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Article

Antimicrobial Activity and Phytochemical Characterization of *Baccharis concava* Pers., a Native Plant of the Central Chilean Coast

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Abstract: Few Sclerophyllous plants from the central coast of Chile have been systematically studied. This work describes the phytochemical composition and antimicrobial properties of *Baccharis concava* Pers. (sin. *B. macraei*), a shrub found in the first line and nearby the pacific coast between latitudes 29-38°S. *B. concava* has been traditionally used by indigenous inhabitants of today central Chile for their anthelmintic properties, wound healing properties and as a diuretic. Hydro-alcoholic extracts of *B. concava* were prepared from leaves and small branches. The qualitative phytochemical characterization indicated presence of alkaloids, steroids, terpenoids, flavonoids, phenolic and tannins compounds. The antimicrobial activity of this extract was assessed in a panel of microorganisms including Gram-positive, Gram-negative bacteria and pathogenic yeasts. The extract displayed important antimicrobial effects against Gram positive, *Candida albicans*, and *Cryptococcus neoformans*, but not against Gram negative, for which an intact Lipopolysaccharide (LPS) is apparently the determinant of resistant to *B. concava* extracts. The hydroalcoholic extract was then fractionated through a sephadex LH-20/methanol-ethyl acetate column. Followed, the fractions were pooled according to a similar composition that was visualized by TLC/UV analysis. Fractions obtained by this criterion were assessed for their antimicrobial activity against *Staphylococcus aureus*. The fraction presenting the most antimicrobial activity was HPLC-ESI-MS/MS analyzed obtaining a composition rich in structures derived from caffeoylquinic acid, dicaffeoylquinic acid and quercetin, among others. In conclusion, the extracts of *B. concava* show a strong antimicrobial activity probably due to their composition rich in metabolites derived from caffeoylquinic acid and quercetin, that in turn could be responsible for helping with wound healing, as reported from its ancient cultural use. In addition, the development of antimicrobial therapies based in the molecules found in *B. concava* could help to combat infection caused by pathogenic yeasts and Gram-positive bacteria, without affecting the Gram-negative microbiota.

Keywords: baccharis concava; chilean medicinal plant; phytochemical composition; antimicrobial properties; used traditional

1. Introduction

The genus *Baccharis* is part of the Asteraceae family, one of the largest families in the Plantae kingdom. It is estimated that there are around 500 species of the genus *Baccharis* from North to South America, of which 210 inhabit the southern cone (Abstract et al., 2019). It is possible to find members

of *Baccharis* in climatic zones including temperate, tropical, and desert climates, either cool or warm and at any elevation. The species of *Baccharis* genus share many anatomical and histologic characteristics, however few studies extensively describe the genus. The species of the genus *Baccharis* are usually branched leafy shrubs that can measure from 0.5 to 4 meters in height. Some species, such as *Baccharis concava* Pers. (syn. *B. macraei*) can be found at elevations close to sea level. *B. concava* is abundant in the central coastline of Chile and considered a pioneer specie with ability to grow in poor soils, including sands of the first line of the coast. Another example is *Baccharis obovata*, found in Chile and Argentina, closely related to other *Baccharis*, such as *B. concava* and *B. magellanica* (Molares et al., 2009).



Figure 1. Details of *B. concava* small branches and leaves used for the study. Branches and leaves were collected from a female plant located in a private garden around 400 m from San Sebastian beach and 80 m from the Cartagena estuary, during February (middle summer) 2017. Precise location (33°31'46.8"S 71°35'59.3"W) is shown at the right panel, obtained from Google Earth App. By the end of summer (late February and onwards) intense blooming, as seen at the left, massively attract bees and bumblebees (left panel).

Table 1. Qualitative determination of secondary metabolites in hydro-alcoholic extracts of *B. concava*.

Assay	Compounds tested	Positive results	Result
Dragendorff	Alkaloids	red precipitated	+
Borntrager	Free Anthraquinones	red color in aqueous fase	-
Flurescence under UV	Cumarines	blue color under UV light	-
Libermann-Burchard	Steroids y terpenes	green-blue or purple red	+
Aluminum chloride	Flavonoids	yellow green fluorescence under UV light	+
Keller-Killiani	Cardiac glycosides	greenish blue color	-

Foam formation	Saponins	stable foam production (stands 10 min)	-
Ferric chloride	Tannins and phenolic compounds	green or dark blue color	+

While systematic studies provide support for popular and ancestral uses of several species of the genus *Baccharis* from Brazil, Argentina, and Uruguay, few studies describe medicinal properties and the phytochemical characteristics of *B. concava* and other species of *Baccharis* in Chile. In 1985, Houghton and Manby, found from a list of 136 species with medicinal properties use by the native Mapuche people in Chile, three species of the *Baccharis* genus: *B. concava* used as vermifuge; *B. rosmarinifolia*, whose resins has been used for treating rheumatism and respiratory and genitourinary infections; and *B. sagittalis* for the treatment of fractured limbs (Houghton and Manby, 1985). Other species of the *Baccharis* genus such as *B. articulata*, *B. crispa* and *B. dracunculifolia* have been used for the treatment of ulcers, wounds, alleviation of gastrointestinal discomfort and infections, usually used as a decoct of leaves applied on skin or for drinking (Abad and Bermejo, 2007; Desmarchelier, 2015). In 1986, preparing ethanolic extracts of *B. linearis*, *B. rhomboidalis*, and *B. solieri* , Labbé *et al.*described , after partition with solvents of higher polarity, the presence of terpenes, coumarins and fatty esters of different nature.(Labbe *et al.*, 1986).

Preparations of *B. concava*, has been used in pre- and post-Columbian traditional medicine to treat wounds, preventing infections, as diuretic and as a tonic beverage. Up-to-day, there are few scientific studies describing medicinal properties and the phytochemical characteristics of *B. concava*. In 1989, preparing dichloromethane extracts of aerial parts of *B. concava*, Gambaro *et al.* found derivatives of the clerodane diterpenes such as hardwickiic acid, hautriwaic acid, and derivatives of bacchasmacranone (Gambaro *et al.*, 1986). Also using the aerial parts of *B. concava*, but performing a hydromethanolic followed by partitions with petroleum ether, ethyl ether and ethyl acetate, the flavonoids salvigenin, cirsimaritin and pectolinarigenin were purified (Zamorano *et al.*, 1987). In 2012, antimicrobial activity of *B. concava* againsts *Staphylococcus aureus* (Gram-positive) but not for *Escherichia coli* (Gram-negative) was described in essential oils. From a total of 102 identified compound; limonene, miricene, α -pineno, murolene espatulene, δ -cadineno and lachnophyllum ester were the most abundant compounds, with different proportion, depending in the gender of plants (Santander Meyer, 2012).

Antibacterial effects have been reported for species of the *Baccharis* genus, including *B. concava*, the wound healing process is promoted by *B. concava* (according to traditional medicine) and it could be due to antimicrobial effects. Because *B. concava* have not been extensively described in the scientific literature, we sought in this study describing the possible antibacterial effects and phytochemically describing the hydroalcoholic extracts of *B. concava*.

2. Material and Methods

Plant Material.

Leaves and small branches of *B. concava* were collected in San Sebastian Beach, V region, Chile (33°31'46.8"S 71°35'59.3"W) from a female plant which material was validated by MRD and deposited at the herbarium of Facultad de Ciencias Químicas y Farmaceuticas, Universidad de Chile under code number SQF #22.889. Leaves and small branches of *B. concava* were dried for 14 days at 21°C and crushed in a porcelain mortar. 100g of dry plant powder were macerated with 1000 mL of ethanol (70% in water) for 3 days at 50 rpm and 30°C. after 3 days the extract was filtered and rotovaped from 30 to 70°C before further drying with a lyophilizer at -50°C, until the solvents are eliminated. Finally, 42.51 g of dry extract was obtained. The extract was stored in an amber bottle, at a temperature of 4°C, according to methodologies reported and standardized in the literature for this type of study (Tiwari *et al.*, 2011).

Phytochemical characterization

Standard reactions were performed to qualitatively identify the main families of compounds present in the hydroalcoholic extract of *B. concava*. The performed tests include: Dragendorff for alkaloids, Borntrager for anthraquinones, fluorescence for coumarins, Libermann-Burchard for steroids and terpenes, aluminum chloride for flavonoids, Keller-Killiani for cardiac glycosides, foam formation for saponins and Ferric chloride for tannins and phenolic compounds. Details in how to perform these tests have extensively been described in the literature (María et al., 2018; Pandey and Tripathi, 2014).

Antimicrobial activity assays by an agar-diffusion test.

90mm Petri-like plates were filled with 20 mL of Miller-Hinton agar. Once Miller-Hinton agar was jellified, the agar was seeded to form a lawn of bacteria, by zigzag streaking a cotton swab soaked in diluted bacteria (McFarland 0.5 in saline serum). Right after seeding the lawn, each plate was 6mm punched to form 4 holes, 3 peripheric to test 100 μ L of the *B. concava* extract (333.33 mg/mL dried extract dissolved in 70% EtOH) and one at the center to test for 100 μ L of 70% EtOH vehicle. After incubating 16 h, the diameter was measured through 3 different sections for each inhibition halo. Independent experiments were performed at least 3 times. To test susceptibility of yeast such as *Candida albicans* and *Cryptococcus neoformans*, the assay was as described, but Miller-Hinton agar was replaced with Potato-Dextrose agar.

Minimum inhibitory concentration (MIC).

Starting with a solution of 166,67 mg/mL of lyophilized extract, base two serial dilutions were prepared in LB-broth (for bacteria) or Potato-Dextrose broth (for yeasts). 150 μ L of each dilution were loaded through the rows of a flat-bottom 96-well plate to finally seed with 50 μ L of diluted bacteria or yeast (adjusted to McFarland 0.5 in saline serum). After 16 h incubation, turbidity of cultures was measured at a longitude of 600 nm to calculate IC₅₀, which was consider as the MIC. As a control, bacteria were treated with same amount of vehicle to look for possible effects in growth.

Minimum biocidal concentration (MBC).

From the same set of experiments intended to determine the MIC (or IC₅₀), after 16 h incubation, aliquots of 5 μ L from each dilution were seeded on top of a LB agar or Potato-Dextrose agar, incubated for 16 h to finally count the colony forming units (CFU). MBC was estimated as the concentration capable of eliminate 99.9% of a microorganisms compared to the control without treatment.

Thin layer chromatography (TLC).

Characterization of extracts and eluted fractions were characterized by thin layer chromatography using aluminum oxide (Al₂O₃) as the stationary phase and methanol/ethyl acetate (1:1) as the mobile phase. UV (254 nm y 365 nm) and visible longitudes were used to visualize characteristic bands in crude and purified extracts.

Sephadex preparatory column.

A 290 mm long x 21 mm wide LH-20 column was selected as the stationary phase to separate 300 mg of dried *B. concava* extract dissolved in 3 mL of methanol/ethyl acetate (1:1). To elute the column methanol/ethyl acetate (1:1) was used as mobile phase. A total of 40 fractions of approximately 5 mL each were collected, resolved by TLC and pooled by their patterns as shown in Figure 2. Finally, antimicrobial activity was determined to continue with further chemical characterization.

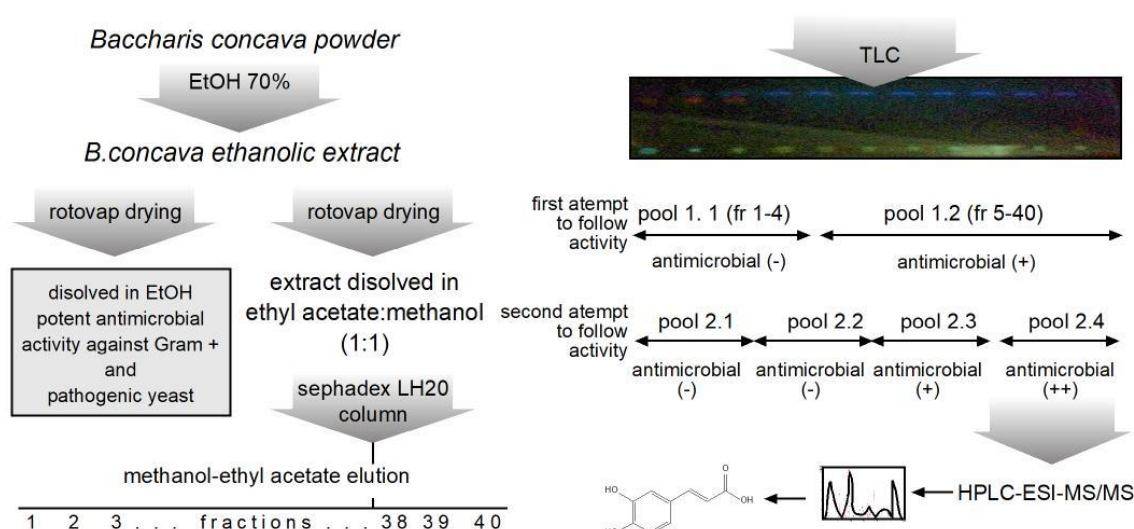


Figure 2. Schematic representation of steps followed from extraction to identification of molecules present in a fraction with antimicrobial activity. After confirming antimicrobial activity in the original hydroalcoholic extract of *B. concava*, dried extract was dissolved in ethyl acetate:methanol (50:50), fractionated in a Sephadex column, analyzed by TLC, and fractions pooled to test antimicrobial activity. The pool preserving the stronger antimicrobial effects was subjected to HPLC-ESI-MS/MS to identify phenolic compounds.

LC-MS analysis.

The *Baccharis* extract was examined on an LC-MS system consisting of the HPLC HP 1100 (Agilent Technologies Inc., CA-USA) coupled to an electrospray ion-trap mass spectrometer Esquire 4000 ESI-IT (Bruker Daltonik GmbH, Germany). For the HPLC separation, a Kromasil 100-5C18 250×4.6 mm, 5 µm, 100 Å column (Eka Chemicals AB, Sweden) was used, the column outlet was connected to a split that divided the flow to the UV detector and the mass spectrometer. The analysis was performed at room temperature by the injection of 20 µL of extract, at a flow rate of 1.0 mL/min. The mobile phase components were formic acid 0.1% v/v (component A) and methanol (component B) according to the following elution gradient: 0-5 min, 5% B; 5-7.5 min, 5-20% B; 7.5-20 min, 20-30% B; 20-40 min, 30-40% B; 40-45 min, 40-60% B; 45-50 min, 60-80% B; 50-55 min, 80% B; 55-57.5 min, 80-5% B; and 57.5-60 min, 5% B. The UV detection was performed at 254 nm. The ionization process (nebulization) by electrospray was performed at 3,000 V assisted by nitrogen as nebulizer gas at a pressure of 50 psi and flow rate of 10 L/min and assisted by nitrogen as drying gas at a temperature of 365°C. Chromatograms and mass spectra were acquired in positive and negative polarity. The trap parameters were set in ion charge control (ICC) using manufacturer default parameters, and maximum accumulation time of 200 ms. Collision induced dissociation (CID) was performed by collisions with the helium background gas present in the trap. Fragmentation was controlled by SmartFrag. All data obtained was analyzed using DataAnalysis 3.2 (Bruker Daltonik GmbH, Germany). The identification of compounds was carried out by precursor and fragmentation pattern comparison with a library from the *Unidad de Espectrometría de Masas* at Universidad de Chile. The identification of compounds was carried out by comparison of their precursors and corresponding fragmentation patterns with a library developed at the Mass Spectrometry Unit of the Universidad de Chile.

3. Results

Alcaloids, steroids and terpenes, flavonoids and phenolic compounds are present the hydroalcoholic extract of B. concave.

Before testing biological activity in the *B. concava*, the phytochemical composition was qualitatively studied. The extract obtained in 70% EtOH, resulted positive for Dragendorff,

Libermann-Burchard, aluminum chloride, and Ferric chloride tests. Indicating presence of detectable amount of alcaloids, steroids and therpenes, flavonoids, and phenolic compounds.

Hydroalcoholic extract of *B. concava* shows potent antimicrobial effects.

In a first attempt to find antimicrobial activity, extracts of *B. concava* were directly tested by the agar diffusion test on a lawn of *S. aureus* and *S. Typhimurium*. Because a discrete, but consistent activity was observed we decided to test antimicrobial effects on a battery of microorganism including Gram-positive, Gram-negative bacteria, and pathogenic yeast by the agar diffusion test. As shown in **Table 2**, second column, all Gram-positive bacteria were sensitive to the *B. concava* extract. In the opposite, all Gran-negative tested (*S. Typhimurium*, *K. pneumoniae* and *A. Baumannii*) were resistant to *B. concava*. The two pathogenic yeast, *C. albicans* and *C. neoformans*, were also susceptible to the *B. concava* extracts with *C. neoformans* as the most sensitive microorganism tested, measured as diameter of inhibitory haloes. Next, minimal inhibitory concentration was assessed to quantify the concentration capable of inhibiting at least half of growth (IC₅₀) of the tested microorganisms compared with the control condition. As shown in **Table 2**, third column, all tested microorganism, except for Gram-negative bacteria, were susceptible to treatment with *B. concava* extract.

Table 2. Antimicrobial effect of a *Baccharis concava* hydro-alcoholic extract.

Microorganism	Inhibition haloes (mm)	MIC-IC ₅₀ (mg/mL)	MBC (mg/mL)
<i>S. epidermidis</i>	30.67 ± 0.58	6.95 ± 3.01	13.89 ± 6.01
<i>S. aureus</i>	20.67 ± 0.58	2.17 ± 0.75	4.34 ± 1.51
<i>B. subtilis</i>	14.67 ± 1.16	13.89 ± 6.01	27.78 ± 12.03
<i>B. cereus</i>	14.67 ± 1.16	20.83 ± 0	41.67 ± 0
<i>S. pyogenes</i>	27.33 ± 0.58	6.95 ± 3.01	13.89 ± 6.01
<i>E. coli</i>	6 ± 0	-	-
<i>S. Typhimurium</i>	6 ± 0	83.33 ± 0	111.11 ± 48.12
<i>K. pneumoniae</i>	6 ± 0	-	-
<i>A. baumannii</i>	6 ± 0	-	-
<i>C. albicans</i>	14.33 ± 0.58	<27.78 ± 12.03	27.78 ± 12.03
<i>C. neoformans</i>	37 ± 0	41.67 ± 0	83.33 ± 0

Results are the average of three independent experiments ± standard deviation. MIC, minimal inhibitory concentration; MBC, minimal biocidal concentration. Strains were *S. epidermidis* ATCC 12228, *S. aureus* ATCC25923, **B. subtilis** ATCC 6633, *B. cereus* ATCC 11778, *S. pyogenes* ISP36900, *E. coli* ATCC25922, *S. enterica* sv. Typhimurium 14028s, *K. pneumoniae* ATCC 13883, *A. baumannii* ATCC 19606, *C. albicans* ATCC 90029, *C. neoformans* GM1.

Using the same samples of the IC₅₀ experiment, treated bacteria with different concentrations were used to prepare base-10 serial dilutions and plated to find the minimal biocidal concentration (MBC) defined as the concentration that kills at least 99,9% of microorganisms (**Table 2**, fourth column), compared to treatment with the vehicle. As expected, MBC were higher than MIC-IC₅₀, followed the same patterns, with *S. aureus* and *S. epidermidis* among the most susceptible microorganism when *B. concave* extract is used directly diluted in the culture media.

Defective lipopolysaccharide renders susceptibility to *B. concava* extract in *S. Typhimurium*.

As all tested Gram-negative bacteria were resistant to *B. concava* extract. two plausible hypotheses come to place. First, there is no active compounds against Gram-negative bacteria in *B. concava* extract or second, Gram-negative bacteria are resistant to active compounds present in *B. concava* extract. although these hypotheses are not mutually exclusive, the effects of altering the lipopolysaccharide and cell wall stability were tested, in regard of their effects on susceptibility to *B. concava* extract. Therefore, resistance of *S. Typhimurium* mutants in genes *rfaD*, *rfaE*, or *ompA* were exposed to *B. concava* extract. *S. Typhimurium rfaD* and *S. Typhimurium rfaE* resulted highly susceptible to *B. concava* extract. this result indicates that affecting the envelope (and therefore the permeability) of *Salmonella* sensitizes the bacteria to the active compound present in the extract. both *rfaD* and *rfaE* genes encode enzymes affecting synthesis of the LPS core, producing a severely defective LPS molecule. The *ompA* gene encodes for the outer membrane protein OmpA an outer membrane protein, important for anchorage and strength of cell wall. Mutant in *ompA* gene of *Salmonella*, did not sensitize the bacteria to the extracts.

Table 3. Mutants of Salmonella with defective LPS become susceptible to *B. concava* extract.

Strain derived from <i>S.Typhimurium</i>	Function/structure affected	Inhibition haloes (mm)
<i>S. Typhimurium</i> 14028s WT	wild type bacteria	6
<i>S. Typhimurium ΔrfaD</i>	intesis LPS	20
<i>S. Typhimurium ΔrfaE</i>	Sintesis LPS	22
<i>S. Typhimurium ΔOmpA</i>	outer membrane protein/channel	6

Mutants derived from *S. Typhimurium* 14028s (a standard strain) were obtained by facilitated allelic exchange and designed based in mutants obtained through a screening. The *rfaD* gene encodes an heptosyltransferase. The *rfaE* gene encodes a bifunctional enzyme. In both mutants, synthesis of LPS core is affected. The gene *ompA* encodes outer membrane protein implicated in transport through membrane and stabilization of cell envelope.

Column fractionation of *B. concava* extract.

Sephadex LH-20 was used to separate 300 mg of dried extract dissolved in 3 mL of methanol/ethyl acetate (1:1). The same solvent was used to elute the column. A total of 40 fractions of approximately 50 mL each were collected, resolved by TLC as described in material and methods and fractions pooled according to presence of similar patterns under UV light. As seen in Figure 2, in a first attempt to follow antimicrobial activity present in the pooled fractions, fractions 1 through 4 and 5 through 40 were pooled, to rescue antimicrobial in the bigger pool. Next, in a second attempt, fractions were pooled in lots of 10 consecutive fractions, this time the fourth pool (pool 2.4 in Figure 2) was the one with the higher antimicrobial activity measured against *S. aureus*. Therefore, pool 2.4 was selected for further chemical composition analysis.

HPLC/mass spectrometry identification of phenolic compounds.

Figure 3 shows the UV chromatogram at 254 nm obtained for the *B. concava* extract, pooled fraction 2.4. The identification of the labeled peaks is detailed in Table 4 which contains the precursors observed in negative polarity, as well as their corresponding fragmentations. The fragmentations are arranged by decreasing intensity from left (base peak) to right.

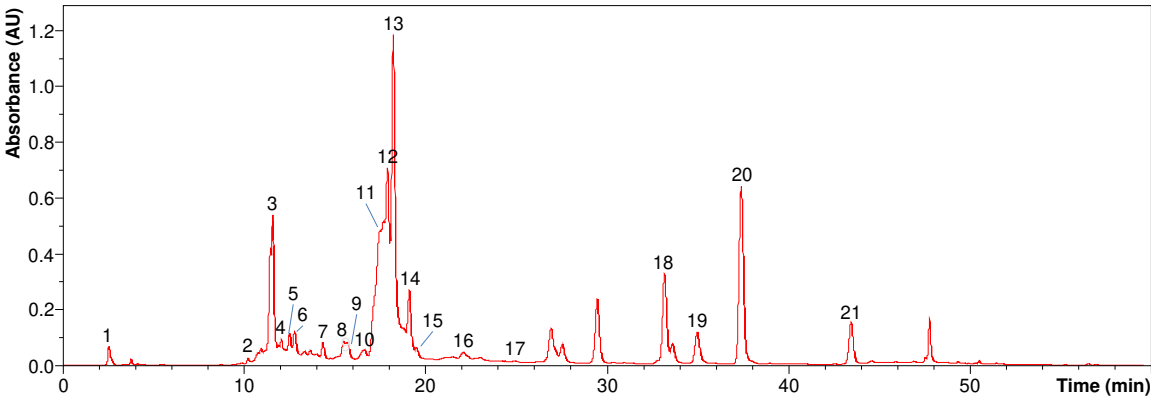


Figure 3. Phenolic profile of the most active fractions of the *Baccharis concava* extract. HPLC-ESI-MS/MS chromatogram identify phenolic compounds in a hydroalcoholic extract of *B. concava*. The extract presenting antimicrobial effects is rich in molecules derived from caffeoylquinic acid, quinic acid, coumaroyl acid and quercetin.

Table 4. identification of phenolic compounds in a *B. concava* extract presenting antimicrobial activity.

Pe ak	RT (min)	[M-H]- (m/z)	Fragments MS2 (m/z)					Compound
1	2.5	192.1	172.4	126.3	84.4	92.3	110.4	Quinic acid
2	10.9	355.9	190.5	178.5	134.4			Coumaroylhexaric acid
3	11.6	353.2	172.6	178.5	190.5			Caffeoylquinic acid
	11.6	593.7	473.6	503.3	407.8	575.1		Apigenin-di-C-hexoside
4	12.2	354.3	190.5	179.0				Caffeoylquinic acid
5	12.8	741.6	300.0	609.2	591.3	475.1	343.1	Rutin-O-pentoside
6	13.3	352.2	190.5	178.4				Caffeoylquinic acid
7	14.3	609.5	300.8					Quercetin-O-rhamnosyl exoside
8	15.4	463.6	300.8					Quercetin-O-hexoside
9	15.6	477.2	300.9					Quercetin-O-glucuronide
10	16.5	515.4	352.9					Dicaffeoylquinic acid
11	17.6	515.1	352.9	190.8	202.8	178.7		Dicaffeoylquinic acid
12	17.8	515.1	352.9					Dicaffeoylquinic acid
13	18.2	515.2	352.9	190.7	178.9			Dicaffeoylquinic acid
14	19.0	515.1	352.9					Dicaffeoylquinic acid

15	19.4	516.6	352.9	202.6	335. 1	19 0.5	17 2.5	Dicafeoylquinic acid
16	22.0	529.4	353.1	366.9	190. 8	17 8.7		Caffeoyl-feruloylquinic acid
17	24.9	677.9	515.0					Dicafeoylquinic acid-O-exoside
18	33.1	269.1	224.7	148.5	200. 6			Apigenin
19	34.9	329.1	313.9					Kaempferol methoxy methyl ether
20	37.3	300.0	283.9					Kaempferol methyl ether
21	43.4	313.8	297.2					Kaempferol dimethoxy

As seen from the table 4, three isomers of caffeoylquinic acid (m/z 353, $[M-H]^-$) were observed at t_R 11.6 min (peak 3), t_R 12.2 min (peak 4) and t_R 13.3 min (peak 6). Peaks 10-15 were identified as isomers of dicafeoylquinic acid as indicated by the observation of signal m/z 515 in negative polarity ($[M-H]^-$) and m/z 499 in positive polarity ($[M-H_2O+H]^+$). Peak 17 (t_R 24.9 min) would correspond to caffeoylquinic acid-O-hexoside as suggested by the observation of signal m/z 677 and its fragment m/z 515. Among the flavones, apigenin was identified at t_R 33.1 min (peak 18) through signal m/z 269 in negative polarity ($[M-H]^-$) and m/z 271 in positive polarity ($[M+H]^+$); and apigenin-di-C-hexoside (probably vicenin 2) at t_R 11.6 min (peak 3) on the basis of signal m/z 593 ($[M-H]^-$) and its corresponding fragmentation. Among the flavonols, several quercetin derivatives were identified as: peak 5 (t_R 12.8 min) identified as rutin-O-pentoside based on the signal m/z 741 ($[M-H]^-$) and its fragmentation; peak 7 (t_R 14.3 min) which was identified as quercetin-O-rhamnosyl hexoside based on the signal m/z 609 ($[M-H]^-$) and m/z 611 ($[M+H]^+$) together with its fragmentations; peak 8 (t_R 15.4 min) identified as quercetin-O-hexoside according to the observation of signal m/z 463 ($[M-H]^-$) and m/z 465 ($[M+H]^+$); and peak 9 (t_R 15.6 min) which was identified as quercetin-O-glucuronide based on the signals m/z 477 ($[M-H]^-$) and m/z 479 ($[M+H]^+$) and their corresponding fragmentations. Several kaempferol derivatives were also identified: peak 19 (t_R 34.9 min) identified as kaempferol methoxy methyl ether based on the signals m/z 329 ($[M-H]^-$) and m/z 331 ($[M+H]^+$); peak 20 (t_R 37.3 min) which according to the signal m/z 300 ($[M-H]^-$) and its fragmentation would correspond to kaempferol methyl ether (probably kaempferide); and peak 21 (t_R 43.4 min) identified as dimethoxy kaempferol according to the signal m/z 313 ($[M-H]^-$) and its corresponding fragmentation. Other compounds identified were quinic acid (peak 1), coumarylhexaric acid (peak 2) and caffeoyl feruloylquinic acid (peak 16).

4. Discussion

This work describes potent antimicrobial activity, of a hydroalcoholic extract of *B. concava* against Gram-positive bacterium, *C. albicans*, and *C. neoformans*. Previously, antimicrobial activity was reported against *S. aureus* in essential oil prepared from *B. concava*. In the same line with our results, the essential oil derived from *B. concava* was inactive against the Gram -negative bacteria *E. coli*. it is possible that the active molecules present in the *B. concava* are inactive against Gram-positive. However, some *Baccharis* have presented discrete activity against Gram-negatives, such is the case of *Baccharis revoluta*, although its activity against Gram-positive is higher (Rodríguez A et al., 2016). Whether the extracts of *B. concava* are inactive against Gram-negative or Gram-negative are just highly resistance to *B. concava* extracts was assessed by using *S. Typhimurium* mutants. Increased permeability turned *S. Typhimurium* susceptible to *B. concava* extracts. Therefore, we speculate that active molecules in the tested extract poorly penetrates the bacterial envelope of Gram-negatives with

an intact LPS. Moreover, new evidence indicates that this kind of mutants accumulate oxidative species, that in turn may facilitate killing by antimicrobial agents (Seregina et al., 2022).

Few reports exist about antimycotic effects within the *Baccharis* genus, and none are published for *B. concava*, in this study we found potent activity against pathogenic yeast *C. albicans* and *C. neoformans*. Previous report indicates some discrete to poor antifungal activity in some *Baccharis* species, however some efforts were made to evaluate synergistic effects of different *Baccharis* extracts in combination with terbinafine against *Trichophyton rubrum*. Some promising synergistic and additive effect were reported in regard of chemical composition of extracts (Rodriguez et al., 2013). Further studies, in regard of antimycotic activity, will be important in the light of emergent yeast and fungi with multiresistant phenotypes (Arendrup and Patterson, 2017).

This study describes the phytochemical composition of a polar extract of *B. concava*. Fifteen compounds were identified and among them fourteen are described for the first time in this specie. These compounds were diterpenoids, flavonoids, and chlorogenic acid derivatives, which are natural products frequently found in *Baccharis* species. The antimicrobial effects of polar compounds such as phenolic acids, flavonoids and heteroside derivatives have been reported. An example is chlorogenic acid, a compound found in extracts of *B. concava* and from which its antimicrobial activity has been reported, probably as a disruptor of bacterial membranes, therefore, disrupting cellular homeostasis (Lou et al., 2011). Other molecules described in the ethanolic extract of *B. concava* are various caffeic acid-derivative molecules. Because of their abundance, pure commercial caffeic acid was tested for antimicrobial activity, against *S. aureus*, using 6mm filter disks impregnated with 0.3 or 0.5 mg. The effect of pure caffeic acid was discrete with haloes ranging from 8 to 9 mm, compared with 6mm in the DMSO control (data not shown). It is plausible that the blend of caffeic acid-derivatives and other molecules account for the potent antimicrobial activity observed in ethanolic *B. concava* extract.

B. concava has been used in traditional medicine for wound healing, as diuretic and as a tonic in beverages. It is possible that its antimicrobial effects accounts for its curative effect on wounds. The antimicrobial effects, found in the ethanolic extract, are exclusive to Gram-Positive bacteria and pathogenic yeast, with no effect over Gram-negative bacteria. It would be interesting to continue learning about the molecules responsible for the spectrum in the ethanolic extract of *B. concava*, which in the future might allow developing more specific therapies with less unwanted effects on the Gram-negative microbiota.

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