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Research Article

Antifungal and Antibacterial Metabolites from a French Poplar Type Propolis

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During this study, the *in vitro* antifungal and antibacterial activities of different extracts (aqueous and organic) obtained from a French propolis batch were evaluated. Antifungal activity was evaluated by broth microdilution on three pathogenic strains: *Candida albicans*, *C. glabrata*, and *Aspergillus fumigatus*. Antibacterial activity was assayed using agar dilution method on 36 Gram-negative and Gram-positive strains including *Staphylococcus aureus*. Organic extracts showed a significant antifungal activity against *C. albicans* and *C. glabrata* (MIC₈₀ between 16 and 31 μ g/mL) but only a weak activity towards *A. fumigatus* (MIC₈₀ = 250 μ g/mL). DCM based extracts exhibited a selective Gram-positive antibacterial activity, especially against *S. aureus* (SA) and several of its methicillin-resistant (MRSA) and methicillin-susceptible (MSSA) strains (MIC₁₀₀ 30–97 μ g/mL). A new and active derivative of catechin was also identified whereas a synergistic antimicrobial effect was noticed during this study.

1. Introduction

Propolis is a resinous natural substance collected by honeybees from buds and exudates of various trees and plants, mixed with beeswax and salivary enzymes. Bees generally use it as a sealer, to smooth out the internal walls of the hive, as well as a protective barrier against intruders. Propolis has been used in folk medicine since ancient times due to its pharmacological potential associated with antioxidant [1–3], antifungal [4, 5], antibacterial [6–8], and anti-inflammatory [9] properties.

Propolis is generally composed of 50% of resin and balm (including polyphenolic compounds), 30% of wax and fatty acids, 10% of essential oils, 5% of pollen, and 5% of various organic and inorganic compounds. However, the composition of propolis deeply depends on the vegetation at the site of collection [10]. Indeed, propolis from temperate climatic zones, like in Europe, North America, or nontropical regions of Asia, mainly originates from the bud exudates

of *Populus* species (*Salicaceae*) and consequently is rich in flavonoids and phenolic acids and their esters [11]; however tropical propolis, originating from regions where neither poplars nor birches grow, is rich in prenylated derivatives of *p*-coumaric acids, benzophenones, or terpenoids [12, 13].

The antifungal, antibacterial properties and chemical composition of propolis from many countries all over the world have been widely studied [6, 8, 14–20] but few reports were already given for European propolis [21, 22]. In 1990, Grange and Davey [23] highlighted for the first time the bactericidal activity of a French propolis against Grampositive strains whereas later on, in 2000, Hegazi et al. [22] could associate this antibacterial activity with the presence of benzyl caffeate, pinocembrin, and *p*-coumaric acid.

During a previous study, we have evaluated the antioxidant and anti-AGEs activities of different solvents extracts [water; 95% EtOH; 70% EtOH; MeOH; dichloromethane (DCM) and DCM/MeOH/ H_2O (31/19/4)] obtained from a French propolis batch and identified their active constituents

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[24]. Here we have investigated the *in vitro* antifungal and antibacterial activities of these extracts. The antifungal activity was studied on three fungal strains (two yeasts, *Candida albicans* and *C. glabrata*, and one filamentous fungus, *Aspergillus fumigatus*). 36 strains of Gram-positive (including *Staphylococcus aureus*) and Gram-negative (including *Escherichia coli*) bacteria were used for the antibacterial assays. During this study, a new secondary metabolite was isolated, namely, 8-[(E)-phenylprop-2-en-1-one]-5-methoxy-(±)-catechin.

2. Materials and Methods

2.1. Reagents and Standards. Formic acid, p-coumaric acid, ferulic acid, isoferulic acid, 3,4-dimethoxycinnamic acid, and prenyl caffeate were purchased from Sigma-Aldrich (L'Isle d'Abeau Chesnes, Saint-Quentin-Fallavier, France). Caffeic acid and chrysin were obtained from Acros Organics (Geel, Belgium). Galangin was purchased from Extrasynthese (Genay, France). Pinocembrin and pinobanksin-3-acetate were isolated from the DCM extract of propolis.

2.2. Instrumentation. Optical rotation was measured on a JASCO P-2000 polarimeter. IR spectra were recorded on a Bruker Vertex 70 spectrophotometer. NMR spectra (1D and 2D) were recorded on a Bruker Avance spectrometer at 500 MHz for ¹H and 125 MHz for ¹³C. MS analyses were performed on an ESI/APCI Ion Trap Esquire 3000+ from Bruker. UV absorbances were obtained from a Tecan Infinite M200 microplate spectrophotometer.

2.3. Propolis Samples. In order to analyze a typical French batch, that is, exhibiting an average chemical composition, a mixture of samples (10 g of each), collected in apiaries originating from different regions of France, was used for this study. These samples were provided by "Ballot-Flurin Apiculteurs," a company specialized in organic beekeeping. Indeed, even collected in the same geographical region, propolis profiles may differ between apiaries and even inside the same apiary from one hive to another one [25]. Keeping in mind any potential economic development, it then appeared more appropriate to study an industrial end-product, that is, a mixture, exhibiting an average chemical composition associated with an average antimicrobial activity, rather than a specific sample. Therefore, 24 batches of propolis collected over two years (2010 and 2011) from different places in France (cf. supporting information 1; see Supplementary Material available online at http://dx.doi.org/10.1155/2015/319240) were homogeneously mixed to undergo this study.

2.4. Extractions. The extraction processes have been already described [24]. Briefly, the propolis batch was homogeneously pulverized in the presence of liquid nitrogen and divided into 1 g samples. Four different extractions were then carried out on 1 g samples with water (E1), 95% EtOH (E2), 70% EtOH (E3), and MeOH (E4). Then, two extractions, preceded by a cyclohexane wax elimination, were independently performed on 1 g samples with DCM (E5) and

a mixture of DCM, MeOH, and $\rm H_2O$ (31/19/4) (E6). For E1, a decoction of 1g of propolis powder was boiled in 20 mL $\rm H_2O$ at 100°C for 15 min. After cooling, the solidified wax and the residue were removed by filtration, and the filtrate was evaporated to dryness. For other solvents, 1g of propolis powder (or residue obtained from a previous extraction) was macerated in 3 × 20 mL of solvent. After stirring for 3 × 2 h at room temperature, the mixture was filtered. The filtrates were gathered and evaporated under vacuum. Extraction yields (dried extract/100 g) were as follows: E1 7%; E2 68%; E3 65%; E4 68%; E5 50%; and E6 59%.

2.5. Antifungal Activity. Antifungal activity was assayed on human pathogenic fungi, including two common yeasts (Candida albicans ATCC 66396 and C. glabrata LMA 90-1085) and an opportunistic mould (Aspergillus fumigatus CBS 11326). The strains were obtained from the Parasitology and Mycology Laboratory at the University Hospital Center of Angers, France. Microorganisms were cultivated at 37°C on yeast extract-peptone-dextrose-agar (YPDA) containing 0.5 g/L chloramphenicol for two (*C. albicans* and *C. glabrata*) or three (A. fumigatus) days. Tests were performed according to a procedure described by Alomar et al. [27], following the guidelines of the approved reference method of the National Committee for Clinical Laboratory Standards (NCCLS) for yeasts [28] and filamentous fungi [29]. Briefly, the yeast suspensions were prepared in RPMI-1640 culture medium and adjusted spectrophotometrically at 630 nm to reach a final concentration of ca. 0.5×10^3 to 2.5×10^3 cells/mL. The tests were performed using sterile 96 flat shaped well microtiter plates. Serial twofold sample dilutions were made in DMSO. Sample solutions were dispensed at a volume of $5 \,\mu$ L in triplicate into the wells to obtain final concentrations from 250 to 1.95 μ g/mL. After 48 h at 37°C for *C. albicans* and C. glabrata and 72 h for A. fumigatus, the spectrophotometric MIC endpoint was calculated from the turbidimetric data as the lowest sample concentration causing a growth inhibition equal to or greater than 80% of the control (MIC₈₀). Amphotericin B was used as a positive control.

2.6. Antibacterial Activity. Antibacterial activity was evaluated on 36 human pathogenic bacterial strains collected by the Laboratory of Bacteriology at the University Hospital Center of Angers, France: seven strains of Acinetobacter baumannii (RCH, SAN008, 12, AYE, CIP7034, 107292, and 5377), two of Escherichia coli (ATCC25922 and a clinical isolate), three of Pseudomonas aeruginosa (ATCC27853 and two clinical isolates), and 4 clinical isolates of Enterobacter cloacae, E. aerogenes, Klebsiella oxytoca, and Salmonella enteritidis (phage type 4) for Gram-negative bacteria; thirteen strains of Staphylococcus aureus (ATCC25923, six methicillinsusceptible clinical isolates, six methicillin-resistant clinical isolates), two clinical isolates of S. epidermidis (methiS and methiR), three clinical isolates of Enterococcus faecalis and one of *E. faecium*, and one clinical isolate of *Corynebacterium* striatum for Gram-positive bacteria. Tests were performed using the methodology described by Alomar et al. [30]. Briefly, a stock solution of each sample was prepared in triplicate at 20 mg/mL in DMSO under sterile conditions. Serial dilutions were prepared (sample concentrations: 10, 20, 30, etc., to $100~\mu g/mL$) and 0.1~mL of each dilution was added to 19.9 mL of Mueller-Hinton agar (Merck, Germany) and transferred to Petri plates. Bacterial strains ($2 \times 10^4~CFU/mL$) were suspended in sterile NaCl aqueous solution (0.15~M) and inoculated on the different Petri plates using the multipoint inoculator (AQS, England). After 24 h of incubation at 37°C, the minimum inhibitory concentration (MIC $_{100}$, $\mu g/mL$) against bacterial strains was defined as the lowest concentration of each sample that inhibited visible growth. A blank was made inoculating the strains on Mueller-Hinton agar without any extract or compound. Oxacillin was used to distinguish the methicillin-resistant from the susceptible staphylococcal strains.

2.7. HPLC-DAD and HPLC-MS Procedures. Dry extracts were dissolved in MeOH (5 mg/mL for the aqueous extract and 10 mg/mL for the organic solvents ones) and centrifuged at 13000 rpm for 10 min prior to injection (10 μ L) into the HPLC system. Analytical HPLC was run on a 2695 Waters separation module equipped with a diode array detector 2996 Waters. Separation was achieved on a LiChrospher column 100 RP-18 (125 × 4 mm i.d., 5 μ m) protected with a LiChrocart 4-4 guard cartridge (4 × 4 mm i.d.) at a flow rate of 1 mL/min. The mobile phase consisted of 0.1% formic acid in water (solvent A) and MeOH (solvent B) and the separation was performed using the linear gradient: 25–100% B in 40 min. UV detection was achieved at two wavelengths: 254 and 280 nm.

The mass analyses were performed with an ESI interface coupled to an ion trap mass analyzer in both positive and negative modes, with the following conditions: collision gas, He; collision energy amplitude, 1.3 V; nebulizer and drying gas, N_2 , 7 L/min; pressure of nebulizer gas, 30 psi; dry temperature, 340°C; flow rate, 1.0 mL/min; solvent split ratio 1:9; scan range, m/z 100–1000.

2.8. Identification of Propolis Constituents. 18 and 22 were directly identified in the DCM extract by HPLC/UV/MS and comparison with the literature data [31, 32], whereas 3, 6, 7, 8, 10, and 32 were compared with authentic standards (Sigma-Aldrich and Acros organics, cf. Section 2.1). A flash chromatography was then carried out in order to identify the other phenolic constituents. 50.0 g of pulverized propolis was firstly extracted with cyclohexane $(3 \times 200 \text{ mL}, 2 \text{ h}, \text{ room})$ temperature) to eliminate waxes. After filtration, the residue was extracted with DCM (5×200 mL, 2 h, room temperature) to give 25.0 g of dry DCM extract (50% yield). 21.0 g of this extract was fractionated using a CombiFlash Teledyne ISCO apparatus and a prepacked silica gel column (Interchim PF-50SI HC/300 g, 50 μ m), at a flow rate of 100 mL/min and with the following gradient elution system: cyclohexane (C_6H_{12}) 100% (2.0 L), C_6H_{12} : EtOAc 90:10 (1.7 L), C_6H_{12} : EtOAc 90:10 to 80:20 (2.2 L), 80:20 to 70:30 (2.5 L), C_6H_{12} : EtOAc 70:30 to 60:40 (2.2 L), and C_6H_{12} : EtOAc 60:40 to 50:50 (3.0 L) then DCM: MeOH 96:4 (2.2 L). UV detection (λ 254 and 280 nm) and TLC monitoring allowed collecting

21 fractions (F1-21). 48 [33] was identified in F1, 14 [34] and 17 [31] were identified in F11, 1 and 2 [19] were identified in F13, and finally 23 [31] was identified in F15 by HPLC/UV/MS and comparison with the literature data. The remaining constituents were isolated and identified through 1D and 2D NMR analysis (cf. Section 2.2). 200 mg of F1 was chromatographed on a silica gel column (Grace, 24 g) by flash chromatography at a flow rate of 25 mL/min with a mixture of C₆H₁₂ and EtOAc (B) [gradient: 1% B (30 min), 2% B (5 min), 2-5% B (2 min), 5% B (2 min), 5-30% B (1 min), and 30% B (5 min)] to give 46 [33, 35] (5 mg), 47 [33, 36] (3 mg), and **49** [37] (5 mg). F2 (1.5 g) gave **43** [38] whereas F4 (696 mg) and F5 (384 mg) allowed us to, respectively, identify 44 [37, 39] and 45 [40]. 500 mg of F6 (1.6 g) was chromatographed on reverse-phase- (RP-) Flash chromatography (Interchim column PF-30C18 HC/6 g, 30 μ m) at a flow rate of 15 mL/min with water and MeOH (B) [gradient: 25–30% B (20 min), 30– 40% B (2 min), 40% B (8 min), 40–45% B (1 min), 45% B (12 min), 45–50% B (2 min), and 50% B (20 min)] to give 9 [31] (10 mg), 25 [38] (126 mg), and a mixture of 35 and 36 [34] (15 mg). Similarly 500 mg of F7 was fractionated [gradient: 30-50% B (25 min), 50-60% B (25 min), and 60-65% B (20 min)] to give 11 [41] (2 mg), 33 [31, 42] (128 mg), 34 [38] (65 mg), and **42** [31] (41 mg) whilst 500 mg of F8 [gradient: 30-45% B (20 min), 45% B (30 min), 45-48% B (5 min), 48-55% B (5 min), 55% B (8 min), 55-60% B (1 min), and 60% B (12 min)] gave 13 [38] (3 mg), 28 [31, 43] (224 mg), and 41 [34] (28 mg). 500 mg of F9 [gradient: 30–60% B (50 min) and 60-65% B (5 min)] yielded mixture of 4 [34] and 5 [44, 45] (8 mg), **15** [43] (33 mg), **24** [32, 46] (16 mg), **32** [38] (150 mg), and another mixture of 37 [16] and 38 [32] (3 mg). 27 [34] and **29** [31] were identified from *F10* (1.3 g). 500 mg of *F11* [flow rate of 20 mL/min, gradient: 25-75% B (55 min)] gave 26 [47] (82 mg), **31** [31] (56 mg), and **39** [32] (70 mg). 300 mg of *F13* [flow rate of 15 mL/min, gradient: 30–45% B (20 min), 45% B (20 min), 45–50% B (10 min), 50–60% B (5 min), and 60% B (5 min)] gave **10** [31] (2 mg) and **16** [38] (3 mg). **30** [48] was directly identified in F17 (543 mg). 500 mg of F18 [gradient: 40% B (7 min), 40–50% B (2 min), and 50% B (30 min)] gave a mixture of 8 and 12 (cf. F19) together with the new compound **40** (12 mg). 500 mg of *F19* was chromatographed [gradient: 25% B (25 min), 25–35% B (1 min), and 35% B (20 min)] to give 8 (8 mg) and 12 (26 mg) [31]. Finally 500 mg of F20 allowed us to isolate [gradient: 30-40% B (20 min), 40-43% B (20 min), and 43-50% B (10 min)] 19 [32] (3 mg) and a mixture of 20 [49] and 21 [50] (7 mg).

3. Results and Discussion

3.1. Antifungal and Antibacterial Activities. Table 1 shows the minimum inhibitory concentration of at least 80% of fungal growth (MIC $_{80}$) obtained with E1–6 extracts for Candida albicans, C. glabrata, and Aspergillus fumigatus. E1 did not exhibit any interesting antifungal activity (MIC $_{80}$ > 250 µg/mL for the three strains) whereas E2–6 showed significant antifungal activities (MIC $_{80}$ between 16 and 31 µg/mL) on both C. albicans and C. glabrata. These results are in agreement with those previously obtained for an Argentinian propolis

Amphotericin B

| - | | | | | | | |
|---------|----------------|--------------------------|-------------|--------------|--|--|--|
| Extract | Solvent | C. albicans | C. glabrata | A. fumigatus | | | |
| | Solvent | MIC_{80} (μ g/mL) | | | | | |
| E1 | H_2O | >250 | >250 | >250 | | | |
| E2 | 95% EtOH | 31.25 | 15.63 | 250 | | | |
| E3 | 70% EtOH | 31.25 | 31.25 | 250 | | | |
| E4 | MeOH | 31.25 | 31.25 | 250 | | | |
| E5 | DCM | 31.25 | 31.25 | 250 | | | |
| E6 | Mixed solvents | 15.63 | 31.25 | 250 | | | |

TABLE 1: Antifungal activity against Candida albicans, C. glabrata, and Aspergillus fumigatus.

on several *Candida* species (MIC₁₀₀ in a range of 31 to 125 μ g/mL) [51] as well as with Greece and Cyprus ones (MIC₁₀₀ 20 μ g/mL) [21]. E2–6 also exhibited a weak activity towards *A. fumigatus* (MIC₈₀ 250 μ g/mL).

0.125

0.125

6

According to Ríos and Recio [52] a MIC $_{100}$ < 100 μ g/mL should be considered as a promising value for a crude extract (versus 10 μ g/mL for pure compounds). This is the reason why Table 2 gives the results of the antibacterial activity of E1–6 at the concentration of 100 μ g/mL for 28 strains of Gramnegative and Gram-positive bacteria.

Results showed that Gram-negative bacteria were not susceptible to E1-6 at this concentration. In contrast, organic solvents extracts were active on several Gram-positive bacteria such as Corynebacterium striatum (sometimes involved in pleuropulmonary infections) (E2–5) and especially *Staphy*lococcus aureus, including for the latter several methicillinresistant (MRSA) and methicillin-susceptible (MSSA) clinical isolates (E5-6). Sometimes called "golden staph," S. aureus is the most pathogenic species of Staphylococcus genus. It might cause food poisoning, skin infections, abscesses, and diseases like pneumonia, meningitis, and sepsis. S. aureus is additionally one of the major causes of hospital-acquired infections, and the treatment of some multiresistant strains has become quite problematic. Among them, MRSA appears in France as one of the most commonly multiresistant strains encountered in hospitals.

 ${
m MIC_{100}}$ of El–6 were determined on the 6 susceptible Gram-positive strains as well as on 8 other MRSA and MSSA strains. Results are given in Table 3.

El did not show any interesting activity on the 14 studied strains (MIC $_{100}$ > 100 μ g/mL). E2–6 showed interesting activities against *Corynebacterium striatum* with MIC $_{100}$ ranging from 63 to 90 μ g/mL. E5 and E6 exhibited the best antibacterial activities against the *Staphylococcus* strains with MIC $_{100}$ up to 57 and 30 μ g/mL, respectively. Among the alcoholic extracts, only E4 showed a moderate activity (MIC $_{100}$ 90 μ g/mL) against *S. aureus* and one MRSA whereas E2 and E3 appeared as inactive. These overall activities therefore appeared to be better than those reported by Grange and Davey for the antibacterial activity of a French propolis on *S. aureus* and MRSA (MIC $_{100}$ 188–375 μ g/mL) [23]. Our global antibacterial activity against MRSA and MSSA could be compared with those reported for propolis collected in Solomon Islands, exhibiting MIC $_{100}$ between

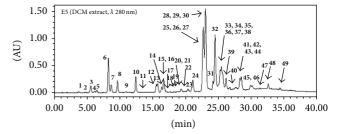


FIGURE 1: HPLC chromatograms of E5: 1 3,4-dihydroxybenzaldehyde, 2 4-hydroxybenzoic acid, 3 caffeic acid, 4 vanillin, 5 4hydroxyacetophenone, 6 p-coumaric acid, 7 ferulic acid, 8 isoferulic acid, 9 benzoic acid, 10 3,4-dimethoxycinnamic acid, 11 3phenylpropanoic acid, 12 pinobanksin-5-methyl ether, 13 cinnamic acid, 14 4-methoxycinnamic acid, 15 pinobanksin, 16 naringenin, 17 quercetin, 18 quercetin-3-methyl ether, 19 pinocembrin-5methyl ether, 20 1,3-di-p-coumaroylglycerol, 21 1-p-coumaroyl-3feruloylglycerol, 22 kaempferol, 23 apigenin, 24 cinnamylidene acetic acid, 25 pinocembrin, 26 benzyl caffeate, 27 isopent-3-enyl caffeate, 28 pinobanksin-3-acetate, 29 prenyl caffeate, 30 2-acetyl-1,3-dicoumaroylglycerol, 31 phenylethyl caffeate (CAPE), 32 chrysin, 33 benzyl p-coumarate, 34 galangin, 35 benzyl ferulate, 36 prenyl ferulate, 37 kaempferide, 38 rhamnocitrin, 39 cinnamyl caffeate, 40 8-[(E)-phenylprop-2-en-1-one]-5-methoxy-(±)-catechin (new), 41 cinnamyl isoferulate, 42 cinnamyl p-coumarate, 43 pinostrobin, 44 alpinone-3-acetate, 45 tectochrysin, 46 benzyl cinnamate, 47 cinnamyl benzoate, 48 cinnamyl cinnamate, and 49 cinnamyl cinnamylidene acetate.

64 and 128 μ g/mL [6]. Similarly E4 was more active than a methanolic propolis extract from Jordan (585 μ g/mL against *S. aureus* and 4700 μ g/mL against MRSA) [20].

These results suggested that antifungal and antibacterial activities of propolis extracts could be related to their flavonoids contents [24]. Indeed, whereas E1–6 exhibited high total polyphenol contents (239–281 mg GAE/g), only those showing both high flavone/flavonol (FF) and flavanone/dihydroflavonol (FD) contents (i.e., E5–6) were active on the studied strains. In addition the higher the cumulative contents FF+FD were, the stronger the antibacterial activity was, as shown with E5 (254 mg/g) and E6 (236 mg/g) > E2–4 (220–228 mg/g). These results are in agreement with those reported by Velazquez et al. [15] for different Mexican propolis collected in Sonora State where EEP from the areas of Ures (410 mg/g), Caborca (332 mg/g), and Pueblo de Alamos (209 mg/g) showed MIC₁₀₀ against *S. aureus* of 100, 200, and >400 μ g/mL, respectively.

3.2. Chemical Composition. Figure 1 shows the HPLC chromatogram of the DCM extract E5. 48 compounds were identified by comparison with the literature data (UV/MS) and pure standards or, when needed, through ¹H and ¹³C (1D and 2D) NMR analysis after compound isolation.

Additionally a new flavan-3-ol was identified as the 8-[(E)-phenylprop-2-en-1-one]-5-methoxy-(\pm)-catechin **40** (Figure 2).

Compound 40 was obtained as a yellow amorphous solid (0.6 μ g/g of DCM extract). The molecular formula was

 ${\it Table 2: Antibacterial\ activity\ of\ El-6\ against\ 28\ Gram-negative\ and\ Gram-positive\ strains.}$

| Number | Bacterial strains | Extracts (100 μ g/mL) | | | | | | |
|--------|--|---------------------------|--------------|--------------|----------|---------|--------------------|--|
| rumber | Dacterial strains | E1: H ₂ O | E2: 95% EtOH | E3: 70% EtOH | E4: MeOH | E5: DCM | E6: mixed solvents | |
| | Gram-negative: | | | | | | | |
| 1 | Acinetobacter baumannii (RCH) | _ | _ | _ | - | _ | _ | |
| 2 | Acinetobacter baumannii (SAN008) | _ | _ | _ | - | _ | _ | |
| 3 | Acinetobacter baumannii (12) | _ | _ | _ | - | _ | _ | |
| 4 | Acinetobacter baumannii (AYE) | _ | _ | _ | - | _ | _ | |
| 5 | Acinetobacter baumannii (CIP7034) | _ | _ | _ | _ | _ | _ | |
| 6 | Acinetobacter baumannii (CIP107292) | _ | _ | _ | _ | _ | _ | |
| 7 | Acinetobacter baumannii (CIP5377) | _ | _ | _ | _ | _ | _ | |
| 8 | Enterobacter cloacae (0705A1743) | _ | _ | _ | _ | _ | _ | |
| 9 | Enterobacter aerogenes (0705A0867) | _ | _ | _ | _ | _ | _ | |
| 10 | Escherichia coli (ATCC25922) | _ | _ | _ | _ | _ | _ | |
| 11 | Escherichia coli (0705A0434) | _ | _ | _ | _ | _ | _ | |
| 12 | Klebsiella oxytoca (0705C0187) | _ | _ | _ | _ | _ | _ | |
| 13 | Pseudomonas aeruginosa (ATCC27853) | _ | _ | _ | _ | _ | _ | |
| 14 | Pseudomonas aeruginosa (0704C0134) | _ | _ | _ | _ | _ | _ | |
| 15 | Pseudomonas aeruginosa (0703C0259) | _ | _ | _ | _ | _ | _ | |
| 16 | Salmonella enteritidis (4) | _ | _ | _ | _ | _ | _ | |
| | Gram-positive: | | | | | | | |
| 17 | Corynebacterium striatum (56) | _ | + | + | + | + | + | |
| 18 | Enterococcus faecalis (11003508001) | _ | _ | _ | _ | _ | _ | |
| 19 | Enterococcus faecalis (11003492701) | _ | _ | _ | _ | _ | _ | |
| 20 | Enterococcus faecalis (11004774001) | _ | _ | _ | _ | _ | _ | |
| 21 | Enterococcus faecium (11502441401) | _ | _ | _ | _ | _ | _ | |
| 22 | Staphylococcus aureus (ATCC25923) | _ | _ | _ | + | + | + | |
| 23 | MRSA (0706C0025) | _ | _ | _ | + | + | + | |
| 24 | MRSA (0702E0196) | _ | _ | _ | _ | + | + | |
| 25 | MSSA (0703H0036) | _ | _ | _ | _ | _ | _ | |
| 26 | MSSA (0701A0095) | _ | _ | _ | _ | + | + | |
| 27 | S. epidermidis methiS ^a (12004523201) | _ | _ | _ | _ | _ | _ | |
| 28 | S. epidermidis methiR ^b (12552599902) | _ | _ | _ | _ | _ | _ | |

 $^{-:} no\ antibacterial\ activity, +: antibacterial\ activity, {}^amethic illin-Susceptible, {}^bmethic illin-Resistant.$

Table 3: MICs of E1–6 against 14 Gram-positive strains including MRSA and MSSA.

| Number | Bacterial strains | MIC ₁₀₀ (μg/mL) | | | | | | |
|----------|-----------------------------------|----------------------------|--------------|--------------|-------------|-------------|--------------------|-----------|
| Nullibei | | E1: H ₂ O | E2: 95% EtOH | E3: 70% EtOH | E4: MeOH | E5: DCM | E6: mixed solvents | Oxacillin |
| 17 | Corynebacterium striatum | >100 | 83 ± 6 | 90 ± 0 | 77 ± 12 | 63 ± 15 | 87 ± 21 | _ |
| 22 | Staphylococcus aureus (ATCC25923) | >100 | >100 | >100 | 90 ± 0 | 60 ± 10 | 67 ± 15 | ≤0.25 |
| 23 | MRSA (0706C0025) | >100 | >100 | >100 | 90 ± 0 | 57 ± 12 | 30 ± 0 | ≥4 |
| 24 | MRSA (0702E0196) | >100 | >100 | >100 | >100 | 80 ± 10 | 77 ± 23 | ≥4 |
| 25 | MSSA (0703H0036) | >100 | >100 | >100 | >100 | >100 | >100 | ≤0.25 |
| 26 | MSSA (0701A0095) | >100 | >100 | >100 | >100 | 87 ± 6 | 83 ± 29 | ≤0.25 |
| 29 | MRSA (11004533801) | >100 | >100 | >100 | >100 | 80 ± 0 | 87 ± 21 | ≥4 |
| 30 | MRSA (11004691801) | >100 | >100 | >100 | >100 | 77 ± 6 | 67 ± 23 | ≥4 |
| 31 | MRSA (11004787401) | >100 | >100 | >100 | >100 | 97 ± 6 | >100 | ≥4 |
| 32 | MRSA (11006153901) | >100 | >100 | >100 | >100 | 77 ± 6 | 73 ± 29 | ≥4 |
| 33 | MSSA (11004327701) | >100 | >100 | >100 | >100 | 77 ± 6 | 73 ± 12 | 0.25 |
| 34 | MSSA (11004480701) | >100 | >100 | >100 | >100 | 80 ± 0 | 97 ± 12 | 0.5 |
| 35 | MSSA (11004691801) | >100 | >100 | >100 | >100 | 77 ± 6 | 90 ± 17 | 0.5 |
| 36 | MSSA (11004010401) | >100 | >100 | >100 | >100 | 77 ± 6 | 90 ± 0 | ≤0.25 |

FIGURE 2: Chemical structure of the new compound 40.

determined as C₂₅H₂₂O₇ by HRESIMS (found for [M+H]⁺ 435.1436; calculated 435.1438). The UV spectrum showed an absorption maximum at 350 nm. The IR spectrum indicated the presence of OH (3400 cm⁻¹) as well as conjugated ketone carbonyl (1610 cm⁻¹) groups. The ¹H NMR spectrum exhibited signals due to a hydrogen-bonded OH at $\delta_{\rm H}$ 14.49, two *trans*-olefinic protons ($\delta_{\rm H}$ 8.06 and 7.63, 2 d, J=15.7 Hz), aromatic rings (9H, $\delta_{\rm H}$ 6.15–7.30), and one methoxyle ($\delta_{\rm H}$ 3.92). It also showed the characteristic signals of a flavan-3ol moiety at $\delta_{\rm H}$ 4.68 (1H, d, J = 8.9 Hz, H₂), 4.21 (1H, m, H₃), 3.07 (1H, dd, J = 16.2, 5.7 Hz, H_{4a}), and 2.53 (1H, dd, J = 16.2, 9.5 Hz, H_{4b}). The ¹³C NMR and HMQC spectra confirmed the presence of 25 carbons with typical flavan-3-ol signals at $\delta_{\rm C}$ 84.2 (C₂), 66.8 (C₃), and 30.6 (C₄). In the ¹H NMR spectrum, the signals at $\delta_{\rm H}$ 7.11 (1H, d, $J=1.4\,{\rm Hz}$), 6.96 (1H, dd, J = 8.4, 1.4 Hz), and 6.91 (1H, d, J = 8.4 Hz) suggested the presence of a 1',3',4'-trisubstituted ring B whereas a singlet at $\delta_{\rm H}$ 6.15 (1H) indicated a pentasubstituted ring A. Two multiplets at $\delta_{\rm H}$ 7.17 (2H) and 7.29 (3H) revealed the presence of a phenyl residue. The HMBC spectrum showed a longrange correlation between the two *trans*-olefinic protons [$\delta_{\rm H}$ 7.63 (1H, d, J = 15.7, H_{α}) and 8.06 (1H, d, J = 15.7, H_{β})] and the ketone carbon at δ_{C} 193.2. This correlation revealed the presence of an α,β -unsaturated ketone group. The *trans*-olefinic proton H_{β} at δ_{H} 8.06 was also correlated with the phenyl quaternary carbon at δ_C 136.2 ($C_{1''}$). This correlation implied the presence of a (2E)-4-phenylprop-2-en-1-one moiety. A correlation between the methoxyle protons ($\delta_{\rm H}$ 3.92) and the carbon at $\delta_{\rm C}$ 165.1 (${\rm C}_{\rm 5}$) proved that the OCH₃ was attached to C₅. The NOESY spectrum showed that this methoxyle was spatially close to the proton at $\delta_{\rm H}$ 6.15 (H₆), whereas a long-range COSY indicated a correlation between H₆ and one of the hydroxyl groups at $\delta_{\rm H}$ 14.49 (OH₇). Therefore a (2E)-4-phenylprop-2-en-1-one moiety was located at C_8 (δ_C 105.9). Finally, it appeared that the aromatic ring B was substituted at $C_{3'}$ and $C_{4'}$ by two hydroxyl groups (NMR spectra cf. supporting information 2). ¹H and ¹³C NMR data together with 2D NMR correlations for 40 are summarized in Table 4 and Figure 3.

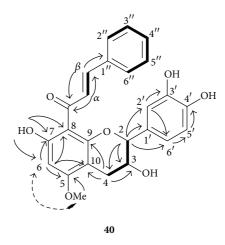


FIGURE 3: 2D NMR studies of compound **40**: COSY (bold lines), selected HMBC (solid arrows: $^{1}H \rightarrow ^{13}C$), and NOESY (dashed arrows) correlations.

Table 4: ¹H and ¹³C NMR data of the new compound **40** (in acetone-*d*6).

| Docition | 40 | | | | | |
|---------------------|--|-------------------------|--|--|--|--|
| Position | δ_{H} , mult. (J in Hz) | $\delta_{ m C}$, mult. | | | | |
| 2 | 4.68, d (8.9) | 84.2, CH | | | | |
| 3 | 4.21, m | 66.8, CH | | | | |
| 4 | a 3.07, dd (16.2, 5.7) | 30.6, CH ₂ | | | | |
| | b 2.53, dd (16.2, 9.5) | | | | | |
| 5 | | 165.1, qC | | | | |
| 6 | 6.15, s | 93.5, CH | | | | |
| 7 | | 168.1, qC | | | | |
| 8 | | 105.9, qC | | | | |
| 9 | | 157.6, qC | | | | |
| 10 | | 102.4, qC | | | | |
| 1' | | 130.7, qC | | | | |
| 2' | 7.11, d (1.4) | 116.2, CH | | | | |
| 3' | | 146.6, qC | | | | |
| 4' | | 146.2, qC | | | | |
| 5' | 6.91, d (8.4) | 116.0, CH | | | | |
| 6' | 6.96, dd (8.4, 1.4) | 121.3, CH | | | | |
| 1" | | 136.2, qC | | | | |
| 2" | 7.28, m | 129.2, CH | | | | |
| 3" | 7.16, m | 129.7, CH | | | | |
| 4" | 7.28, m | 130.8, CH | | | | |
| 5" | 7.16, m | 129.7, CH | | | | |
| 6" | 7.28, m | 129.2, CH | | | | |
| OH-7 | 14.49, s | | | | | |
| α | 7.63, d (15.7) | 143.1, CH | | | | |
| β | 8,06, d (15.7) | 128.4, CH | | | | |
| C=O | | 193.2, qC | | | | |
| OCH ₃ -5 | 3.92, s | 56.5, CH ₃ | | | | |

40 had no optical rotation and, thus, was isolated here as a racemate mixture of 8-[(E)-phenylprop-2-en-1-one]-(2R,3S)-5-methoxycatechin (**40a**) and 8-[(E)-phenylprop-2-en-1-one]-(2S,3R)-5-methoxycatechin (**40b**). Sha et al.

>100

FIGURE 4: Structures of 8-[(E)-phenylprop-2-en-1-one]-(2R,3S)-5-methoxycatechin (40a) and 8-[(E)-phenylprop-2-en-1-one]-(2S,3R)-5-methoxycatechin (40b) and a similar pair of racemates isolated by Sha et al. 2009 [26].

Antifungal activity Antibacterial activity Fungal strains $MIC_{80} (\mu g/mL)$ Number Bacterial strains $MIC_{100} (\mu g/mL)$ C. albicans 22 S. aureus (ATCC25923) >250 >100 C. glabrata 23 MRSA (0706C0025) >250 ≤10 A. fumigatus 24 MRSA (0702E0196) >250 ≤10 25 MSSA (0703H0036) >100

26

TABLE 5: Antifungal and antibacterial activities of 40.

already isolated a similar compound, only differing from 40 by a 1',3',5'-trisubstituted aromatic ring B, from a Chinese propolis [26] (Figure 4).

3.3. Antifungal and Antibacterial Activities of **40**. The new flavan-3-ol **40** did not show any antifungal activity on the three strains studied (Table 5). However, though active neither on *S. aureus* nor on MSSA, its MIC_{100} on MRSA numbers 23 and 24 were lower or equal to $10 \,\mu g/mL$ (close to oxacillin: $\geq 4 \,\mu g/mL$).

3.4. Major Compounds Activities. Antifungal and antibacterial activities were then individually evaluated for the five major compounds identified in E2–6, namely, pinobanksin-3-acetate (28), pinocembrin (25), chrysin (32), galangin (34), and prenyl caffeate (29) [24]. Their MIC₈₀ towards *C. albicans, C. glabrata*, and *A. fumigatus* as well as their MIC₁₀₀ towards *S. aureus*, MRSA, and MSSA are given in Table 6.

Pinobanksin-3-acetate (28), chrysin (32), and galangin (34) appeared as inactive. Pinocembrin (25) showed a moderate activity towards *Candida albicans*, *C. glabrata* (MIC $_{80}$ 62–125 μ g/mL), and *S. aureus* (MIC $_{100}$ 100 μ g/mL). Overall prenyl caffeate (29) exhibited the best activities (MIC $_{100}$ up to 16 μ g/mL against *C. glabrata* and up to 63 μ g/mL against *S. aureus* and MRSA). Even so it appeared that these compounds were not individually as active as it could be expected from E5-6 results (MIC $_{100}$ 30–97 μ g/mL). As far as *S. aureus* and MRSA are concerned, this kind of synergistic effects was recently pointed out by Darwish et al. [20] who evaluated the antibacterial activities of pinobanksin-3-acetate, pinocembrin, and chrysin isolated from a Jordanian propolis. Therefore these results are also in agreement with Kujumgiev

et al. stating that, in spite of a great chemodiversity, no specific compounds can be associated with the antimicrobial activities of propolis extracts whereas, obviously, different flavonoid combinations are essential for these activities [7]. The antimicrobial of propolis extracts most probably involves a complex mechanism. It can be attributed to the synergistic effects of phenolic compounds such as cinnamic acid and ester derivatives, including caffeic acid and CAPE, as well as flavonoids including quercetin and naringenin [17, 53, 54]. Indeed, each of these compounds would be able to increase membrane permeability and inhibit bacterial mobility [54], thus contributing to the antimicrobial activity of propolis but also to its synergism with other antibiotics [53, 55, 56]. It is the reason why Stepanović et al. could notice the antibacterial and synergistic actions of propolis extracts with ampicillin, ceftriaxone, and doxycycline towards Staphylococcus aureus and with nystatin towards Candida albicans, stating that the bacterial resistance to antibiotics had no influence on the susceptibility to propolis extracts [57]. In vitro studies of synergism carried by Fernandes Jr. et al. also revealed synergistic effects of EEP with chloramphenicol, gentamicin, netilmicin, tetracycline, vancomycin, and clindamycin [58]. Therefore our findings are in total accordance with these results and, now that antibiotic resistance to bacteria has become a major public health concern [59], could bring valuable knowledge to develop new antimicrobial drugs for challenging S. aureus infections.

MSSA (0701A0095)

4. Conclusions

On the basis of these results, it may be concluded that organic solvents extracts of a French poplar type propolis are

Table 6: Antifungal and antibacterial activities of five major compounds.

| | | | _ | | | |
|-----------------------|-----------------------------------|----------------------------|--------------|---------------|----------------------|----------------|
| Antifungal strains | Pinobanksin-3- acetate (28) | Pinocembrin (25) | Chrysin (32) | Galangin (34) | Prenyl caffeate (29) | Amphotericin B |
| Candida albicans | 250 | 62 | >250 | >250 | 62 | 0.125 |
| Candida glabrata | 250 | 125 | >250 | >250 | 16 | 0.125 |
| Aspergillus fumigatus | >250 | 250 | >250 | >250 | 125 | 6 |
| | | MIC ₁₀₀ (μg/mL) | | | | |

Pinobanksin-3-Prenyl caffeate Number Bacterial strains Pinocembrin Galangin (34) Oxacillin acetate Chrysin (32) (25)(29)(28)Staphylococcus aureus 22 >100 100 ± 0 >100 >100 63 ± 6 ≤0.25 (ATCC25923) 23 MRSA (0706C0025) >100 >100 >100 70 ± 0 >100 ≥ 4 >100 24 MRSA (0702E0196) >100 >100 >100 70 ± 0 \geq 4 25 MSSA (0703H0036) >100 >100 >100 >100 93 ± 6 ≤0.25 MSSA (0701A0095) >100 >100 >100 >100 ≤0.25 26 93 ± 6

associated with a good antifungal activity towards *Candida albicans* and *C. glabrata*, correlated with high flavonoid contents. However only DCM based extracts (E5-6) showed a significant antibacterial activity against both methicillinresistant and methicillin-susceptible *Staphylococcus aureus* strains. Unfortunately these extracts are not compatible with a pharmaceutical use because of their toxicity, whereas EtOH based extracts were not as active as expected. Therefore it would be interesting to develop some alternative extraction of propolis using a nontoxic solvent such as subcritical water. In addition, it should be noticed that, as an intrinsic polytherapy, propolis may also circumvent the development of drug resistance by bacteria [60].

Abbreviations

APCI: Atmospheric pressure chemical ionization

ATCC: American Type Culture Collection

AYE: Patient code

CAPE: Caffeic acid phenylethyl ester

CBS: Centraal Bureau voor Schimmelcultures

CIP: Collection de l'Institut Pasteur
COSY: Correlation spectroscopy

DCM: Dichloromethane DMSO: Dimethyl sulfoxide

EEP: Ethanolic extract of propolis ESI: Electrospray ionization FD: Flavanone/dihydroflavonol

FF: Flavone/flavonol GAE: Gallic acid equivalent

HMBC: Heteronuclear multiple bond correlation

HMQC: Heteronuclear multiple quantum

correlation

HPLC: High performance liquid chromatography HRESIMS: High resolution electrospray ionization

mass spectrometry

IR: Infrared

LMA: Laboratoire de Mycologie d'Angers MIC: Minimum inhibitory concentration MRSA: Methicillin-resistant *Staphylococcus aureus*

MS: Mass spectrometry

MSSA: Methicillin-susceptible *Staphylococcus aureus* NCCLS: National Committee for Clinical Laboratory

Standards

NMR: Nuclear magnetic resonance

NOESY: Nuclear Overhauser effect spectroscopy

RCH: Patient code RP: Reversed phase

RPMI: Roswell Park Memorial Institute

UV: Ultraviolet

YPDA: Yeast peptone dextrose agar.

Conflict of Interests

This study was financed by Ballot-Flurin Apiculteurs Cie. There are no other competing interests.

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