



Full Length Article

Antifungal Activity of Root Extracts from *Baccharis salicina* on Germination of Uredospores of *Hemileia vastatrix*

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Received 04 November 2020; Accepted 25 January 2021; Published 16 April 2021

Abstract

Hemileia vastatrix is a fungus associated with coffee leaf rust, the most destructive disease of *Coffea arabica*. The objective of this work was to evaluate the antifungal activity of alcoholic extracts from roots of *Baccharis salicina* and to determine the metabolites present in these fractions. Antifungal activity was evaluated under *in vitro* conditions by monitoring the germination ability of *H. vastatrix*, the coffee leaf rust pathogen. In order to determine the presence of metabolites, chemical characterization of fractions obtained from methanolic root extracts was performed with help of an untargeted metabolomic approach and by using high-resolution mass spectrometry (MS) and MS² based on direct-injection electrospray mass spectrometry (DIESI-MS) and hierarchical cluster analysis (HCA). Germination percentage was evaluated by leaves fixation technique. The MEBs significantly decreased the percentage of germination of *H. vastatrix* to levels below 5% as the dose increased. The multivariable analysis confirmed that the distribution of three fractions of methanolic extracts belonged to polyketides, organoheterocyclic compounds, fatty acyls, prenol-lipids, organo-oxygen and farnesene classes. This report comprises the first study of the metabolomic profile and biological activity study of roots from *B. salicina* against coffee leaf rust pathogen *H. vastatrix*. © 2021 Friends Science Publishers

Keywords: Antifungal activity; MS-MS analysis; *Hemileia vastatrix*; Multivariate analysis; *Baccharis salicina*

Introduction

Asteraceae is regarded as the largest family of widely distributed flowering plants in the world, being absent only in Antarctica. This family contains mostly herbaceous plants and small shrubs and rarely trees, with an estimated 23,600 species belonging to 1600 genera with 12 subfamilies and 28 tribes (Ramos *et al.* 2016). The Astereae tribe includes the subtribe Baccharidinae Less., which is exclusively on the American continent and includes the genera *Archibaccharis* and *Baccharis*. *Baccharis* plants are usually studied due to their biological activities such as antibacterial, antifungal, antiprotozoal, antiviral, antioxidant, antidiabetic, immunomodulatory, antimutagenic and chemopreventive properties (Abad and Bermejo 2007).

In Southern México, *Baccharis salicina*, is known as chilcanka' or chilca in the Tzotzil language. In this region, it has been used as a traditional medicinal plant. The locals use infusions from these species' leaves to treat gynecological and digestive disorders (Abad and Bermejo 2007; Ramos *et al.* 2016). Methanolic extracts of its leaves contain antifungal properties against phytopathogens, but few reports regarding the chemical composition of the plant are available. It has been reported that methanolic extracts of its leaves contain antifungal properties against phytopathogens. In this regard, significant effects on the growth of mycotoxigenic fungi have been found (Tequida-Meneses *et al.* 2002; Rosas-Burgos *et al.* 2011). Likewise, the methanolic extract from roots has reportedly inhibited sporulation and mycelial growth of *Aspergillus ochraceus* and *Fusarium moniliforme*.

This was reported for the first time in a chemical composition study by GC-MS (Lam-Gutiérrez *et al.* 2019). Thus, due to the importance of this genus, phytochemical studies and the search for new active molecules of more *Baccharis* species offer promising outcomes.

Coffee leaf rust, the most destructive disease in the cultivation of coffee worldwide, is caused by the fungus *H. vastatrix*, mainly affecting *C. arabica*, which represents an estimated 70% of the production worldwide (Talinhas *et al.* 2016). *H. vastatrix* is an obligate biotrophic fungus, penetrating and sporulating through stomata (López-Bravo *et al.* 2012). The need to have less toxic and more environment friendly fungicides is a challenge when it comes to searching novel antifungal agents (Naqvi *et al.* 2019; Javaid *et al.* 2020). Nowadays, the finding relies mainly on ethnobotanical information and exploration of new bioactive compounds obtained from native plants (Yoon *et al.* 2013; Akhtar *et al.* 2020), which are increasingly being studied in a holistic way. Thus, omics approaches are being used to provide qualitative and quantitative descriptions. Metabolomics is a well-established research field that is developing rapidly for the study of metabolites, relying on fast and broad screening (Cox *et al.* 2014) and supported by modern instruments. These instruments include mass spectrometry (MS), which has allowed the detection of many chemical compounds with high sensitivity and precision, thanks to direct-injection electrospray mass spectrometry (DIESI-MS). This technology has proven to be an important tool in the high throughput of non-targeted metabolite fingerprint (García-Flores *et al.* 2015). Therefore, the objective of this study was to evaluate the inhibitory effect of *B. salicina* root extracts on the germination of coffee rust (*H. vastatrix*) uredospores and to identify the composition of the most abundant metabolites in MEBs through use of DIESI-MS and MS².

Materials and Methods

Plant material

Plants were collected in Chiapas, México, in San Cristóbal de las Casas (coordinates 16°45' N and 92°38' O and 2255 m altitude), where Mexican pine-oak forest vegetation and a temperate climate predominates. The *B. salicina* was authenticated and registered under number 49985 at the herbarium of Dr. Faustino Miranda Botanical Garden, in Chiapas, Mexico.

Methanolic extracts of roots from *B. salicina* (MEBs)

The roots of *B. salicina* were dried at room temperature, ground to a fine powder, and sifted using a 2 mm mesh sieve. Subsequently, 300 g of pulverized roots were macerated in 3 L of methanol, as reported by Lam-Gutiérrez *et al.* (2019).

Fraction preparation of MEBs

As an analysis approach, the MEBs were partitioned into three fractions. MEBs were suspended in hexane and the soluble organic layer was separated and concentrated until drying under vacuum conditions at 55°C (hexane fraction). The insoluble residue was suspended in ethyl acetate, after which the soluble organic layer was removed and treated under vacuum conditions in order to concentrate the ethyl acetate fraction at 55°C. Finally, the last insoluble residue was the methanolic fraction. Phytochemical composition of hexane, ethyl acetate and methanol fractions were analyzed by spectrometer mass.

Mass spectrometry (MS) and tandem mass spectrometry (MS²)

Screening tests of spectrum mass analysis were carried out using a mass analyzer quadrupole Micromass ZQ 2000 Waters ®. For all measurements, ionization by electrospray (ESI) was used, both in positive and negative modes. The voltages for the ionization source were established at 3 KV for the capillary, 60V for cone and 3 V at the extractor. The RF lens (orientation range-finding) was used at 0.5 V. Temperatures were set to 80°C and 150°C for the source and the desolvation, respectively. The used gas was nitrogen with a flow of 250 L/h for desolvation and 50 L/h for cone. Multiplier was set to a value of 650 V. The spectra were obtained continuously in a range of 50–2000 m/z. Each run had duration of 1 min, with a scan time of 10 s and an inter-scan time of 0.1 s. The samples were injected at a rate of 10 µL/min. The fragmentation of the molecules (MS²) with the highest concentration in each fraction obtained was performed with the LCQ Fleet™ Ion Trap Thermo Scientific ion trap mass spectrometer, with 6 microscans, at 300°C capillary temperature, 35 V capillary voltage, 4.42 kV spray voltage, 45 V tube lens voltage and 30 units of nitrogen gas (AU) with ionization energies from 20 to 40 eV, in positive and negative mode.

Raw data processing

A revision of the mass spectra was obtained *via* a bioinformatic analysis. Alignment and correction of the baseline was done by extracting data relating m/z and relative intensity by use of M mass V. 5.5.0 software to generate the study matrix. Subsequently, the Metaboanalyst platform version 4.0 was used, with functions written in R software version 3.4.3 for multivariate statistical analysis (Xia and Wishart 2016).

The putative identification of metabolites was cross-referenced against information related to monoisotopic mass, mass spectra, most probable molecular mass and a mechanism for breaking-fragmentation patterns in online libraries (ChemSpider, HMDB, LIPID MAPS, MassBank, MetFrag, METLIN, PubMed, SpiderMass).

Fungi

Samples of rust (*H. vastatrix*) were isolated from the *C. arabica* var. Bourbon located in Ocuilapa, Municipality of Ocozocoutla, Chiapas, Mexico, with the coordinates 16°52'22.46"N and 93°24'39.39"W. Previously, the presence of the fungus was qualitatively identified on infected leaves for a characteristic yellow or orange powder, reniform, smooth on the inside and rough on the outside (Fig. 1; McCain and Henen 1984). The uredospores were scraped from the surface of the collected leaves and stored at 4°C until ready to be used.

Separate leaf test

A suspension of 1×10^5 uredospores of *H. vastatrix* in 50 mL of distilled water was prepared previously, which also included a drop of Tween 20. Afterwards, healthy and young leaves from *C. arabica* plants were cut and immersed for 3 sec in MEBs at 100, 270 and 750 mg mL⁻¹, respectively. They were then placed in Petri dishes with a base of wet filter paper, and the undersides of the exposed leaves were inoculated by dripping 1 mL of the previously obtained spore suspension. Petri dishes were kept in darkness for 24 h. Subsequently, they underwent a 12/12 photoperiod at 22°C and 65% relative humidity. Monitoring of the Petri dishes occurred at 4, 8, 14, 20.5, 32.5, 45.5, 69.5 and 165.5 h after inoculation, which consisted on counting the germinated spores fixated on the leaves. Each coffee leaf was cut from the central part into pieces of approximately 1 cm². The pieces were then collected in a beaker and added 5 mL of Carnoy solution, and let to incubate for 4 h at 60°C.

Subsequently, the Carnoy solution was drained, 5 mL of lactophenol solution were added to the fragment, and let for a 5 h incubation at 60°C. After this time, the lactophenol was strained off of the leaf fragments; the fragments were stained with 0.2% trypan blue and left at room temperature for 1.5 h, then washed with sterile distilled water until the dye was completely washed off. Smears of the washed fragments were placed on each slide with two drops of polyvinyl alcohol, then covered with slips, and let to dry at room temperature.

The same procedure was carried out for each sample. The number of germinated spores was counted at each monitoring time with help of a Primo StarZeiss microscope and a Canon G10 digital camera.

Statistical analysis

The experiments were performed by following a completely randomized design with 3 treatments and three replicates. In addition, two controls were including, one without application of fungicide (negative control) and another with the application of the commercial fungicide used by the producers (positive control). Statistical analyses consisted of an analysis of variance (ANOVA) with $\alpha = 0.05$ and Tukey

test was used to determine the statistically different treatments, with a total of 144 experimental units. This was conducted using the Statgraphic Centurion version XV software.

Results

The experimental tests *in vitro* for antifungal activity allowed the observation of the infective process of the uredospore from *H. vastatrix* on leaves of *C. arabica* var. Bourbon. According to Table 1, in the positive control treatment B (application of inoculum only on leaves) a 15% of spore germination demonstrated after 32.5 h.

MEBs proved to have an inhibitory effect on the germination percentages of uredospores (Table 1). It was found that, regardless of the sampling time, as that of 32.5 h, leaves treated with MEBs significantly decreased the percentage of germination by < 5% as the dose increased ($P < 0.05$).

Antifungal activity: germination percentage

The application of MEBs at concentrations of 250 and 750 mg mL⁻¹ allowed to obtain the highest inhibition percentage of spore germination compared with copper oxychloride used as negative control (Table 1). The latter is actually the most common chemical treatment to prevent leaf rust disease spreading on Coffee arabica crops throughout the American continent. In the compatible interactions between plant and uredospores, a penetration hypha differentiates from an appressorium and develops into the substomatal chamber. At the growing tip of the penetration hypha, two thick lateral branches are formed (anchor), and differentiate into a haustorial mother cell (HMC), which gives rise to haustorium in the stomata subsidiary cells prior to the formation of haustoria in mesophyll cells. This is a unique feature of *H. vastatrix*.

Mass spectrum

In order to know the main chemical components in MEBs with likely antifungal potential, an analytical strategy was proposed for improving their study. The strategy consisted of partition in the chemical components into three fractions, according to their polarity. Fractioning was achieved with use of the following solvents: ethyl acetate, hexane and methanol. In the pre-analytical sample preparation, before instrumental analysis, an important step was carried out to increase detection of low-abundance from roots plant metabolites. Two of the fractions were dissolved once again in a compatible solvent with the ion trap spectrometer aiming to obtain the highest intensity detection. The hexane fraction was dissolved in ACN/MeOH (50:50), the ethyl acetate fraction in ACN, and the methanolic fraction remained suspended in the organic solvent without alterations. Two features *viz.*, 5386 and 1189 in ESI (+) and (-) ionization mode, respectively, were obtained after

Table 1: *In vitro*: Effect of MEBg on the germination capacity of uredospores from *H. vastatrix*. Tukey test ($P < 0.05$). The uppercase letters indicate fixed leaves: A) Without inoculum and treatment; B) Positive control (Inoculum); C) Negative control: copper oxychloride; D) 100 mg mL⁻¹; E) 270 mg mL⁻¹; F) 750 mg mL⁻¹

Treatment	Time (h after inoculation)							
	4	8	14	20.5	32.5	45.5	69.5	165.5
A	0.00 ^a	0.00 ^a	0.00 ^a	0.00 ^a	0.00 ^a	0.00 ^a	0.00 ^a	0.00 ^a
B	0.00 ^a	0.00 ^a	0.00 ^a	0.69 ^b	14.86 ^b	21.66 ^b	37.52 ^b	37.35 ^b
C	1.33 ^b	0.00 ^a	4.44 ^b	0.90 ^b	2.75 ^c	3.83 ^c	23.21 ^c	38.54 ^b
D	1.95 ^b	1.51 ^b	1.92 ^b	2.48 ^c	4.45 ^c	3.52 ^c	16.43 ^c	7.01 ^c
E	2.29 ^b	0.00 ^a	0.00 ^a	1.23 ^{bc}	0.00 ^a	1.33 ^{cd}	5.03 ^d	1.67 ^d
F	2.56 ^b	0.00 ^a	0.00 ^a	8.98 ^c	3.61 ^c	0.00 ^a	4.32 ^d	2.20 ^d

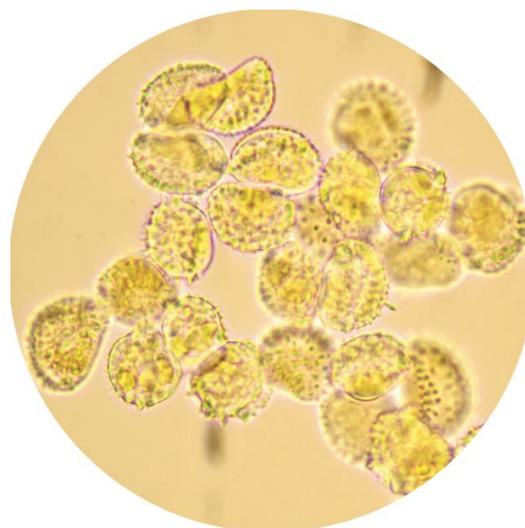
Latin letters a, b, c, d (superscript) in the same column indicates significant differences

Table 2: Significant compounds detected in roots of *Baccharis salicina* by DIESI-MS, positive mode

Fraction	m/z [m + H] + (ionization mode)	Fragment pattern [m+H] ⁺ (MS ²)	The most intense compounds in <i>Baccharis glutinosa</i> roots			Putative compound	Class	Data base	ID data base
			Ionization energy (V)	Molecular formula					
Ethyl acetate	284.1249 (+)	228.17, 157.17, 170.08, 198	35	C ₁₇ H ₃₂ O ₃	5-(1-hydroxytridecyl)oxolan-2-one	Fatty Acyls	Metlin	93340	
	380.3365 (+)	252.25, 266.25, 238.50, 280.25, 308.25, 197.25	35	C ₂₂ H ₃₀ O ₆	Multijuginol	Polyketides: flavonoids	Metlin	48617	
	174.0679 (+)	156.17, 97.92, 115.92, 132.08, 86	25	C ₉ H ₁₈ O ₃	3-hydroxy-nonanoic acid	Fatty Acyls	Metlin	35407	
	156.8757 (+)	138.17, 113.83, 74.00, 97.92, 30, 55.08, 128	30	C ₆ H ₄ O ₅	2,5-Furandicarboxylic acid	Organoheterocycli c compounds: furans	MoNA Spectrum	HMDB0004812 _ms_ms_2403	
	190.9425 (+)	174.17, 156.00	25	C ₆ H ₈ O ₇	Glucaric acid lactone; (2S)-[(2S,3R,4R)-3,4-dihydroxy-5-oxotetrahydrofuran-2-yl](hydroxy)ethanoic acid	conjugate acid	Metlin	44758	
Hexane	380.9073 (+)	252.17, 325.17, 366.33, 123	30	C ₂₂ H ₃₀ O ₆	Multijuginol	Polyketides: flavonoids	Metlin	48617	
	240.0411 (+)	143.08, 97.17, 222.08, 183.08, 30, 80.83	30	C ₁₅ H ₁₂ O ₃	(Z)-3-hydroxy-1-(2-hydroxyphenyl)-3-phenylprop-2-en-1-one; también se llama 2'-beta-Dihydroxychalcone	polyketides: flavonoids: chalcones	Metlin	43644	
	333.0255 (+)	96.75, 309.33, 180.00	-	-	-	-	-	-	
Methanolic	381.0341 (+)	252.17, 325.17, 366.33, 123	35	C ₂₂ H ₃₀ O ₆	Multijuginol	Polyketides: flavonoids	Metlin	48617	
	218.8543 (+)	111, 147.17, 159.17, 204.08, 30, 135.08, 119	30	C ₁₅ H ₂₂ O	Dendrolasin (3-(4,8-Dimethyl-3,7-nonadienyl)-Furan)	Farnesenes	Metlin	71335	
	365.1163 (+)	203.08, 184.92	25	C ₁₄ H ₁₇ NO ₉	(2S)-7-Hydroxy-3-oxo-3,4-dihydro-2H-1,4-benzoxazin-2-yl glucopyranoside	O-glycosyl beta-D- compounds	MoNA Spectrum	CCMSLIB00000 849622	
	332.8352 (+)	240.25, 296.42, 270.42, 254.08	30	C ₁₉ H ₂₄ O ₅	Gibberellin A51	Prenol-lipids: isoprenoids	Metlin	41234	
	155.8606 (+)	111.92	30	-	-	-	-	-	

alignment and integration correction of the set of samples fingerprint by DIESI-MS. The data matrix was used for statistical evaluation. Representative peaks of samples in positive mode are shown in Fig. 2.

The MEBs were found to have various functional groups of compounds. The most abundant putative secondary metabolites identified (DIESI-MS in positive mode) were polyketides (flavonoids), such as multijuginol with the peak corresponding to 380 m/z in ethyl acetate and hexane fractions, 2'-beta-Dihydroxychalcone with peak at 240.041 m/z in hexane fraction, these were followed by furans, such as 2,5-Furandicarboxylic acid in ethyl acetate fraction (Fig. 2 and Table 2). On the other hand, the most abundant compounds identified (DIESI-MS in negative mode) were polyketides (flavonoids), such as 3'-methoxyquercetin with the peak corresponding to 316(-) m/z in ethyl acetate fraction, 12a β-hydroxydeguelin with its peak at 411(-) m/z in hexane fraction and 3,4-dihydroxycinnamic acid with the peak at 179(-) m/z in methanolic fraction (Table 3).

**Fig. 1:** Uredospores of *H. vastatrix*, contrast microscope: Objective 100X. Canon camera G10

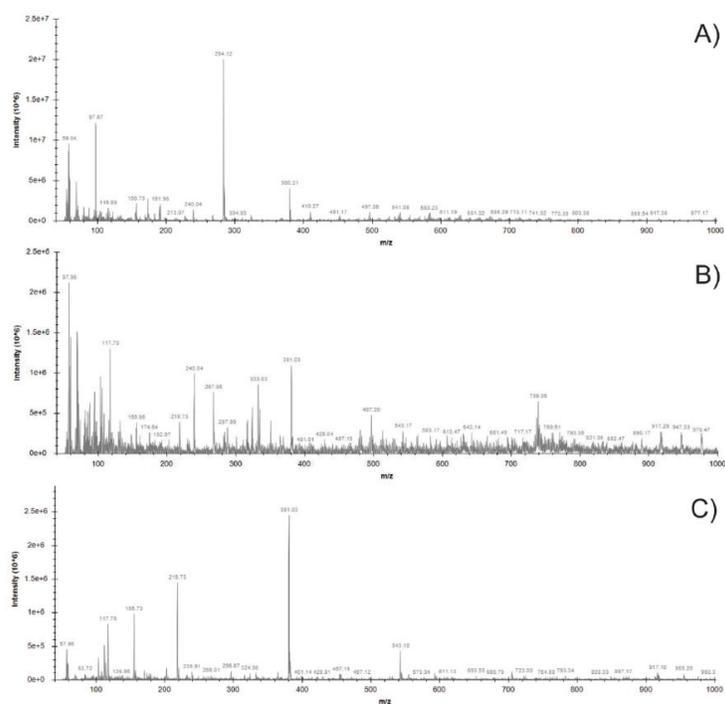


Fig. 2: Mass spectrum of MEBs partitioned: **A)** Ethyl acetate, **B)** Hexane and **C)** Methanolic fractions. Acquired with DIESI-MS in positive mode

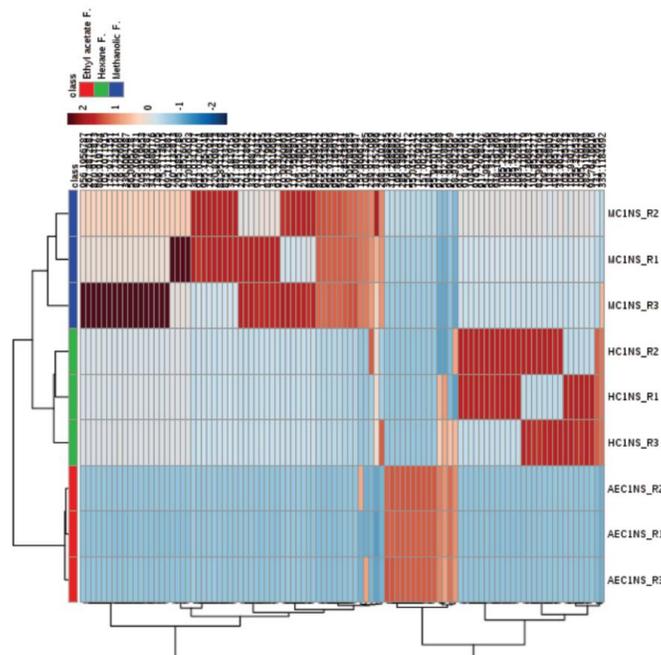


Fig. 3: Heatmap clustering analysis performed in MetaboAnalyst 4. Using DIESI (+) MS spectra of the metabolites obtained in three fractions different of methanolic extracts roots of *Baccharis salicina*. Analysis was performed using Euclidian distance method with ward clustering algorithm after normalization to sample median and Pareto scaling

Multivariate analysis

The hierarchical classification by "clusters" can be observed in the graphical representation of a heat map

(Fig. 3). On the right side, there are 100 representative m/z of all the samples studied and identified via mass spectrometry. In the lower part of the graph, the fractions in which the crude methanolic extract was partitioned are represented.

Table 3: Significant compounds detected in roots of *Baccharis salicina* by DIESI-MS, negative mode

Fraction	[m-H] ⁺ (ionization mode)	m/z	Fragment pattern [m-H] ⁺ (MS ²)	The most intense compounds in <i>Baccharis glutinosa</i> roots			Class	Data base	ID data base
				Ionization energy (V)	Molecular formula	Putative compound			
Ethyl acetate	174 (-)	156, 98, 116, 138, 86, 69, 54		25	C ₇ H ₁₀ O ₅	(3R,4S,5R)-3,4,5-Trihydroxycyclohex-1-enecarboxylic acid	Cyclohexanecarboxylic acid.	MoNA	Spectrum PR100485
	265 (-)	96		35	C ₁₈ H ₁₈ O ₂	2-(4-hydroxy-3-prop-2-enyl-phenyl)-4-prop-2-enyl-phenol	Benzene and substituted derivatives	MoNA	Spectrum FiehnHILIC001252
	297 (-)	182.92, 196.92, 211.83, 233		30	C ₁₄ H ₁₉ N O ₆	(3,4,5-trihydroxy-6-methyloxan-2-yl)-2-(methylamino)benzoate	Organooxygen compounds:	MoNA	Spectrum VF-NPL-QEHF019193
	316 (-)	106, 298.33, 189.25, 272.25		30	C ₁₁ H ₁₂ O ₇	3'-Methoxyquercetin	Carbohydrates and carbohydrate conjugates	MoNA	Spectrum PR100640
	325 (-)	182.92, 196.92, 169.92, 224, 261		30	C ₂₇ H ₄₂ O ₂	Heineicosanoic acid	Flavonoids: flavonols	MoNA	Spectrum EXPO_D10_5
	241 (-)	171, 79.8, 145, 196.92		25	C ₁₅ H ₁₂ O ₃	2,4-Dihydroxychalcone, también se llama 3-(2,4-Dihydroxyphenyl)-1-phenyl-2-propen-1-one	Fatty acyls: Fatty acids and conjugates	MoNA	ppm_NEG_iTree 39612
	439 (-)	337.17, 424.25, 395, 368, 323, 293.42, 162.92, 176.92		25	C ₂₆ H ₃₂ O ₆	Oxireno(o)phenanthro(1,2-d)pyran-3,8(3aH,4bH)-dione, 1-(3-furanyl)-1,5,6,6a,7,10a,10b	Phenylpropanoids and polyketides	MoNA	Spectrum NP_C3_114_p2_A05_iTree_NEG_46
	410 (-)	330.33, 258.25, 392.33, 272.33, 244.17, 340.25		30	-	-	-	-	-
	514 (-)	353, 172.92, 203		25	-	-	-	-	-
	255 (-)	175, 197, 210.92, 236.83		25	-	-	-	-	-
311 (-)	182.92, 196.92, 247.17, 169.92, 267.25, 280		30	-	-	-	-	-	
581 (-)	501.25, 563.17, 416.5, 374.25		25	-	-	-	-	-	
Hexane	182.92 (-)	138.92, 118.92, 166.75, 155		25	C ₉ H ₁₀ O ₄	4-O-Methylphloracetophenone	Organooxygen compounds:	MoNA	Spectrum VF-NPL-QEHF011678
	325.17 (-)	182.92, 197, 169.92, 261.08, 239.08		30	C ₁₅ H ₁₆ O ₅	(Z)-3-hydroxy-1-(2-hydroxyphenyl)-3-phenylprop-2-en-1-one; también se llama 2'-beta-Dihydroxychalcone	Phenylketones	MoNA	11688418
	383 (-)	337.42, 365.25		30	C ₂₃ H ₃₂ O ₄	5-[(Z)-12-(3,5-dihydroxyphenyl)dodec-8-enyl]benzene-1,3-diol	Carboxylic acids and derivatives	PubChem	
	411 (-)	365.42, 337.08, 382, 395.25, 349.17, 251.08		30	C ₂₃ H ₃₂ O ₇	12aβ-hydroxydeguelin	Phenols: Resorcinols	MoNA	Spectrum VF-NPL-QEHF021775
	142.75 (-)	99.83, 98.92, 110.75, 72.75, 123, 125.17		25	-	-	-	-	-
	255.25 (-)	227, 211, 196.83, 236.92		25	-	-	-	-	-
	311 (-)	482.92, 196.92, 247.08, 169.92, 196.92		30	-	-	-	-	-
	377 (-)	219, 340.75, 303.08, 314.92, 191, 182.92, 331.08, 159		25	-	-	-	-	-
	455.33 (-)	419.42, 393.17, 407.33, 353.08, 332.83		25	-	-	-	-	-
	499 (-)	397.25, 382.08, 367, 484.17, 255, 453.42, 407.67, 25		25	-	-	-	-	-
	671 (-)	469.08, 243.25, 271.08, 425.25		30	-	-	-	-	-
	281.33 (-)	391.42, 256.25, 409.17, 610.58, 627.58, 581.5, 278.233		25	-	-	-	-	-
	325 (-)	263.08, 239.08, 209.08, 124.92, 111, 191, 252.92		25	-	-	-	-	-
		182.92, 310.17, 261.17, 281, 225, 238		30	-	-	-	-	-
	Methanol	233 (-)	143.08, 171.08, 127, 156.83, 186.75		25	C ₁₃ H ₁₂ O ₄	Goniothalenol o bien 3-hydroxy-2-phenyl-2,3,3a,7a-tetrahydrofuro[3,2-b]pyran-5-one	Furofurans	MoNA
515 (-)		353, 191, 427.58, 334.92, 317, 399, 172.75		20	C ₂₅ H ₃₂ O ₁₂	3,5-Dicafeoylquininic acid	Organooxygen compounds	MoNA	Spectrum CCMSLIB00000081755
793 (-)		647.25, 571.42, 393.83, 585.33, 629.25, 603.08, 553.08		25	C ₃₆ H ₅₆ O ₁₁	Olean-12-en-28-oic acid, 3-(beta-D-glucopyranuronosyloxy)-23-hydroxy-, (3beta,5xi,9xi,18xi)-	Phenol lipids	MoNA	Spectrum CCMSLIB00000849986
179 (-)		134.92, 163.92, 88.67, 68.83		20	C ₉ H ₈ O ₄	3,4-Dihydroxycinnamic acid	Prenol lipids	MoNA	Spectrum KO000511
311 (-)		182.83, 250.92, 197, 280.83, 170.92, 210.75, 140.50, 122.67		35	C ₁₉ H ₂₁ NO ₄	1,10-Dimethoxy-6-methyl-5,6,6a,7-tetrahydro-4H-dibenzo[de,g]quinoline-2,9-diol	Phenylpropanoids and polyketides	MoNA	Spectrum 248507
353 (-)		191, 178.92, 172.92, 131, 196.92,		35	C ₁₆ H ₁₈ O ₉	3-(3,4-Dihydroxycinnamoyl)quinic acid	Aporphines	PubChem	1794427
835 (-)		585.25, 647.33, 107.17, 817.42, 634.75		30	-	-	-	-	-
915 (-)		835.33, 727.17, 647.33, 629.17, 254.92, 769.33		30	-	-	-	-	-
191 (-)		92.83, 126.83, 108.83, 154.83, 172.83		20	-	-	-	-	-
568 (-)		489, 416.83, 367.25, 365.50, 527.25, 479.17		25	-	-	-	-	-
877 (-)		585.17, 513.25, 629.33, 471.33, 859, 479.25		45	-	-	-	-	-
369 (-)		323.33		30	-	-	-	-	-
383 (-)		337.42		30	-	-	-	-	-
398 (-)		352.33, 396.17, 382.75		25	-	-	-	-	-
472 (-)		408.50, 444.08, 410.25, 316.25, 454.25, 288.33, 426.58, 370, 158.92, 300, 311, 324, 370		25	-	-	-	-	-
411 (-)		365.33		25	-	-	-	-	-

The color scale (-2 to 2) indicates the percentage of relative abundance of the m/z detected for each fraction. For equality of observable characteristics, the heat map classified the study fractions into a dendrogram. Partitioning the crude extract into three fractions (methanol, hexane and ethyl acetate), allowed to separate metabolites attracted by different polarities. The greatest number of peaks (m/z) was found to be in the methanolic fraction, followed by hexane and ethyl acetate fractions (Fig. 3).

In total, 539 features in ESI (+) and 119 in ESI (-) mode were found to be statistically significant. The cross-validation based on the filtered data matrix was performed to estimate the predictive ability of the multivariate Partial Least Squared Discriminant Analysis (PLS-DA) models (Fig. 4). In the score plot, the two main components explain the greater effect of analyzing the methanolic extracts of Chilca roots (*B. salicina*) in the three fractions that were considered. The analysis showed the separation of variables

organized into clusters, in a two-plane graph. The observed trend indicates that, depending on the nature of the fraction, there are characteristic m/z for each (Fig. 4A). The main 25 m/z identified in positive mode represent the repeated metabolites in the study fractions, considering the Variable Influence on the Projection (VIP) (Fig. 4B).

Tandem mass spectrometry (MS²)

The screening analysis allowed a total of 54 compounds to be identified and characterized in both positive and negative mode from MEBs. Fragmentation patterns (MS²) were used to determine the identity of each metabolite, based on the most probable molecular mass. This was also verified with on-line libraries and breaking mechanisms. The list of putatively identified compounds is shown in Table 2 and 3.

Discussion

Loureiro *et al.* (2015) found 43% germinated uredospore in the *in vitro* study on *C. arabica* H147/1 24 h after inoculation (Diniz *et al.* 2012; Yoon *et al.* 2013) and 34% in the *in vivo* condition. In a natural situation, for infection to be completed, spores germinate better on young leaves than on intermediate and old leaves, as they require free water. A temperature between 22–28°C is another important factor influencing germination along with moisture. A single lesion on a leaf produces 4 to 6 spore crops over a 3 to 5-month period, releasing 300–400,000 spores (Yoon *et al.* 2013).

Uredospore germination and fungal penetration are key stages for the initial development of the coffee leaf rust pathogen *H. vastatrix* (Talinhas *et al.* 2016). Fungi infecting from uredospores are particularly sensitive during the initial stages of the disease cycle, especially between spore germination and host penetration (Thines *et al.* 2004). The presence of antifungal metabolites from MEBs suggests a possible breakdown in the first communications between plant and uredospore.

The growth cycle of *H. vastatrix* is known to begin at the spotting of the host surface, in which uredospores germinate inserting appressoria into the stomata (Talinhas *et al.* 2016). Therefore, the surface hydrophobicity is an important stimulus for inducing the infection structure in the plant (Kou and Naqvi 2016). It has also been detected at this stage that spores release an adhesive drop from a periplasmic compartment, which attaches the uredospore to hydrophobic surfaces. This glue is likely to contain glycoproteins. It has been observed that plant lectin concavalin A inhibits binding (Thines *et al.* 2004).

The results presented herein indicate that full inhibition in germination percentage is not achieved. However, there is a trend linking increased treatment doses with reduced germination percentages (Table 1). This means that some important proteins for generating growth could be lost, thus delaying the biotrophic development and growth

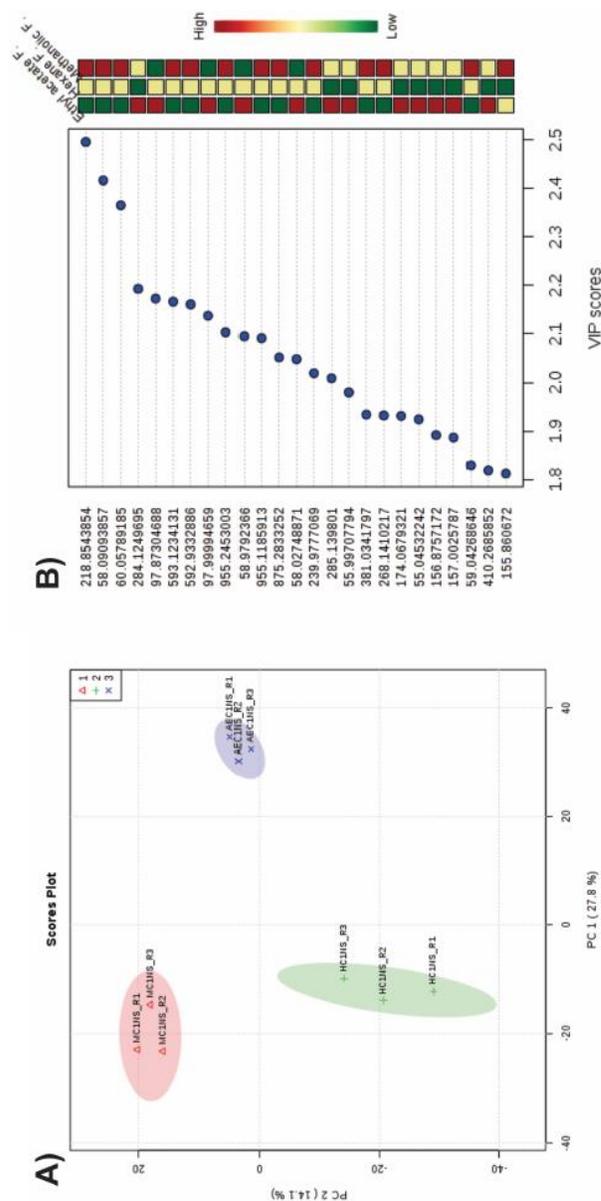


Fig. 4: A) Score plot of the first two principal component of PLS-DA using normalization to sample sum and Pareto scaling. B) Significant features identified by PLS-DA performed in MetaboAnalyst 4.0. Positive mode

(Meyer *et al.* 2009). Deising *et al.* (2000) and Bölker (1998) reported that chitin deacetylases (CD), G-protein subunits and MAP Kinases (MAPKs) were associated with early developmental phases of several pathogenic fungi. The genes involved in the early infection process of *H. vastatrix* on *C. arabica* leaves were confirmed by Vieira *et al.* (2011) with an expression profile (Vieira *et al.* 2012).

In this study, the flavonoids putatively found were multijuginol, (Z)-3-hydroxy-1-(2-hydroxyphenyl)-3-phenylprop-2-en-1-one, 3'-Methoxyquercetin and 12aβ-

hydroxydeguelin. The incidence of 3'-Methoxyquercetin was previously reported in leaves of *B. pseudotenuifolia* by Moreira *et al.* (2003). In addition, flavonoids, phenolic acids and triterpenes extracted from leaves and roots from *B. dracunculifolia* were reported with immunomodulatory biological activity (Figueiredo-Rinhel *et al.* 2013; Ramos *et al.* 2016). An important molecule included in the tetracyclic diterpenoid class are Gibberelins (signaling compounds or plant hormones), which regulate developmental processes such as seed germination, root and shoot elongation, flowering and fruit patterning (Shani *et al.* 2013). They therefore coordinate the synthesis of other molecules. In this study, these were detected in roots. *In vitro* studies have shown that phenolic compounds like flavonoids, coumarins and phenolic acid can have antioxidant activity, which allows them to act as reducing agents (Barchan *et al.* 2014).

On the other hand, the presence of furans 2, 5-furandicarboxylic acid, dendrolasin (3-(4,8-dimethyl-3,7-nonadienyl)-furan) and 3-hydroxy-2-phenyl-2,3,3a,7a-tetrahydrofuro[3,2-b]pyran-5-one, coincide with previous work by Lam-Gutiérrez *et al.* (2019) where this molecule class was found *via* GC-MS, albeit with a different molecular mass. (2E,6E)-2-(4-methylpent-3-en-1-yl)-6-[3-(2,5-dihydro-2-oxofuran-3-yl)-propylidene]hept-2-enedioic acid was found in aerial parts from *B. thymifolia* (Hikawczuk *et al.* 2008). According to Konovalov (2014), a type of polyacetylene commonly spotted in the Asteraceae family is furanopolyacetylene.

In the study of MEBs, compounds such as 3, 4-dihydroxycinnamic acid and 3-(3,4-dihydroxycinnamoyl)quinic acid belonging to phenylpropanoids and polyketides class (including hydroxycinnamic acids) could be observed. These kind of molecules have been reported in aerial parts of *B. grisebachii*, *B. chilco*, *B. trimera*, *B. retusa*, *B. incarum* (leaves and stems) and *B. dracunculifolia* (leaves, stems and roots) (Ramos *et al.* 2016). García-Jimenez *et al.* (2018) showed that cinnamic acid, 2-hydroxycinnamic, 2,3 and 4-methoxycinnamic acids were tyrosinase inhibitors. In the same way, it was reported on the fungal β -1,3-glucanase and chitinase activities of maize phytopathogenic fungi and extracts of aerial parts of *B. salicina* (Buitimea *et al.* 2013), was important for the development of fungi. Additionally, Taofiq *et al.* (2017) found that the derivatives of cinnamic acid had antifungal, antibacterial, antioxidant, anti-inflammatory and antitumoral activities.

The type of extraction solvent and its polarity are known to potentially have a significant impact on the number of metabolites extracted (Thouri *et al.* 2017). Methanol molecules are proton donors, making it the most used solvent in the crude extraction of metabolites, considering a broad-molecule spectrum. Nevertheless, the present study included also a first extraction with methanol, a partition with two other organic solvents that was able to separate metabolites attracted according to the polarity of each one (hexane > ethyl acetate). Hence, the separation of metabolites could be observed. This behavior is largely due

to the chemical nature of the organic solvent used. Methanol is a protic polar solvent with the capacity of binding by hydrogen bridges, whereas hexane is non-polar, as it lacks functional groups capable of yielding protons and has a high dielectric constant. On the other hand, ethyl acetate is known for being an aprotic polar solvent and for having a low dielectric constant. Just as hexane, it is lacking in functional groups.

The proposed analysis allowed the finding of major metabolites, which has been suggested as belonging to polyketides (flavonoids, flavonols, isoflavones and chalcones), organoheterocyclic compounds (furans and furofurans), fatty acyls, prenol-lipids (isoprenoids), organooxygen and farnesene classes. Ramos *et al.* (2016) reported that the main constituents of *Baccharis* genus are phenolic and terpenoid compounds, demonstrating that flavonoids and diterpenoids have been the major classes of metabolites, as this study reveals as well. Likewise, antifungal activity in *Baccharis* genus has been reported in these compounds, as coumarins, triterpenes, sesquiterpenes and flavonoids are present in *B. darwini* (aerial parts), *B. dracunculifolia* (aerial parts), *B. elaeoides* (leaves), *B. semiserrata* (twigs) and *B. retusa* (twigs) (Kurdelas *et al.* 2010; Johann *et al.* 2012; Vannini *et al.* 2012; Grecco *et al.* 2014). Furthermore, antifungals have been reported as being present in roots of *B. salicina* for the first time. Therefore, the fingerprinting by use of DIESI-MS offers an important tool for pioneering descriptive analysis of metabolites in a reliable screening.

Conclusion

Concentrations of MEBs > 100 mg mL⁻¹ on leaves of coffee reduce the germination of uredospores from *H. vastatrix*, with significant statistical differences. Using the DIESI-MS tools, a general screening was obtained that allowed the generation of metabolite fingerprints. Greater relative abundance was present in MEBs in an m/z range of 50-2000. The previous partition of crude extract methanolic of root from Chilca into three fractions improved the chemical composition study conducted with successive MS-MS analyses, thus identifying an estimated 50 of the most important metabolites. We also speculated briefly about the possible roles of such substances interrupting communication between uredospores and plant in early stages of infection process or reducing synthesis of important proteins which can affect fungal growth and morphological differentiation. Therefore, the study contributes to composition and properties analysis of bioactive phytochemicals from *B. salicina* against coffee leaf rust pathogen *H. vastatrix*. This research should be furthered to better assess the potential activity of *B. salicina* on multiple other applications in the future.

Acknowledgments

This research was funded by Project number '6211.17'

(TecNM, Mexico). The authors wish to thank the ‘Instituto Nacional de Investigaciones Forestales, Agrícolas y Pecuarias (INIFAP)’. ALG wishes to acknowledge CONACYT Mexico for her postgraduate scholarship.

Author Contributions

AL-G and VMR-V conceptualized the work, ERG-R and RW provided laboratory facilities for analyzing the MEBs, RR-R and FAG-M reviewed the manuscript, AL-G statistically analyzed the data, BAP-O and JMG-A analyzed the final version of the graphics.

Conflicts of Interest

The authors declare that they have no conflicts of interest concerning this article.

Data Availability

The data will be made available on acceptable requests to the corresponding author.

Ethics Approval

Not applicable.

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