no extrato polar, que foi analisado por HPLC-ESI-MS/MS. A presença de flavonoides glicosídeos e amidas derivadas do ácido hidroxicinâmico, ácidos graxos e esteroides foi reportada em muitas amostras de pólen, por vários autores. Este é o primeiro estudo que avalia a composição química de uma amostra de pólen originada de *Cocos nucifera* L, uma planta que é cultivada para a produção do coco.

Palavras-chave: Pólen apícola; *Cocos nucifera*; Ácidos graxos; Flavonoides glicosídeos; Esteroides.



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1 Introduction

Bee pollen is an apicultural product of commercial interest due to its high nutritional value. As a source of energy and proteins for human nutrition, it is used to produce dietary supplements in the form of tablets, capsules and granulates. Bee pollen extracts may also be prepared in the form of alcohol and aqueous extracts (FEÁS et al., 2012; KOMOSINSKA-VASSEV et al., 2015; BÁRBARA et al., 2015; ZHOU et al., 2015). Pollen is the male gametophyte of flowers. Bee pollen is pollen collected by the honey bee *Apis mellifera* for the purpose of feeding its larvae in the early stages of development (CAMPOS et al., 2008). As such, bee pollen is a mixture of materials from different plant species containing pollen mixed with nectar and bee secretions that bind the grains together (FEÁS et al., 2012; KOMOSINSKA-VASSEV et al., 2015; BÁRBARA et al., 2015; ZHOU et al., 2015). Bees collect pollen from a large number of taxa, but only a few of these contribute significantly to their nutritional requirements (LIOLIOS et al., 2015).

The chemical composition of bee pollen depends largely on the plant source and geographical origin, as well as other factors such as climate, soil type, and bee species (KOMOSINSKA-VASSEV et al., 2015; BÁRBARA et al., 2015). Approximately 250 substances, including amino acids, lipids (triglycerides, phospholipids and fatty acids), phenolic compounds, vitamins, and flavonoid glycosides have been detected in bee pollen (KOMOSINSKA-VASSEV et al., 2015; BÁRBARA et al., 2015; ZHOU et al., 2015). Bee pollen also contains macronutrients (calcium, phosphorus, magnesium, sodium and potassium) and micronutrients (iron, copper, zinc, manganese, silicon and selenium) (KOMOSINSKA-VASSEV et al., 2015; ZHOU et al., 2015).

The therapeutic properties of bee pollen, such as antioxidant and antimicrobial activities, amongst other properties, have been attributed to the presence of flavonoids, phenolic acids and hydroxycinnamic acid amides (GABRIELE et al., 2015; CHEN et al., 2015; KOMOSINSKA-VASSEV et al., 2015; FEÁS et al., 2012).

Some authors have reported the presence of rutin as one of the flavonoids present in pollen. Markham and Campos (1996) pointed out that when identification was carried out using the correct procedures with NMR, the linkage between rhamnose and glucose was 1 ↔ 2 (neohesperidose), and not 1 ↔ 6 (rutinoside), as is evident in the rutin sugar moiety. From a nutritional standpoint this is not relevant, but for taxonomic and biosynthesis studies, it is a very important issue. This preponderance of flavonol 3-O-di- and tri-glycosides containing a 1,2 interglycosidic linkage is unlikely to be coincidence, especially as it encompasses such a wide range of taxa (MARKHAM; CAMPOS, 1996; CAMPOS et al., 2003; CAMPOS; MARKHAM, 2007).

Triacylated spermidines or HCA are predominant in pollen from oak species and in *Arabidopsis thaliana* pollen (HANDRICK et al., 2010), while diacylated spermidine conjugates are found in pollen from Betulaceae and Juglandaceae species (FELLENBERG et al., 2009).

The focus of this study was on the physicochemical and phytochemical analyses of *Apis mellifera* bee pollen collected in the municipality of Canavieiras, Bahia State-Brazil.

2 Material and methods

The bee pollen analysed in the current study was collected in an area with a high predominance of *Cocos nucifera* L. (Arecaceae) in the municipality of Canavieiras, located in Southern Bahia State-Brazil (15°41'S, 38°57'W). The area is comprised of a mangrove, large tracts of Atlantic forest, and also "restinga" and dune vegetation, with a predominance of palms such as *Cocos nucifera* L. (Arecaceae). *C. nucifera* blooms throughout the year, peaking between January and April (LORENZI, 2002). Other plant species found in this ecosystem are important polliniferous and nectariferous plant species. They also bloom throughout the year but peak between August and September (DÓREA et al., 2010). *C. nucifera* is traditionally cultivated for its coconut fruit.

2.1 Palynological analysis: identification of pollen taxa in the sample

The sample was prepared by the acetolysis method (ERDTMAN, 1960), and for the quantitative analysis, 1,500 pollen grain samples were used (VERGERON, 1964). The relative and average abundances and relative frequency of the pollen grains in the samples were calculated according to Louveaux et al. (1978).

2.2 Bee pollen sample

The *Apis mellifera* bee pollen sample (Natuflores – P-433) was provided by a consumer of the product. The bee pollen (200 g) was manually cleaned and oven-dried at 45 °C to constant weight (160 g). It was then ground to pass through a size 30 mesh and stored in sealed glass vials at 5 °C.

2.3 Physicochemical analyses

The physicochemical analyses, such as the moisture, protein, lipids, ash and sugar contents were carried out using previously described methodologies.

In order to determine the moisture content, 2.0 g of bee pollen was dried at 105 °C to constant weight to determine the water loss. The ash content was determined after incineration at 560 °C. The pH was measured after mixing 2.0 g of bee pollen with 5 ml of milli-Q

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water (BÁRBARA et al., 2015). The protein content was determined using the standard determination of %N by the Kjeldahl method, with a conversion factor of 6.25 or 5.6 (CAMPOS et al., 2008). The total sugars were determined using the dinitrosalicylic acid method (BÁRBARA et al., 2015).

The lipid content was determined by extraction with dichloromethane using a soxhlet reflux apparatus, as outlined by Archini et al. (2006). The total acidity was determined using 2.0 g of bee pollen mixed with 5.0 mL milli-Q water and titrating with 0.1 M NaOH 0.1 M to a pH of 8.5 (BÁRBARA et al., 2015).

2.4 Preparation of the non-polar and polar *Apis mellifera* bee pollen extracts

The pollen sample was powdered using liquid nitrogen with a mortar and pestle. Two extracts were prepared using this pollen sample: a non-polar extract using dichloromethane as the solvent; and a polar extract using a methanol/water mixture (1:1) as the solvent. The non-polar extract was prepared with 200 mg aliquots of bee pollen and dichloromethane (30 mL) as the solvent, refluxing in a Soxhlet, apparatus for 3 hours at approximately 60 °C. The yield was 30 mg of dichloromethane (non-polar) extract.

The polar extract was prepared with 200 mg of bee pollen macerated in methanol/water (30 mL, 1:1) for 24 hours. The yield was 70 mg of hydromethanolic (polar) extract. The dry extracts were maintained in amber flasks at 5 °C.

2.5 Analyses of the non-polar extract using Fourier Transform Infrared spectrometry (FTIR)

The basic functional groups of the *Apis mellifera* bee pollen non-polar extract were analysed qualitatively by the Fourier Transform Infrared method (ZIMMERMANN; KOHLER, 2014; HUCK, 2015). The FT-IR spectrum was recorded using a Bomem spectrometer at room temperature (ca. 25 °C). Triplicate 2 mg aliquots of the non-polar extract were homogenised with 100 mg of KBr powder and the mixture pressed into a thin pellet using a manual hydraulic press. The spectrum was obtained from the average of 32 scans in the range from 4000 to 400 cm^{-1} , with 5 cm^{-1} spectrum resolution and normalization at 1030 cm^{-1} . The influences of H_2O and CO_2 were subtracted automatically by the instrument.

2.6 Analysis of the non-polar extract using GC-EI-MS

A 1 μL aliquot of the non-polar extract, at a concentration of 3.33 mg/ml, was analysed using the Shimadzu GCMS-QP505A gas chromatograph equipped with a DB-5HT fused silica capillary column BPX5 (non-polar 5% phenyl polysilphenylene) (30 m x 0.25 mm internal

diameter x 0.25 μm film thickness) and an ion-trap mass detector.

Mass spectra were acquired in the electron-impact (EI) mode with an ionization voltage of 70 eV, between m/z 50 and 600. The oven was programmed as follows: the initial temperature of 100 °C was maintained for 5 min and then increased to 320 °C at the rate of 6 °C/min. The final temperature was maintained for 10 min. Helium was used as the carrier gas at a flow rate of 2.1 mL/min, ion source temperature of 300 °C; interface temperature of 300 °C; and scan speed of 2 scans s^{-1} .

Identification of the constituents, whenever possible, was carried out using fatty acid, hydrocarbon and steroid standards. The mass spectral data were also compared with the following computerized MS-databases: Wiley 275, Wiley 229 and the NIST 21 library at 95% confidence, and with data in the literature (KAFFARNIK et al., 2014; OLVERA-GARCIA et al., 2015). The concentrations of the constituents (relative contents) were obtained by calculating the percentage peak areas on the GC chromatograms.

2.7 Analyses of the polar extract using RPHPLC-DAD-ESI-MS/MS

Rutin (PubChem CID:5280805, Quercetin-3-O-Rhamnosyl-(1-2)Glucosyl), Quercetrin (PubChemCID:5280459, Quercetin – 3 – O – L-rhamnoside), Quercetin (PubChem CID:5280343), and Isorhamnetin (PubChem CID:5281654, 3'-Methoxyquercetin) were obtained from Sigma-Aldrich Chemical CO. (St. Louis, MO, USA) and used as the standards. The RPHPLC-DAD-ESI-MS/MS analysis was carried out using the Shimadzu DADSPD-M10AVP5 system equipped with a photodiode array detector coupled to Amazon Speed ETD, Bruker Daltonics (Shimadzu Corporation Kyoto, Japan).

The mass detector was a quadrupole ion trap equipped with an atmospheric pressure ionization source with an electrospray ionization interface, which was operated in the full scan MS/MS mode. All the operations, acquisitions and data analyses were controlled by the Shimadzu CBM-20A software. The polar extract (3.33 mg/ml) was filtered through a 0.45 μm PTFE filter and a 30.0- μL aliquot of the filtrate injected into the HPLC system. Spectral UV data was collected in the range from 240 to 400 nm, and the chromatograms were recorded at 360 nm and 270 nm.

The mixture of polar constituents was separated using a reverse phase Gemini Phenomenex – C18 RP-18 (Hewlett Packard) column (4.6 mm x250 mm, 5 μm) and a mobile phase composed of eluent A (0.1% aqueous. formic acid) and eluent B (methanol) at a constant flow rate of 1.0 mL min^{-1} with an oven temperature of 40 °C. The following elution programme was used: 0 min – (20% B); 10 min – (30% B), 20 min – (50% B); 30 min – (70% B); 40 min– (90% B); and 45 min – (40% B), then returning to

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the initial conditions (20% B) to re-equilibrate the column prior to the next run.

The following parameters were used for mass spectrometry: electrospray voltage of the ion source at -38 V, 4,000 V capillary voltage, end plate set at 500 V and capillary temperature of 300 °C. Helium was used as the collision gas and nitrogen as the nebulizing gas. Dissolution was facilitated using a counter-current nitrogen (dry gas) flow set at 9.0 litres/ minute. The spectra were acquired over a mass-to-charge (m/z) ranging between 100 and 1200 Da with resolution set at 30,000, based on an average of 10 scans.

The constituents were characterized from their mass spectral (MS) data in the negative and positive ionization modes and using on-line diode array ultraviolet visible spectroscopy, comparing the results with MS data reported in the following computer databases: SciFinder Scholar (<https://scifinder.cas.org>), Phenol-Explorer (www.phenol-explorer.eu), ChemSpider (<http://www.chemspider.com>), Knap Sack Core System Knap Sack Core System (<http://kanaya.naist.jp>) and HMDB (www.hmdb.ca) and also the methods reported by Negri et al. (2011), Yang et al. (2012), Khallouki et al. (2015) and Mihajlovic et al. (2015).

The flavonol-O-glycosides were hydrolysed in an acidic medium, according to the methodology described by Negri et al. (2012), in order to identify the aglycone. This was carried out using the standards quercetin and isorhamnetin from Sigma-Aldrich Chemical Company.

3 Results and discussion

The palynological analysis was able to identify *Cocos nucifera* as the major *taxon* in the bee pollen sample from Canavieiras (99.0%), which is explained by the fact that this pollen sample was collected in an area where this plant predominates from January to April, the period in which *C. nucifera* is in full bloom (LORENZI, 2002). The results differ from those of the bee pollen studied by Dórea et al. (2010) which included polliniferous plant species from the Atlantic Forest biome.

The physicochemical data obtained are shown in Table 1. In the physicochemical analysis, the sample showed moisture (24.4%, crude sample), ash (2.54%), protein (13.15%), total sugars (33.27%), and lipids (2.70%) contents, total acidity of (32.0 meq / kg) and pH of (4) (Table 1). The protein content was over 8%, which is in accordance with the Brazilian technical regulation (BRASIL, 2001). According to Komosinska-Vassev et al. (2015), pollen contains an average of 22.7% protein, including 10.4% of essential amino acids. In the sample from Canavieiras, the amount of protein was lower, which is unusual for bee pollen samples, since this product is a protein source for the hive. According to Liolios et al. (2015) pollens from

Table 1. Physicochemical analyses of a dried *Apis mellifera* bee pollen sample from the municipality of Canavieiras, Bahia State, Brazil.

Analyses	Contents	Acceptable level
Moisture	24.4%*	Maximum 30%
Ash	2.54%	Maximum 4% (m/m) dry basis
Protein	13.15%	Minimum 8% (m/m) dry basis
Total sugar	33.27%	14.5 to 55 (%) dry basis
Total acidity	32.0 meq/kg	Maximum 300 meq/kg
pH	4	4 to 6
Lipid	2.70%	Minimum 1.8% (m/m) dry basis

*non dried sample.

plants blooming in the spring have higher protein contents than those from the summer and autumn.

In the present sample, the protein content and total acidity were lower than the values reported by Bárbara et al. (2015), in which bee pollen samples, also from Bahia-Brazil, but obtained from another two regions: João Dourado and Uibaí, were analysed. This result is not surprising because these authors analysed bee pollen samples from other *taxa* and another bee species, *Melipona mandacaia* (Mandacaia stingless bee), not *Apis mellifera*.

Pollen generally contains 30.8% of digestible carbohydrates and approximately 25.7% of reducing sugars, mainly fructose and glucose (KOMOSINSKA-VASSEV et al., 2015). According to Carpes et al. (2009), the total sugar content in the bee pollen is associated with the amount of honey or nectar used by the bees to aggregate the pollen into pellets, allowing for transport to the hives.

The raw bee pollen is a hygroscopic product easily affected by environmental conditions (MARTINS et al., 2011). In the present sample, the moisture content (24.4%) was lower than the maximum limits established by the Brazilian technical regulation (BRASIL, 2001), which is 30% for fresh bee pollen.

The non-polar extract was analysed using Infrared (IR) spectroscopy and GC-EI-MS analyses. Infrared spectroscopy has been used in the analysis of natural products, since their biological properties are attributed to certain classes of compounds (ZIMMERMANN; KOHLER, 2014; HUCK, 2015; CAREDDA et al., 2016). Molecules containing the functional groups C-H, O-H, N-H, C=O, C-N and C-O are stimulated to stretch-, deform- and carry out scissor vibrations in the 4,000 to 400 cm⁻¹ spectral regions (MULARCZYK-OLIWA et al., 2012; ZIMMERMANN; KOHLER, 2014; CAREDDA et al., 2016).

The IR spectrum exhibited bands at 3369, 2924, 2852, 1742, 1464, 1377, 1241, 1171 and 1090 cm⁻¹, bands associated with different molecular groups. The broad band at 3369 cm⁻¹ was attributed to the presence of hydroxyl groups. The symmetrical and asymmetrical C-H stretching bands were at 2852 cm⁻¹ and 2924 cm⁻¹, respectively;

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the C-H bending bands at 1464 cm^{-1} and 1377 cm^{-1} ; the C=O stretching band of the carboxyl group was at 1742 cm^{-1} and the C-O stretching band of the carboxyl group at 1241 cm^{-1} , indicated the presence of fatty acids, hydrocarbons, phytosterols and esters containing long carbon chains. The strong vibration of the C-O single bond of the phytosterols was found in the fingerprint region (1171 cm^{-1} to 1090 cm^{-1}). Thus the bands exhibited an IR spectrum that indicated the presence of fatty acids, hydrocarbons and phytosterols, which was corroborated by the results obtained in the GC-EI-MS analysis.

The constituents detected in non-polar extract are shown in Table 2. Constituents 1-25 were detected in the non-polar extract by means of GC-EI-MS. Fatty acids (1, 3-5, 8, 11), hydrocarbons (6, 9, 13-16, 19), phytosterols (17, 18, 20 – 23), as well as the diterpene kaur-16-ene (2) and esters containing long carbon chains, such as the hexanoic acid undecyl ester (7), the hexanoic acid nonadecyl ester (10), the hexadecanoic acid octadecyl ester (24) and the octadecanoic acid octadecyl ester (25) (Table 2), were identified from their respective MS data, which were compared with MS data previously reported in the literature (KAFFARNIK et al., 2014; OLVERA-GARCIA et al., 2015).

The main constituents detected in the non-polar extract were fatty acids such as palmitic acid (1) (13.33%), oleic acid (4) (10.56%), behenic acid (8) (7.18%), hexanoic acid nonadecyl ester (10) (9.73%), and also hydrocarbons such as 9-heptacosene (15) (4.95%), 9-hentriacontene (16) (6.07%) and 13-tritriacontene (19) (8.21%). Phytosterols were detected in smaller amounts, and the most abundant were sitosterol (20) (3.24%) and fucosterol (21) (2.89%) (Table 2). Thus a predominance of saturated fatty acids was observed in the bee pollen sample under analysis. The most abundant n-alkanes detected in bee pollen samples were the odd-numbered carbon atom series C27, C29, C23, C17, and C21, while the sterol composition consisted mainly of β -sitosterol and stigmastan-3,5-diene (BASHIR et al., 2013).

According to Komosinska-Vassev et al. (2015), fatty acids were the main lipid constituents of bee pollen and constituted approximately 5.1%. Linoleic, γ -linoleic and arachidic acids constitute 0.4%, whereas phytosterols, especially sitosterol, constituted 1.1% (KOMOSINSKA-VASSEV et al., 2015). Palmitic, linoleic and α -linolenic acids were the main fatty acids found in bee pollen samples from *M. mandacai* collected in another two regions of Bahia State in Brazil (BÁRBARA et al., 2015). The dichloromethane extract from Greek bee pollen contained fatty acids and fatty acid esters (GRAIKOU et al., 2011).

On average, unsaturated fatty acids constitute about 70% of the total lipids (CAMPOS et al., 2008). There are generally greater amounts of unsaturated fatty acids in bee pollen than saturated fatty acids, with levels between 43% and 70% of total fatty acids (BASHIR et al., 2013;

BÁRBARA et al., 2015). According to Estevinho et al., (2012), the bees choose pollens with larger amounts of unsaturated fatty acids because these are more adequate for their metabolism. The hypoglycemic activity of the pollen was mainly attributed to the presence of unsaturated fatty acids, phospholipids and phytosterols (KOMOSINSKA-VASSEV et al., 2015).

The polar extract of the bee pollen sample evaluated here was analysed by means of HPLC-DAD-ESI-MS/MS. The constituents, shown in Table 3, were identified based on their UV spectra and MS data, which were compared with previously reported MS data for bee pollen constituents. Many bee pollen samples exhibited high contents of hydroxycinnamic acid amide derivatives or triacylated spermidines, and flavonol glycosides.

Thus the MS data for the flavonol glycosides and hydroxycinnamic acid amides detected in this polar extract of the bee pollen were compared with the MS data reported by Sobolev et al. (2008), Fellenberg et al. (2009), Freire et al. (2012), Goupy et al. (2013), Khallouki et al. (2015) and Mihajlovic et al. (2015).

Compounds 28-31, 33 and 34 were characterized as flavonol glycosides according to the typical maximum UV absorption at 255 – 355 nm (Table 3). Flavonoids in the form of aglycones are rarely found in pollens and in plants, since they are generally linked to sugars. Flavonol-*O*-glycosides with a free hydroxyl group in the C3-*O* position exhibit maximum UV band I at 374 nm, whereas flavonols with a blocked hydroxyl at the C3-*O* position exhibit maximum UV band I between 340 nm to 360 nm (CAMPOS; MARKHAM, 2007; RODRIGUEZ-PÉREZ et al., 2013; SIMIRGIOTIS et al., 2013). Band I at 355 nm indicated that the sugar substitution occurred on the hydroxyl group at the C3-*O* position of flavonols 28-31, 33 and 34 (CAMPOS; MARKHAM, 2007). Quercetin and isorhamnetin were identified as aglycones in the flavonol-3-*O*-glycosides 28-31, 33 and 34 after hydrolysis, which was confirmed by matching the retention time, the UV data, the MS spectral data and the co-injection with standards (data not shown).

Compound 28 exhibited a $[M - H]^-$ ion at m/z 609 in the ESI-MS spectrum and the ion m/z 301 in the MS/MS spectrum, attributed to deprotonated quercetin as the aglycone. Compound 28 was identified as quercetin-3-*O*-rhamnosylglucoside. Compound 29 exhibited a $[M - H]^-$ ion at m/z 639 and the ion m/z 477 in the MS/MS spectrum, which was formed by the loss of the glucose moiety (162 Da). The *O*-glycosidic bond is more labile at position 7, and consequently its breakdown is favoured (GOUPY et al., 2013). The base peak at m/z 315 corresponds to deprotonated isorhamnetin (3'-*O*-methyl quercetin), formed after the loss of the second glucose unit. These MS data agree with those reported by Goupy et al. (2013), and compound **29** was assigned as isorhamnetin-di-3,7-*O*-glucoside.

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Table 2. Constituents found in the analysis of the *Apis mellifera* bee pollen non-polar extract using GC-IE-MS.

N°	Tr Min.	IE - [M] ⁺ m/z	Main mass fragments		Identification
			m/z	m/z	
1	24.61	256 (13) C ₁₆ H ₃₂ O ₂	213 (14), 185 (11), 171 (11), 157 (13), 129 (34), 115 (13), 97 (18), 87 (21), 73 (100)		Palmitic acid - (n-hexadecanoic acid) ^a
2	25.87	272 (26) C ₂₀ H ₃₂	257 (54), 229 (47), 213 (40), 201 (11), 187 (21), 175 (25), 161 (23), 147 (51), 133 (36), 125 (64), 119 (55), 109 (42), 105 (84)		Kaur-16-ene ^b
3	27.86	280 (3) C ₁₈ H ₃₂ O ₂	151 (3), 137 (4), 123 (8), 110 (12), 107 (12), 95 (51), 93 (28)		Linoleic acid - (9,12-octadecenoic acid) ^a
4	27.99	282 (1) C ₁₈ H ₃₄ O ₂	264 (4), 139 (3), 125 (5), 121 (7), 111 (11), 98 (21), 95 (30)		Oleic acid - (9(Z)-octadecenoic acid) ^a
5	28.33	284 (21) C ₁₈ H ₃₆ O ₂	241 (13), 185 (18), 129 (33), 97 (25), 83 (28), 73 (100)		Stearic acid - (n-octadecanoic acid) ^a
6	34.02	352 (1) C ₂₅ H ₅₂	127 (5), 113 (7), 99 (13), 85 (40), 71 (57)		n-pentacosane ^a
7	34.24	270 (3) C ₁₇ H ₃₄ O ₂	239 (24), 134 (34), 129 (12), 112 (24), 98 (100)		hexanoic acid undecyl ester ^b
8	35.25	340 (26) C ₂₂ H ₄₄ O ₂	297 (10), 185 (13), 129 (31), 97 (25), 83 (29), 73 (100)		Behenic acid - (docosanoic acid) ^a
9	37.19	380 (1) C ₂₇ H ₅₆	130 (26), 117 (27), 111 (7), 99 (14), 97 (13), 85 (38), 71 (58), 57 (100)		n-heptacosane ^a
10	37.64	382 (2) C ₂₅ H ₅₀ O ₂	285 (6), 267 (24), 134 (28), 112 (26), 98 (100)		hexanoic acid nonadecyl ester ^b
11	38.32	368 (25) C ₂₄ H ₄₈ O ₂	325 (7), 185 (10), 129 (30), 97 (25), 85 (30), 73 (100)		tetracosanoic acid ^a
12	38.80	410 (1) C ₃₀ H ₅₀	121 (10), 95 (15), 81 (55), 69 (100)		squalene ^b
13	39.85	350 (1) C ₂₅ H ₅₀	139 (8), 125 (17), 111 (37), 97 (68), 83 (90)		9-pentacosene ^a
14	40.08	408 (1) C ₂₉ H ₆₀	99 (13), 97 (11), 85 (38), 83 (11)		n-nonacosane ^a

^a Compounds identified by comparing the electron impact (EI) mass spectra and the retention times with those of the corresponding pure standards; ^b Compounds identified based on the comparison between the electron impact (EI) mass spectra, the mass spectral data reported in the GC-MS computer databases (Wiley 275, Wiley 229 and NIST 21), and the mass spectral data reported by Kafarnik et al. (2014) and Olvera-Garcia et al. (2015).

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Table 2. Continued...

N°	Tr Min.	IE - [M] ⁺ m/z	Main mass fragments m/z	Identification
15	40.18	378 (1) C ₂₇ H ₅₄	139 (8), 125 (21), 111 (44), 97 (100), 85 (21)	9-heptacosene ^a
16	42.54	434 (2) C ₃₁ H ₆₂	139 (9), 125 (21), 111 (37), 97 (100)	9-hentriacontene ^a
17	44.25	398 (5) C ₂₈ H ₄₆ O	383 (4), 365 (4), 314 (55), 299 (21), 281 (31), 271 (28), 229 (22), 173 (10), 161 (13), 159 (19), 145 (28), 133 (29), 105 (43)	Ergosta-5,24(28)-dien-3-ol ^b
18	44.40	400 (21) C ₂₈ H ₄₈ O	385 (9), 382 (10), 367 (10), 315 (21), 281 (15), 213 (16), 163 (20), 159 (21), 147 (27), 145 (30), 133 (28), 121 (25), 117 (45), 109 (29), 107 (38),	Campesterol - (ergost-5-en-3-ol) ^a
19	45.16	462 (2) C ₃₃ H ₆₆	153 (5), 139 (11), 125 (22), 111 (36), 97 (100)	13-tritriacontene ^a
20	45.54	414 (40) C ₂₉ H ₅₀ O	399 (15), 396 (21), 381 (15), 329 (35), 303 (26), 163 (28), 157 (14), 143 (23), 149 (18), 145 (42), 133 (36), 119 (34), 107 (50), 105 (45), 95 (90)	sitosterol - (stigmast-5-en-3-ol) ^a
21	45.72	412 (2) C ₂₉ H ₄₈ O	314 (100), 299 (17), 281 (34), 229 (21), 211 (13), 145 (17), 133 (17), 117 (22), 105 (20)	Fucoesterol - (5,24(28)-Stigmastadien-3β-ol) ^b
22	46.33	410 (11) C ₂₉ H ₄₆ O	367 (8), 297 (10), 281 (14), 269 (17), 245 (10), 207 (30), 147 (20), 133 (21), 123 (17), 117 (23), 109 (17), 105 (18)	5,24(28)-Stigmastadien-3-one ^b
23	47.09	412 (14) C ₂₉ H ₄₈ O	397 (4), 289 (14), 281 (14), 229 (34), 218 (33), 147 (26), 135 (30), 124 (100), 109 (30)	Stigmast-4-en-3-one ^b
24	48.56	508 (6) C ₃₄ H ₆₈ O ₂	281 (15), 257 (30), 207 (33), 111 (20), 109 (10), 97 (90)	hexadecanoic acid octadecyl ester - (stearyl palmitate) ^b
25	51.10	536 C ₃₆ H ₇₂ O ₂	281 (25), 264 (15), 111 (21), 109 (20), 97 (80)	octadecanoic acid octadecyl ester - (stearyl stearate) ^b

^a Compounds identified by comparing the electron impact (EI) mass spectra and the retention times with those of the corresponding pure standards; ^b Compounds identified based on the comparison between the electron impact (EI) mass spectra, the mass spectral data reported in the GC-MS computer databases (Wiley 275, Wiley 229 and NIST 21), and the mass spectral data reported by Kaffarnik et al. (2014) and Olvera-Garcia et al. (2015).

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Table 3. Data of constituents 26-41 including the retention times, UV-vis absorptions, molecular ion peaks and fragmentation patterns in both the negative and positive ionization modes obtained from the MS and HPLC profiles.

No	Tr Min.	HPLC/DAD UV nm	HPLC/(+)ESI-MS/MS m/z (%base peak)	HPLC/(-)ESI-MS/MS m/z (%base peak)	Proposed structure
26	5.9	300		[M - H] ⁻ - 341 MS/MS - 179 (100), 161 (50)	6-O-caffeoyl glucoside
27	6.5			[M - H] ⁻ - 195 [M - H] ⁻ - 609 MS/MS - 300	trihydroxycinnamic acid quercetin-3-O-rhamnosylglucoside
28	22.0	260-355		[M - H] ⁻ - 639 MS/MS - 477 (50), 315 (100)	isorhamnetin-di-3,7-O-glucoside ^a
29	22.6	260-355		[M - H] ⁻ - 639 MS/MS - 477 (50), 315 (100)	isorhamnetin-di-3,7-O-glucoside ^a
30	23.1	260-355	[M + H] ⁺ - 771 MS/MS - 625 (90), 479 (50), 317 (100) [M + Na] ⁺ - 647	[M - H] ⁻ - 769 MS/MS - 623 (50), 605 (90), 315 (100)	isorhamnetin-3-O-(2'',3''-O-dirhamnosyl) glucoside ^a
31	24.6	260-355	MS/MS - 501 (30), 331 (100) [M + H] ⁺ - 625 [M + Na] ⁺ - 654	[M - H] ⁻ - 623 MS/MS - 459 (60), 314 (100), 299 (40)	isorhamnetin-3-O-(2''-O-rhamnosyl) glucoside ^a
32	27.4	330	MS/MS - 479 (20), 317 (100) [M + Na] ⁺ - 654 MS/MS - 492 (100), 330 (50) [M + H] ⁺ - 632 MS/MS - 470 (100), 452 (20)	[M - H] ⁻ - 630 MS/MS - 494 (80), 468 (100), 358 (40), 332 (35)	N',N''-tris-caffeoyl spermidine ^a
33	26.8	260-355		[M - H] ⁻ - 447 MS/MS - 301	quercetin-3-O-rhamnoside - (quercetin) ^b
34	27.4			[M - H] ⁻ - 665 MS/MS - 623 (40), 501 (50), 314 (100), 299 (60)	isorhamnetin-3-O-(2''-O-rhamnosyl acetyl) glucoside ^a
35	27.8	330	[M + Na] ⁺ - 638 MS/MS - 476 (100), 314 (20) [M + H] ⁺ - 616 MS/MS - 454 (100), 436 (18)	[M - H] ⁻ - 614 MS/MS - 478 (100), 452 (80), 358 (70)	N',N''-dicafeoyl,N'''-coumaroyl spermidine ^a
36	28.7	330	[M + H] ⁺ - 646 MS/MS - 484 (50), 470 (50), 452 (100)	[M - H] ⁻ - 644 MS/MS - 508 (100), 482 (30), 468 (30)	N',N''-dicafeoyl,N'''-feruloyl spermidine ^a

^a Identification based on both spectroscopic and mass spectrometric data and from the comparison with MS spectral data and the methods reported by Sobolev et al. (2008), Fellenberg et al. (2009), Negri et al. (2011), Freire et al. (2012), Goupy et al. (2013), Khalilouki et al. (2015) and Minajovic et al. (2015). The aglycones of compounds **28-31**, **33** and **34** were identified using quercetin and isorhamnetin standards from Sigma-Aldrich Chemical CO; ^b quercetin was also identified using standards.

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Table 3. Continued...

No	Tr Min.	HPLC/DAD UV nm	HPLC/(+)ESI-MS/MS m/z (%base peak)	HPLC/ (-)ESI-MS/MS m/z (%base peak)	Proposed structure
37	29.6	330	[M + H] ⁺ - 630 MS/MS - 468 (30), 454 (50), 436 (100)	[M - H] ⁻ - 628 MS/MS - 508 (50), 492 (100), 452 (30), 372 (40), 358 (20)	N'-caffeoyl-N''-feruloyl,N'''-coumaroyl spermidine ^a
38	30.2	330	[M + H] ⁺ - 600 MS/MS - 454 (10), 438 (100), 420 (30) [M + Na] ⁺ - 606	[M - H] ⁻ - 598 MS/MS - 478 (100), 358 (60)	N'-caffeoyl-N''-N'''-dicoumaroylspermidine ^a
39	31.0	330	MS/MS - 460 (10) [M + H] ⁺ - 584 MS/MS - 438 (80), 420 (100) [M + Na] ⁺ - 647	[M - H] ⁻ - 582 MS/MS - 462 (100), 342 (50)	N',N'',N'''-tris-p-coumaroyl spermidine ^a
40	31.6	330	MS/MS - 331 [M + H] ⁺ - 625 MS/MS - 317 (100), 309 (70) [M + H] ⁺ - 674	[M - H] ⁻ - 623 MS/MS - 315 (100), 300 (50)	isorhamnetin-3-O-(6''-O-p-coumaroyl)-glucoside ^a
41	32.0	330	MS/MS - 498 (60), 480 (100)	[M - H] ⁻ - 672 MS/MS - 536 (40), 522 (100)	N',N'',N'''-tris-p-feruloyl spermidine ^a

^a Identification based on both spectroscopic and mass spectrometric data and from the comparison with MS spectral data and the methods reported by Sobolev et al. (2008), Fellenberg et al. (2009), Negri et al. (2011), Freire et al. (2012), Goupy et al. (2013), Khalouki et al. (2015) and Mihajlovic et al. (2015). The aglycones of compounds **28-31**, **33** and **34** were identified using quercetin and isorhamnetin standards from Sigma-Aldrich Chemical CO; ^b quercetin was also identified using standards.

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Isorhamnetin-3-*O*-(2'',3''-*O*-dirhamnosyl) glucoside and isorhamnetin-3-*O*-(2''-*O*-rhamnosyl) glucoside have been previously reported in bee pollen (NEGRI et al., 2011). Compound 30 exhibited a $[M - H]^-$ ion at m/z 769, which, after the MS/MS experiments produced an ion fragment at m/z 623, indicating the loss of the rhamnose moiety (146 Da). It also produced an ion fragment at m/z 605 $[M - H - 164]^-$, which was attributed to the loss of rhamnose and water moieties, characteristic of *O*-glycosylation at the hydroxyl group in position 2'' of the glucose in *C*-glycosyl flavonols-*O*-glycosylated (FERRERES et al., 2010; DINELLI et al., 2011). The base peak at m/z 315 corresponded to deprotonated isorhamnetin, which was formed after the loss of the glucosyl unit. Moreover, compound 30 exhibited a $[M + H]^+$ ion at m/z 771, which produced ion fragments at m/z 625 and at m/z 479, resulting from the loss of two rhamnose moieties, and it also produced a base peak at m/z 317 (protonated isorhamnetin). These MS spectral data agree with those reported by Negri et al. (2011), and compound 30 was assigned as isorhamnetin-3-*O*-(2'',3''-*O*-dirhamnosyl) glucoside.

Compound 31 was the main constituent detected in the polar extract and showed molecular ion peaks at m/z 623 and m/z 625 and a sodium adduct at m/z 647. In the negative ionization mode, the MS/MS experiments produced an ion fragment at m/z 459, resulting from the loss of rhamnose and water moieties. The ion fragments at m/z 314 (100) and 299 (40) indicated isorhamnetin as the aglycone. According to Rodríguez-Pérez et al. (2013) and Simirgiotis et al. (2013), the ion fragment at m/z 314 was produced by the homolytic cleavage of the *O*-glycosidic bond at the C3-*O* position of flavonol, and resulted in the formation of a aglycone anion radical of isorhamnetin. On the other hand, the MS/MS experiments in the positive ionization mode produced ion fragments at m/z 479 and a base peak at m/z 317 (protonated isorhamnetin), corresponding to the loss of rhamnose and glucose, respectively (KHALLOUKI et al., 2015). Protonated molecules are able to form adducts, clusters and/or molecular complexes with the mobile phase, such as the sodium adduct $[M + Na]^+$ at 22 Da instead of the proposed protonated molecule in the positive ionization mode. The sodium adduct exhibited $[M + H + Na]^+$ at m/z 647, which, after the MS/MS experiments, yielded a base peak at m/z 331, representing the protonated rhamnosyl glucoside moiety plus sodium. These MS spectral data agree with those reported by Negri et al. (2011), and compound 31 was assigned as isorhamnetin-3-*O*-(2''-*O*-rhamnosyl) glucoside.

Compound 40 also exhibited molecular ion peaks at m/z 623 and m/z 625 and a sodium adduct at m/z 647, as did compound 31; however it exhibited maximum UV absorption at 330 nm. Flavonoid glycosides esterified with aromatic acids exhibit longer R_t on the RP-HPLC column

and their UV spectra exhibit intense band I (approximately 330 nm) and a small band II at 270 nm, resulting from the UV absorption overlap of the flavonoid and the cinnamoyl groups. The predominant bonding site of the acyl groups was usually the 6''-position of hexose; however, other positions should not be excluded (ABAD-GARCÍA et al., 2009). The MS/MS experiments in the negative ionization mode produced a base peak at m/z 315 (deprotonated isorhamnetin), resulting from the loss of 308 Da (*p*-coumaroyl-glucoside moiety). The MS/MS experiments in the positive ionization mode exhibited a base peak at m/z 317 (protonated isorhamnetin) and an ion fragment at m/z 309, attributed to the protonated *p*-coumaroyl-glucoside moiety. The sodium adduct provided a molecular ion peak at m/z 647, which fragmented to yield a base peak at m/z 331, representing the protonated *p*-coumaroyl-glucoside moiety plus sodium. Compound 40 was tentatively assigned as isorhamnetin-3-*O*-(6''-*O*-*p*-coumaroyl)-glucoside, from its UV and mass spectral fragmentation patterns.

Compound 34 exhibited a $[M - H]^-$ ion at m/z 665, which after the MS/MS experiments exhibited ion fragments at m/z 623 and m/z 501, along with ion fragments attributed to isorhamnetin at m/z 314 and m/z 299. Compound 34 had another 42 Da indicating the presence of an acetyl group, which was also observed from the ion fragment at m/z 623. Acylated flavonol glycosides eluted after their corresponding flavonol glycosides under reversed phase conditions (RODRÍGUEZ-PÉREZ et al., 2013). Based on its mass spectral data, compound 34 was tentatively assigned as isorhamnetin-3-*O*-(2''-*O*-rhamnosyl acetyl) glucoside. Compound 33 exhibited a $[M - H]^-$ ion at m/z 447, which after the MS/MS experiments produced a base peak at m/z 301, indicating quercetin as the aglycone. Compound 33 was identified as quercetin-3-*O*-rhamnoside, also known as quercetrin, based on an authentic standard.

Compounds 32, 35-39 and 41 were characterized as hydroxycinnamic acid amide (HCA) derivatives, according to the typical maximum UV absorption at 330 nm (Table 3). Hydroxycinnamic acid amides (HCA) may undergo isomerization of their double bonds, and yield *E/Z* isomers. In addition, they may undergo positional isomerism and present complex fragmentation behaviours (YANG et al., 2012). MS/MS experiments with protonated HCA produce abundant ion fragments attributed to neutral acyl losses; for example, 176 Da for feruloyl, 162 Da for caffeoyl, and 146 Da for the coumaroyl moieties, followed by neutral water loss (NEGRI et al., 2011). MS/MS experiments with deprotonated HCA produce abundant ion fragments attributed to the loss of 120 Da, corresponding to the coumaric acid moiety; to the loss of 136 Da corresponding to the caffeic acid moiety; and to the loss of 150 Da corresponding to the ferulic acid moiety (YANG et al., 2012; MIHAJLOVIC et al., 2015).

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Compounds 32 and 35 exhibited a $[M + H]^+$ ion at m/z 632 and m/z 616, respectively. Their MS/MS fragmentation patterns were similar in both the positive and negative ionization modes and exhibited base peaks derived from neutral caffeoyl losses at m/z 470 and m/z 454 and ion fragments derived from water loss at m/z 452 and m/z 436, respectively, in the positive ionization mode. The sodium adduct at m/z 654 (for 32) and at m/z 638 (for 35) produced ion fragments derived from two neutral caffeoyl losses at m/z 492 and m/z 330 and at m/z 476 and m/z 314, respectively. Compounds 32 and 35 exhibited $[M - H]^-$ ions at m/z 630 and m/z 614, respectively. Their MS/MS fragmentation produced an ion fragment derived from neutral caffeoyl losses at m/z 468 and m/z 452, corresponding to the loss of two caffeic acid moieties (136 Da) at m/z 494 and m/z 358 and at m/z 478 and m/z 358, respectively. Based on the mass spectral data and the method reported by Negri et al. (2011), Yang et al. (2012), and Mihajlovic et al. (2015), compound 32 was assigned as N',N'',N'''-tris-caffeoyl spermidine and compound 35 as N',N''-dicaffeoyl,N'''-coumaroyl spermidine.

Compounds 36 and 37 exhibited molecular ion peaks at m/z 646 and m/z 630, respectively. MS/MS experiments in the positive ionization mode, produced ion fragments derived from neutral caffeoyl losses at m/z 484 and m/z 468, derived from neutral feruloyl losses at m/z 470 and m/z 454 and base peaks derived from water loss at m/z 452 and m/z 436, respectively. In the negative ionization mode, the fragmentation patterns for compounds 36 and 37 were different. Compound 36 exhibited a $[M - H]^-$ ion at m/z 644, which produced an ion fragment derived from neutral caffeoyl loss at m/z 482, at m/z 468, derived from neutral feruloyl losses at m/z 468 and a base peak at m/z 508, resulting from the loss of a caffeic acid moiety. On the other hand, compound 37 exhibited a $[M - H]^-$ ion at m/z 628, which produced an ion fragment derived from neutral feruloyl loss at m/z 452, corresponding to the loss of a coumaric acid moiety at m/z 508, and a base peak at m/z 492, corresponding to the loss of a caffeic acid moiety. Based on the MS data and the method reported by Negri et al. (2011), Yang et al. (2012), and Mihajlovic et al. (2015), compound 36 was assigned as N',N''-dicaffeoyl,N'''-feruloyl spermidine and compound 37 as N'-caffeoyl-N''-feruloyl, N'''-coumaroyl spermidine.

Compound 38 showed a $[M + H]^+$ ion at m/z 600, which produced an ion fragment at m/z 454 derived from neutral coumaroyl losses and a base peak at m/z 438, derived from neutral caffeoyl losses, as well as an ion fragment at m/z 420. Besides this, compound 38 exhibited a $[M - H]^-$ ion at m/z 598, which produced a base peak at m/z 478 and an ion fragment at m/z 358, both corresponding to the loss of a coumaric acid moiety. These MS data agree with those reported by Mihajlovic et al. (2015) and compound 38 was assigned as N'-caffeoyl-N'',N'''-dicoumaroyl spermidine.

Compound 39 exhibited a $[M + H]^+$ ion at m/z 584, which produced an ion fragment derived from neutral coumaroyl losses at m/z 438, and a base peak at m/z 420. The MS/MS experiments on the sodium adduct at m/z 606 produced a base peak derived from neutral coumaroyl losses at m/z 460. Besides this, compound 39 showed a $[M - H]^-$ ion peak at m/z 582, which produced a base peak at m/z 462 and an ion fragment at m/z 342, both corresponding to the loss of a coumaric acid moiety. Since the ion fragments of compound 39 matched those reported by Negri et al. (2011) and Mihajlovic et al. (2015), it was assigned as N',N'',N'''-tris-*p*-coumaroyl spermidine.

Compound 41 exhibited a $[M + H]^+$ ion at m/z 674, which produced an ion fragment derived from neutral feruloyl losses at m/z 498, and a base peak at m/z 480. Besides this, compound 41 showed a $[M - H]^-$ ion at m/z 672, that produced an ion fragment corresponding to the loss of a ferulic acid moiety at m/z 536 and a base peak at m/z 522. These MS spectral data agree with those reported by Negri et al. (2011), and compound 41 was assigned as N',N'',N'''-tris-*p*-feruloyl spermidine. Tables 1 - Physicochemical analyses of a dried *Apis mellifera* bee pollen sample; 2 - Constituents detected in the analysis of the non-polar extract using GC-IE-MS and 3 - Constituents detected in the analysis of the polar extracts in both the negative and positive ionization modes obtained from the MS and HPLC profiles, together with the mass spectra of the constituents detected in this bee pollen sample (Supplementary Material, Figures 1S-48S) were provided as supporting information. The structures of these compounds will require confirmation in future NMR experiments when sufficient pure material is available.

Of the compounds present in the pollen, the phenolic acids and polyphenolic compounds such as flavonoids are a very important group of bioactive structures, useful in a vast range of applications. One point four percent (1.4%) are flavonoids occurring mainly as glycosides of kaempferol, quercetin, and isorhamnetin, whereas 0.2% of the phenolic acids correspond mainly to chlorogenic acid (KOMOSINSKA-VASSEV et al., 2015) and caffeic acid derivatives (CAMPOS et al., 1997, 2003). In *Typha angustifolia* L. pollen, the main active compounds found were flavonoids (TAO et al., 2011). High contents of flavonol glycosides and aglycones were found in the bee pollen produced in Bahia State, Brazil (FREIRE et al., 2012). Quercetin and kaempferol glycosides were detected in many taxa, for example: *Eucalyptus globulus*, *Salix atrocinera*, *Helianthus annuus*, *Raphanus raphanistrum*, *Erica australis* and *Cistus angustifolia*. The taxa originated from different countries ranging from Portugal to New Zealand (CAMPOS et al., 1997). They were also detected in the rape bee pollen collected from the Qinghai-Tibetan Plateau (LV et al., 2015). Benzoic acid derivatives and hydroxycinnamic acids have also been identified in pollen

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(GRAIKOU et al., 2011). Castanea pollen exhibited high polyphenol, flavonoid and anthocyanin contents, while a high content of flavonols was detected in Cistus pollen (GABRIELE et al., 2015).

Spermidine derivatives are compounds not always identified in pollen samples and give extra information that helps to compliment research into structures involved in the bioactivities explored in this raw material. In *Ambrosia artemisiifolia* L. pollen, flavonoids were detected as monoglycosides and malonyl-mono- and diglycosides of isorhamnetin, quercetin and kaempferol, whereas the spermidine derivatives were identified as the dominant polyamides (MIHAJLOVIC et al., 2015). N',N'',N'''-Tris-p-feruloyl spermidine and N',N'',N'''-tris-p-coumaroyl spermidine were found in seven bee pollen samples from different palynological sources harvested in the municipality of Pindamonhangaba (South-eastern Brazil) (NEGRI et al., 2011).

Fatty acids and esters, phytosterols, flavonol glycosides and triacylated spermidines exhibit important biological activities such as antioxidant activity, free radical scavenging capacity, coronary heart disease prevention, as well as hepatoprotective, anti-inflammatory, anticancer and antimicrobial activities, amongst others. Flavonoids acting as antioxidant compounds can prevent oxidative stress, which leads to disorders that affect the central nervous system. In addition, flavonoids can also modulate both enzyme and receptor activities (GROSSO et al., 2013). Quercetin and isorhamnetin derivatives exhibit free radical scavenging capacity, coronary heart disease prevention, and hepatoprotective, anti-inflammatory, anticancer (KUMAR; PANDEY, 2013), antioxidant and antiviral (FRIEDMAN, 2014) activities.

Spermidine is an endogenous polyamine with a polycationic structure found in the central nervous system of mammals and it regulates biological processes such as the Ca²⁺ influx by the glutamatergic N-methyl-D-aspartate receptor (CARVALHO et al., 2012). Several important control functions in cells, ranging from the basic DNA synthesis to the regulation of cell proliferation and differentiation, were attributed to polyamines such as spermidine and spermine and their precursor putrescine (MINOIS et al., 2011; CHOI; PARK, 2012). Spermidine derivatives are also able to improve locomotor performance in the aging (MINOIS, 2014; MINOIS et al., 2014). In addition they exhibited antimicrobial activity against viruses, bacteria and fungi, and have been related to protection against pathogens (MIHAJLOVIC et al., 2015).

4 Conclusion

Bee pollen has been used for many years as a beneficial dietary supplement. Fatty acids and esters were the main constituents found in the dichloromethane extract of the bee pollen sample collected in the municipality of

Canavieiras, in an area with the predominance of *C. nucifera*. They were followed by smaller amounts of phytosterols and hydrocarbons, whereas large amounts of flavonol glycosides and triacylated spermidines were found in the polar extracts. Flavonoids and di- and triacylated spermidines were reported as the main constituents found in many other pollen and bee pollen samples.

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Supplementary Material

The following online material is available for this article:

Figures 1S-48S

This material is available as part of the online article from <https://www.scielo.br/bjft>

Erratum

In the article "**Phytochemical analysis and botanical origin of *Apis mellifera* bee pollen from the municipality of Canavieiras, Bahia State, Brazil**", DOI <http://dx.doi.org/10.1590/1981-6723.17616>, published in *Brazilian Journal of Food Technology*, vol. 21, 16 pp., on page **1**,

where it reads:

Giuseppina Negri^{1*}, Lidia Maria Ruv Carelli Barreto², Fábila Lugli Sper³, Claudemir de Carvalho³,
Maria da Graça Rodrigues Campos⁴

It should be read:

Giuseppina Negri^{1*}, Lidia Maria Ruv Carelli Barreto², Fábila Lugli Sper³, Claudemir de Carvalho³,
Maria da Graça Ribeiro Campos⁴

