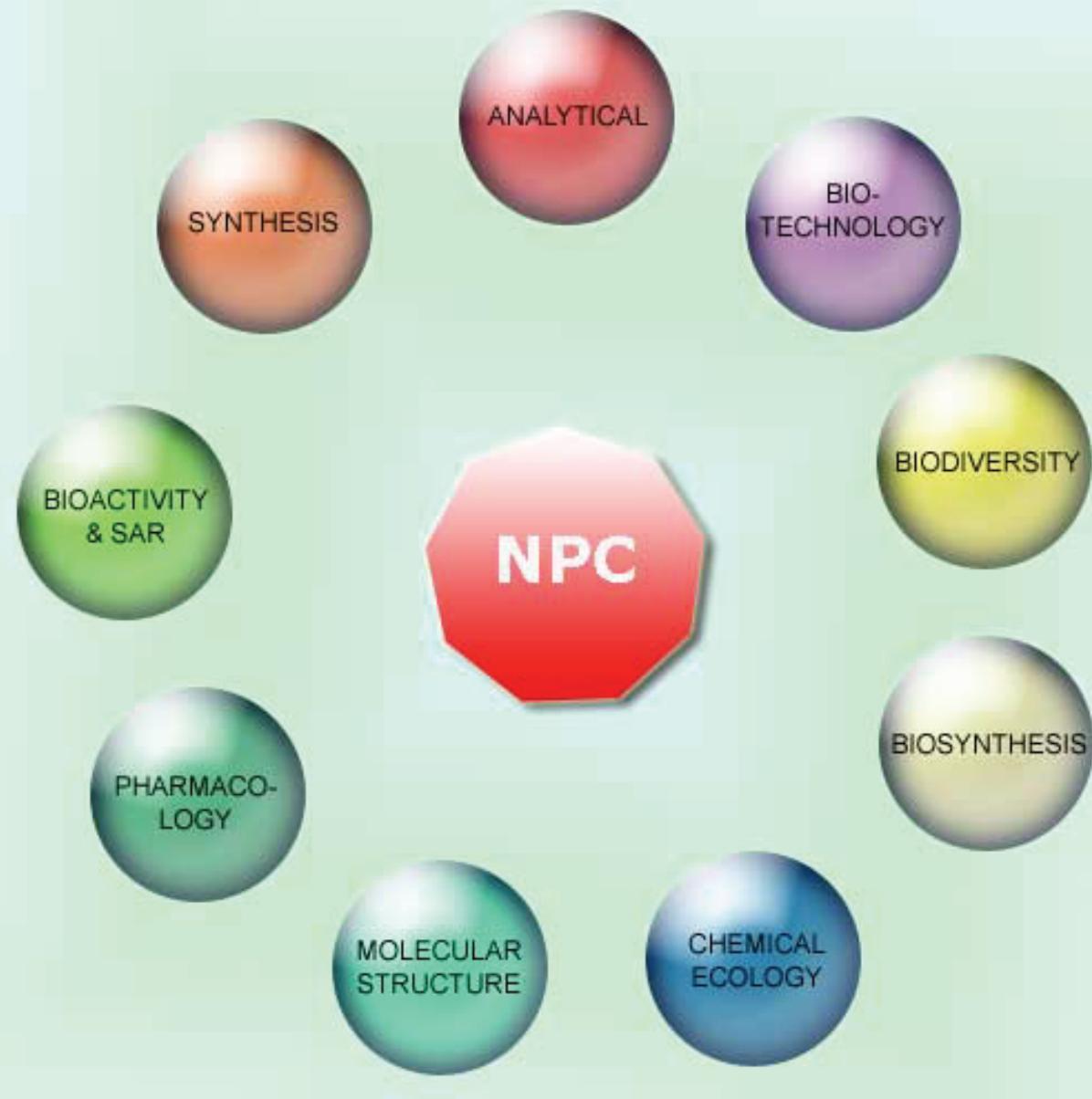


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Editorial Message

The broad scientific scope of *Natural Product Communications (NPC)* means that we publish high quality papers across a wide spectrum of natural products research, including organic, medicinal, pharmaceutical, bioorganic, analytical, theoretical, biotechnological and chemotaxonomic aspects. We welcome all types of manuscripts: communications, full papers and reviews. We have a very diverse international authorship, and a readership that is continuously growing, as indicated by the increasing number of submissions and published papers, and also from the web statistics. *NPC* is a SCI-cited journal with an impact factor in 2015 of 0.884. We would like to **thank our authors, readers and referees who are the architects of this success.**

During March 2016, we published a themed issue dedicated to “Traditional Medicines as Sources of Bioactive Compounds (Guest Editors: Profs. Giovanni Vidari and Paola Vita Finzi)”. We had four honorarium issues to recognize internationally renowned natural product researchers, Prof. Dr Motto Tori, for his 65th Birthday (February 2016), Prof. Dr. DHC Yoshinori Asakawa, President of the Phytochemical Society of Asia, for his 75th Birthday (July and August 2016) and Prof. Dr Wilhelm Fleischhacker, for his 85th Birthday (October 2016). We are committed to placing this journal at the forefront for the dissemination of new and exciting natural products research. *NPC* will be proud to recognize eminent natural product scientists by dedicating an issue in their honour. Suggestions for themed and/or honorarium issue(s) would be welcomed. We look forward to receiving your submissions and your feedback.

NPC in collaboration with the Phytochemical Society of Asia (PSA) organized an International Symposium on Natural Products for the Future 2016 (ISNPF2016) that was held between September 1-4, 2016 at Tokushima Bunri University, Tokushima, Japan. This attracted scientists, professionals and representatives of companies working in the exciting field of natural product research not only from Asia but also from all over the world. We have also planned a special NPC-ISNPF issue to be published during 2017.

Professor Stefano Serra, National Research Council (CNR), Istituto di Chimica del Riconoscimento Molecolare (ICRM), Milano, Italy, won the 2015 “Gerald Blunden Award” for the best article “The Co-identity of Lipiarmycin A3 and Tiacumicin B” *Natural Product Communications* 9 (2) 237-240 (2014), and co-authored by Angelo Bedeschi, Piera Fonte, Giovanni Fronza, and Claudio Fuganti. The Award wins a free year's subscription to *Natural Product Communications* and is accompanied by US\$ 500, sponsored by Natural Product Inc. The award was made for the best article published during the last two years (2014-2015). In 2017, the award will be made for the best review article published during 2015-2016. Each year, the winner will be announced in May at the time of Prof. Blunden's Anniversary, and a similar pattern will be followed in consecutive years.

For the information of our authors and readers, we have to announce some changes to the Editorial Board and Editorial Board of Advisors. Professor De-an Guo, Shanghai Institute of Materia Medica, Shanghai 201203, P. R. China, has been an editor from the time of inception of the journal. He has expressed a wish to step down from his editorial role. His high level of commitment and expertise will be missed, and the editors and publisher thank him for his many valuable contributions. Leaving the Editorial Advisory Board are Professors Viqar Uddin Ahmad, Josep Coll, Imre Mathe and Luca Rastrelli; we thank them for their support and their service. We are, however, pleased to report that as of January 2017, Professor Ping-Jyun Sung, National Museum of Marine Biology and Aquarium Checheng, Pingtung 944, Taiwan, will be joining the Editorial Board, bringing considerable expertise and experience, particularly in marine natural products. Professors Norbert Arnold, Vassya Bankova and Lindros A. Skaltsounis will be joining the Editorial Advisory Board. We welcome them all and look forward to working with them.

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Cytotoxic and Antileishmanial Components from the Bark Extract of *Ruyschia phylladenia* from Monteverde, Costa Rica

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The bark of *Ruyschia phylladenia* was collected from Monteverde, Costa Rica, and extracted with acetone. Bioactivity-directed chromatographic separation of the crude acetone bark extract of *R. phylladenia* led to isolation and identification of lupeol, betulinic acid, and isofraxidin. Lupeol and betulinic acid showed *in-vitro* cytotoxic activity to MCF-7, MDA-MB-231, and 5637 human tumor cell lines. Isofraxidin was not cytotoxic, but did show antileishmanial activity to *Leishmania amazonensis* promastigotes.

Keywords: *Ruyschia phylladenia*, Cytotoxic, *Leishmania amazonensis*, Lupeol, Betulinic acid, Isofraxidin.

Ruyschia phylladenia Sandwith is a Neotropical liana belonging to the Marcgraviaceae that is found in Costa Rica and Panama at an altitude of about 1200 m [1]. Twenty-eight different species of *Ruyschia* are currently recognized [2]. The crude bark extracts of *R. phylladenia* had previously shown antibacterial activity against Gram-positive *Bacillus cereus* and *Staphylococcus aureus* and cytotoxic activity against Hep-G2 (human hepatocellular carcinoma), MDA-MB-231 (human breast adenocarcinoma), and Hs 578T (human breast ductal carcinoma) cell lines [3], as well as antiparasitic activity against *Leishmania amazonensis* promastigotes and amastigotes ($IC_{50} < 12.5$ and = 22.9 μ g/mL, respectively) [4]. In this work, we present the bioactivity-directed separation and identification of bioactive components from *R. phylladenia*. To our knowledge, no previous phytochemical studies have been carried out on this plant.

The bark of *R. phylladenia* was collected from the Monteverde region of Costa Rica and extracted with acetone. The crude acetone bark extract was subjected to bioactivity-directed preparative chromatography to give lupeol, betulinic acid, and isofraxidin as the biologically active components (Table 1).

Both lupeol and betulinic acid have shown cytotoxic activity against several tumor cell lines [5], so the cytotoxicity exhibited by these lupane triterpenoids in this study was not surprising. Although these compounds also showed antileishmanial activity against *L. amazonensis* promastigotes, the cytotoxicity against BALB/c mouse macrophages indicates a lack of selectivity; these compounds cannot be considered as potential antileishmanial agents. Lupeol [5b] and betulinic acid [6] have also shown antitrypanosomal and antiplasmodial activity.

The coumarin isofraxidin, on the other hand, showed no cytotoxic activity against the tumor cell lines and minimal toxicity to BALB/c mouse macrophages, while also showing excellent antileishmanial activity. The antiparasitic activity of isofraxidin is consistent with a previous report of antiplasmodial activity of this compound [7].

Table 1: *In-vitro* cytotoxic activities of compounds from *Ruyschia phylladenia* acetone bark extract.

Compounds	Cytotoxicity (IC_{50} , μ g/mL)				<i>L. amazonensis</i> promastigotes
	MCF-7	MDA-MB-231	5637	BALB/c mouse macrophage	
Lupeol	76.6±2.8	66.1±1.2	32.9±6.8	19.3±3.8	8.5±0.1
Betulinic Acid	15.5±3.9	3.16±1.38	1.29±0.12	10.5±2.9	11.3±4.9
Isofraxidin	inactive	inactive	inactive	51.9±6.2	11.1±1.3
Control	28.6±1.9 ^a	25.9±2.4 ^a	0.98±0.23 ^a	11.7±1.7 ^b	0.37±0.01 ^b

^aDoxorubicin. ^bPentamidine.

This report presents the isolation and identification of lupeol, betulinic acid, and isofraxidin from a new, here-to-fore uninvestigated plant source and reaffirms the biological activities of these compounds.

Experimental

Plant Material: *Ruyschia phylladenia* was collected from the Monteverde region of Costa Rica ($10^{\circ} 17' 25.24''$ N, $84^{\circ} 49' 0.82''$ W, 1173 m elevation) on 14 May 2008. The plant was identified by W.A. Haber and a voucher specimen (Haber 2921) has been deposited in the herbarium of the Missouri Botanical Garden. The bark was chopped and air dried and the dried bark (462 g) was extracted with refluxing acetone for 5 hours using a Soxhlet extractor. Evaporation of the acetone gave 60.0 g crude bark extract.

Chromatographic Separation: The crude acetone bark extract of *R. phylladenia* (25.0 g) was separated by flash column chromatography (silica gel: 90 cm L × 5 cm D) using a step gradient with *n*-hexane/ethyl acetate, 200-mL fractions: 9:1 *n*-hexane/ethyl acetate, F1-F4; 8:2 *n*-hexane/ethyl acetate, F5-F25; 7:3 *n*-hexane/ethyl acetate, F26-F44; 1:1 *n*-hexane/ethyl acetate, F45-F56; 3:7 *n*-hexane/ethyl acetate, F57-F75; 1:9 *n*-hexane/ethyl acetate, F76-F82; 100% ethyl acetate, F83-F90. Fractions F13-F14 were combined and recrystallized from ethyl acetate/pentane to give 38.8 mg lupeol, which had ¹H and ¹³C NMR and IR spectra in agreement with those previously reported in the literature [8]. Fractions F24-F42 were combined and recrystallized from ethyl acetate/pentane to give 1.70 g betulinic acid, which had ¹H and ¹³C NMR and IR spectra in agreement with those previously reported in

the literature [8ab,9]. Fractions F57-F61 were combined and recrystallized from ethyl acetate/pentane to give 13.8 mg isofraxidin, which had ¹H and ¹³C NMR in agreement with those previously reported in the literature [10].

Cytotoxicity Assay: The isolated compounds from *R. phylladenia* were screened for *in-vitro* cytotoxicity against MCF-7 (human breast adenocarcinoma), MDA-MB-231, and 5637 (human bladder carcinoma) cell lines using the MTT assay as described previously [11].

Antileishmanial Assay: *L. amazonensis* culturing, antileishmanial screening against *L. amazonensis* promastigotes, and cytotoxicity

screening against BALB/c mouse macrophages were carried out as previously described [4].

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A New Cardenolide and Other Compounds from *Salsola tetragona*

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One new cardenolide, 3-*O*-β-D-allopyranosylcoroglaucigenin (1), was isolated from the aerial parts of *Salsola tetragona* Delile with four known cardenolides (2-5), two known flavonoids (6-7), three known phenolic compounds (8-10) and two known fatty acids (11-12). Their structures were identified by spectroscopic analyses and by comparison of their spectral data with those reported in the literature. Compounds 1-5, 7, 10 and 12 were isolated from the genus *Salsola* for the first time. This is the first report on cardenolides identified in the Amaranthaceae family.

Keywords: *Salsola tetragona* Delile, Amaranthaceae, Cardenolides, Phenolic glycosides.

The genus *Salsola* (Amaranthaceae, ex. Chenopodiaceae) consists of over 174 species found in the arid regions of Asia, Europe and Africa [1]. *Salsola* species possess antihypertensive, anti-inflammatory, anticancer, antioxidant, anti-Alzheimer, antidepressant and antimicrobial activities [2-5]. Previous phytochemical investigations of this genus reported the isolation of flavonoids and other phenolic compounds [6], alkaloids [7], fatty acids [8] and triterpene glycosides [9]. In continuation of our works on plants from Algerian Septentrional Sahara [10-11], we report here the isolation and structural characterization of one new cardenolide, 3-*O*-β-D-allopyranosylcoroglaucigenin, named salsotetragonin (1), in addition to eleven known compounds from aerial parts of *S. tetragona* (Figure S1).

The hydromethanolic extract of the aerial parts of *S. tetragona* was partitioned successively with light petroleum, CH₂Cl₂, EtOAc and *n*-butanol to give four extracts. Purification of the *n*-butanol extract through repeated silica gel columns, flash chromatography and semi-prep HPLC led to the isolation and identification of a new cardenolide (1) and seven known compounds (3, 4, 6-10). The dichloromethane extract was fractionated by silica gel chromatography to give four known compounds (2, 5, 11-12). These compounds were identified by extensive spectroscopic methods including 1D-¹H and ¹³C) and 2D-NMR (COSY, HSQC, HMBC and NOESY) experiments, as well as HR-ESI-MS analysis and by comparison of their spectral data with the literature. The eleven known compounds were identified as four cardenolides: uzarigenin (2) [12], desglucouzarin (3) [13], 12-dehydroxy-ghalakinoside (4), and calactin (5) [14] (Figure 1), two flavonoids: kaempferol-3-*O*-β-D-glucopyranoside (6) [15] and quercetin-3-*O*-β-D-glucopyranosyl-(1→6)-glucopyranoside (7) [16], three phenolic compounds: vanillic acid (8) [17], canthoside C (9), and canthoside D (10) [18] and two fatty acids : oleic acid (11) [19], and 2,3-dihydroxypropylpalmitate (12) [20] (Figure S1).

Compound 1 was isolated as a white amorphous powder. Its positive HR-ESI-MS showed a pseudomolecular ion peak at *m/z* 575.2829 [M+Na]⁺, corresponding to the molecular formula C₂₉H₄₄O₁₀. The

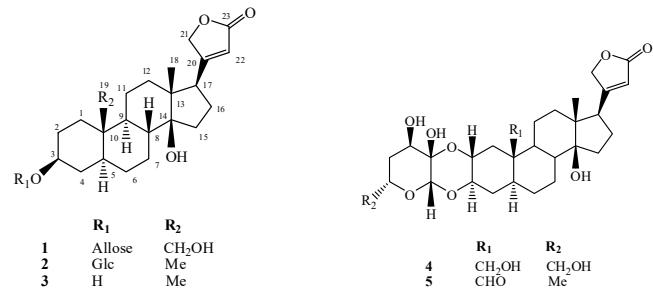


Figure 1: Structures of compounds 1-5.

¹H NMR spectrum of compound 1 (Table 1) showed characteristic signals of a butenolactone ring at δ_H 4.95 (1H, dd, *J* = 18.4, 1.6 Hz, H-21a), 5.07 (1H, dd, *J* = 18.4, 1.4 Hz, H-21b) and 5.93 (1H, s, H-22), one oxymethylene at δ_H 3.74 (1H, d, *J* = 11.7 Hz, H-19a) and δ_H 3.88 (1H, d, *J* = 11.7 Hz, H-19b), one oxymethine at δ_H 3.78 (m, H-3), and one methyl signal at δ_H 0.95 (3H, s, H-18). Its ¹³C NMR spectrum (Table 1) showed twenty-nine carbon signals including six of the sugar part (Figure 1).

The ¹H NMR and ¹³C NMR data of 1 were very similar to those of 3-β-D-glucopyranosylcoroglaucigenin [21]. The COSY and HSQC spectra showed one anomeric proton of a sugar moiety at δ_H 4.78 (d, *J* = 7.9 Hz, H-1') and δ_C 99.8 with five signals of an hexose unit [H-2' [δ 3.28 (dd, *J* = 7.9, 3.0 Hz)], H-3' [δ 4.06 (t, *J* = 3.0 Hz)], H-4' [δ 3.48 (dd, *J* = 9.5, 3.0 Hz)], H-5' [δ 3.69 (m)], and H₂-6', [δ 3.66 (dd, *J* = 11.3, 5.7 Hz) and 3.85 (dd, *J* = 11.3, 1.8 Hz)]; the sugar was identified as an allopyranoside [22]. The large coupling constant (*J* = 7.9 Hz) of the anomeric proton at δ_H 4.78 indicated that the allose had a β-configuration. The linkage of the allose at C-3 of the aglycone was established from the HMBC correlation between allo-H-1' (δ_H 4.78) and C-3 (δ_C 78.9) (Figure 2). The stereochemistry of 1 was confirmed by the NOE effect (Figure 2) between H-5/H-9 and H-3, which clearly established the α-orientation of H-5. The correlation of H₃-18 to H-21 and H-22 in

the NOESY spectrum indicated the β orientation of the γ lactone at C-17. The chemical shifts of the carbons of the C and D rings of **1** are in agreement with those of coroglaucigenin and madagascarrensisid (A) recorded in deuterated pyridine and in CD₃OD [23,24], indicating a 14- β hydroxy orientation. Thus compound **1** was identified as 3-O- β -D-allopyranosyl-coroglaucigenin, which we named salsotetragonin.

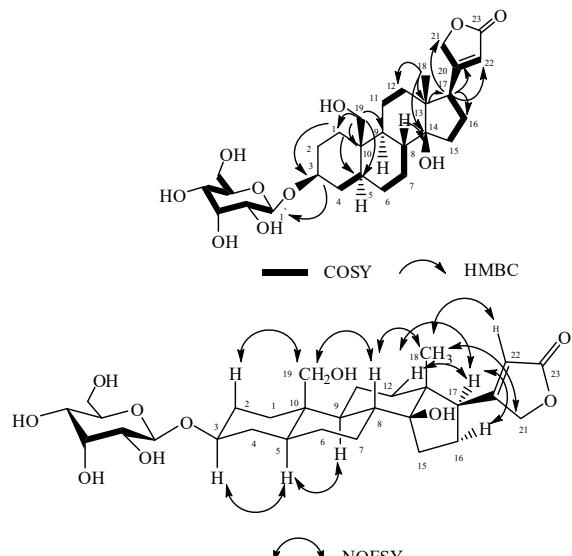


Figure 2: ^1H - ^1H COSY, HMBC and NOESY correlations of **1**.

Table 1: ^1H and ^{13}C NMR data for compound **1** (in CD₃OD; ^{13}C : 150 MHz; ^1H : 600 MHz).

Position	1	δ_{H} (m, J in Hz)	δ_{C}
1	0.81 (td, 13.6, 3.5), 2.34 (dt, 13.3, 3.5)	32.8	
2	1.57 (dm, 14.9), 1.91 (m)	30.9	
3	3.78 (m)	78.9	
4	1.44 (m), 1.82 (dd, 11.9, 3.4)	35.9	
5	1.24 (m)	45.9	
6	1.24 (m), 1.34 (m)	29.6	
7	2.10 (dd, 12.3, 3.1), 1.15 (m)	28.8	
8	1.82 (td, 11.9, 3.4)	43.1	
9	1.03 (td, 13.5, 4.5)	51.5	
10	-	40.6	
11	1.65 (m)	24.1	
12	1.41 (td, 13.6, 4.3), 1.53 (dt, 13.5, 3.0)	41.5	
13	-	51.2	
14	-	86.3	
15	1.73 (dd, 11.6, 8.5), 2.14 (m)	33.5	
16	2.16 (m), 1.90 (m)	28.1	
17	2.85 (dd, 9.5, 5.8)	52.2	
18	0.95 (s)	16.7	
19	3.74 (d, 11.7), 3.88 (d, 11.7)	59.9	
20	-	178.5	
21	4.95 (dd, 18.4, 1.6), 5.07 (dd, 18.4, 1.4)	75.3	
22	5.93 (s)	117.8	
23	-	176.9	
Allose (at C-3)			
1'	4.78 (d, 7.9)	99.8	
2'	3.28 (dd, 7.9, 3.0)	72.4	
3'	4.06 (t, 3.0)	73.1	
4'	3.48 (dd, 9.5, 3.0)	69.1	
5'	3.69 (m)	75.5	
6'	3.66 (dd, 11.3, 5.7), 3.85 (dd, 11.3, 1.8)	63.3	

Experimental

General: Optical rotations were measured in DMSO with a Perkin-Elmer 241 polarimeter. ^1H and ^{13}C NMR spectra were recorded on a Bruker Avance DRX III 500 spectrometer (^1H at 500 MHz and ^{13}C at 125 MHz) and on a Bruker Avance III-600 spectrometer equipped with a cryo platform (^1H at 600 MHz and ^{13}C at 150 MHz). 2D-NMR experiments were performed using standard Bruker-microprograms (TopSpin version 3.2 software). HR-ESI-

MS and ESI-MS experiments were performed using a Micromass Q-TOF micro-instrument (Manchester, UK). TLC was carried out on pre-coated silica gel 60 F₂₅₄ (Merck). Vacuum liquid chromatography (VLC) was carried out on LiChroprep RP-18 (40–63 μm , Merck). HPFC was performed on a Grace® Reveleris System using Grace® cartridges (Silica gel or RP-C₁₈). The semi-preparative HPLC was performed on a Dionex apparatus equipped with an ASI-100 automated sample injector, a STH 585 column oven, a LPG-3400A pump, a diode array detector UVD-340U and Chromeleon® software.

Plant material: *Salsola tetragona* Delile was collected in Ghardaïa (Algerian Septentrional Sahara) in May 2011 and was identified by Pr. Gérard De Belair (University of Annaba, Algeria). A voucher specimen (LOST.St05/09) was deposited at the herbarium of the Faculty of Sciences, University frères Mentouri- Constantine.

Extraction and isolation: Air-dried and powdered aerial parts (1980 g) of *S. tetragona* were extracted with 80% MeOH (10 L). After evaporating the MeOH under *vacuum*, the residue was dissolved in water (900 mL) and partitioned with PE, CH₂Cl₂, EtOAc, and *n*-BuOH, successively (3 x 300 mL). The *n*-BuOH extract (6 g) was subjected to VLC over RP-18 eluted with H₂O-MeOH (6:4, 5:5, 4:6, 3:7, 2:8 and 0:10). Fractions of 150 mL were collected and pooled according to their similarity in TLC profile to give 6 fractions (F1–F6 respectively). F1 (1.2 g) was submitted to RP-18 flash chromatography eluting with MeCN:H₂O (5:95 to 10:0) to afford 19 sub-fractions. Sub-fraction [20-23] (35 mg), was purified by semi-prep HPLC on RP-18 eluted with MeCN:H₂O (6:94 to 1:9), affording compounds **9** (3 mg) and **10** (1.5 mg). F2 (1.5 g) was chromatographed on a silica gel column eluted with EtOAc:MeOH:H₂O (20:2:1), giving 2 main sub-fractions. Sub-fraction [90] (180 mg) afforded **4** (22 mg) by silica gel CC (CHCl₃:MeOH, 9:1). Sub-fraction [115-119] (223 mg) was subjected to silica gel CC eluting with CHCl₃:MeOH (8:2) to afford **1** (27 mg) and sub-fraction [35-48] (100 mg), which was purified over silica gel using EtOAc:MeOH:H₂O (20:2:1) to yield compounds **6** (15 mg) and **7** (18 mg). F3 (750 mg) was fractionated by silica gel CC eluted with CHCl₃:MeOH (0 to 100% of MeOH). The main sub-fraction [59-63] (212 mg) was further separated by silica gel CC (CHCl₃:MeOH, 0 to 100% of MeOH) and **3** (25 mg) was isolated. F4 (302 mg) was subjected to silica gel CC using CHCl₃:MeOH (0 to 100% of MeOH) to give 3 sub-fractions. Sub-fraction [72] (111.2 mg) was selected for silica gel CC eluted with CHCl₃:MeOH (0 to 100% of MeOH) to yield compound **8** (3 mg).

The CH₂Cl₂ extract (2.5 g) was subjected to silica gel CC, eluted with toluene:CHCl₃ (0 to 100% of CHCl₃) and CHCl₃:EtOAc (0 to 100% of EtOAc). Sub-fraction [27-32] (123 mg) was purified over silica gel using diethyl ether:EtOAc (9:1) to yield compound **11** (1.4 mg). Sub-fraction [160-173] (212 mg) was passed through a silica gel column (CHCl₃:EtOAc, 9:1) yielding compounds **2** (15 mg) and **5** (5 mg). Sub-fraction [234-257] (162 mg) was chromatographed on silica gel, eluted with (CHCl₃:MeOH, 9:1) that afforded compound **12** (2 mg).

Acid hydrolysis: The pure compounds were treated with 2 M HCl at 100°C for 1 h. The hydrolysates were extracted with EtOAc and the sugars identified in the aqueous residue by comparison with authentic samples by TLC using silica gel impregnated with 0.2 M NaH₂PO₄ and a solvent system of Me₂CO-H₂O (9:1); the compounds were revealed with aniline malonate. The optical rotation of the purified sugar was measured and compared with an authentic sample of D-allose.

3-O- β -D-Allopyranosylcoroglauconin (salsotetragonin) (1)

White amorphous powder.

[α]_D²⁰: -16.9 (c 0.18, DMSO).¹H NMR (CD₃OD, 600 MHz): Table 1.¹³C NMR (CD₃OD, 150 MHz): Table 1.HR-ESI-MS: [M+Na]⁺ *m/z* 575.2829 (calcd for C₂₉H₄₄O₁₀Na, 575.2832).**References**

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Supplementary data: Figure S1 and Supplementary Data are included in the Supporting information.

Quantification of Saponins in *Asparagus racemosus* by HPLC-Q-TOF-MS/MS

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Asparagus racemosus Willd. or Shatavari (Asparagaceae family) is an important medicinal plant in Ayurvedic medicine as a rejuvenate for women. A method for quantitative analysis of saponin glycosides bioactive constituents in *A. racemosus* is reported. A high performance liquid chromatography quadrupole time of flight mass spectrometry (HPLC-Q-TOF-MS/MS) method was developed and validated for simultaneous determination of five saponin glycosides, asparacoside, shatavarin IX, shatavarin IV, asparanin A and shatavarin V in *A. racemosus* extracted with 70% MeOH. The method was validated through intra- and inter-day precision, with the relative standard deviation (RSD) less than 6%, limits of detection (LOD) and limits of quantification (LOQ) <10 and 50 ng, respectively. Overall recoveries ranged from 95% to 105%, with RSD ranging from 0.7% to 4.5%. The method was applied to saponin glycoside contents in the leaves, stems, and roots of *A. racemosus* sourced from different geographical locations, including four provinces in Thailand, and a sample from India. Saponin glycosides were detected predominantly in the roots, the part used in traditional medicines and these showed wide variations in saponin glycoside profiles from undetectable to 12 mg/g dry weight. The quality control of *A. racemosus* is crucial for reliable and predictable therapies and only methods like the one developed has the necessary flexibility, sensitivity, accuracy, and selectivity for reliable routine quality control.

Keywords: Steroidal saponin, *Asparagus racemosus*, LC-MS/MS, HPLC-Q-TOF-MS/MS, Quantitative analysis.

Asparagus racemosus Willd is among 300 species classified in the family Asparagaceae. The plant is called Shatavari which means “one who possesses a hundred husbands or acceptable to many”, implying its ability to increase fertility and vitality [1]. The plant is commonly found in tropical and subtropical forests in India, Thailand and elsewhere. Its medicinal uses are recorded in Indian and British Pharmacopoeias, and in indigenous systems including Ayurveda, and Thai traditional medicine. The powdered dried root of *A. racemosus* is used in Ayurveda to prevent and for the treatment of gastric ulcers [2] and galactogogue [3] and in Thailand as a galactogogue, to induce menstruation, and as a cosmetic tonic. Contemporary studies reveal estrogenic [4], galactagogic [3] effects in animals, anti-fungal [5], anti-diarrheal [6], anti-oxidant [7], and anti-cancer [8] actions *in vitro*.

Steroidal saponins found throughout the plant is characteristic of the *Asparagus* genus and used for chemotaxonomic classification. Many saponin glycosides have been isolated from *A. racemosus* roots and fruits including shavarins (I, V-X) [9], asparanins [10], asparosides [10], curillins [10], curilosides [10], immunoside [11], racemoside A [12] and shatavaroside C [12]. Among these, Shatavarin I and IV were reported as the major components [9b] in *A. racemosus* collected in India. These saponin glycosides are responsible for *A. racemosus* medicinal affects including, anticancer activity. Immunoside and asparanin A induced apoptosis of cultured tumor cells [8] while shatavarin IV was cytotoxic to several cancer cell lines [13]. Racemoside A, a water-soluble saponin glycoside was a selective and potent inducer of apoptosis in drug resistant

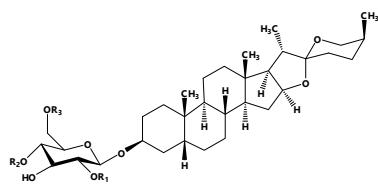
donovani promastigotes [14]. We also demonstrated *in vitro* cytotoxicity of asparacoside, shatavarin IX, shatavarin IV, asparanin A and shatavarin V against human hepato- and prostate-carcinoma cell lines [15].

Clearly, these compounds have considerable clinical potential but there lacks a sensitive and reliable way of quantitating these saponin glycosides in crude *A. racemosus* preparations from different sources. The high performance thin layer chromatography (HPTLC) [16] and high performance liquid chromatography (RP-HPLC) [17] with UV detection showed low sensitivity as the saponin glycosides lack chromophores. Recently, we have developed competitive enzyme-linked and immunosorbent assay (ELISA) using monoclonal antibody against shatavarin IV [18] but the method showed cross-reactivity between glycosides sharing the same aglycone. To address these problems, a HPLC-Q-TOF-MS/MS method was developed such that compounds could be identified and quantitated down to 25 ng through a single step extraction process. This provides a quick and easy testing method for quality control as a herbal medicine.

Asparacoside, shatavarin IX, shatavarin IV, asparanin A and shatavarin V were quantitated by HPLC-Q-TOF-MS/MS in the negative ion mode. Their structures, *m/z*, and retention times are shown in Figure 1 and their total ion current (TIC) chromatograms in Figure 2a. The fragmentation pattern of each saponin glycoside shows sequential loss of single monosaccharides leaving one glucose attached to *m/z* 577 [M+Glc-H]⁻ while the

aglycone is undetectable. Peaks corresponding to monosaccharides, glucose (*m/z* 161), rhamnose (*m/z* 145) and arabinose (*m/z* 131) are apparent (Figure 3). The method was validated for linearity, LOQs and LODs, precision, and recovery based on the International Conference on Harmonization (ICH) Guidelines [19] (Tables 1-3). Standard curves of the five saponin glycosides had acceptable linearity in the range of 2.50 - 30.00 µg/mL with correlation coefficients exceeding 0.997. The minimum concentration levels at which analysts could be reliably detected (LODs) and quantified (LOQs) were 5 to 10 ng and 25 to 50 ng, respectively (Table 1). The %RSDs of intra- and inter-day precisions were less than 4% and 6%, respectively (Table 2) and deemed acceptable according to the Horwitz's equation [20] (for analysts at concentration of 6 µg/mL, %RSD should be <12; for 15 µg/mL analyses, RSD should be <11 and for 25 µg/mL analysts, RSD should be <10). For overall accuracy recoveries ranged from 95% to 105%, with RSDs ranging from 0.7 - 4.5 %, again within ICH acceptable values [19] (Table 3). This suggests that sample extraction was efficient. Thus, the method developed is reliable and accurate for quantitation of the five saponin glycosides in *A. racemosus*.

Five saponin glycosides in different parts of *A. racemosus* from five different growing locations were quantified. None of the saponin glycosides were found in the leaves and stems of all *A. racemosus* tested except for trace amounts (less than LOQ) of asparacoside in the Bangkok sample. For root samples, saponin glycoside contents were diverse. Thus asparacoside from Phetchabun had 12 mg/g but



Compound	Formula	R ₁	R ₂	R ₃	RT, min	[M-H] ⁻
Asparacoside	C ₄₉ H ₈₀ O ₂₁	β-D-glucose	α-L-arabinose	α-L-arabinose	5.38	1003.4966
Shatavarin IX	C ₄₅ H ₇₄ O ₁₈	β-D-glucose	β-D-glucose	H	6.70	901.4766
Shatavarin IV	C ₄₅ H ₇₄ O ₁₇	β-D-glucose	α-L-rhamnose	H	7.04	885.4745
Asparanin A	C ₃₉ H ₆₄ O ₁₃	β-D-glucose	H	H	8.06	739.4162
Shatavarin V	C ₄₅ H ₇₄ O ₁₇	α-L-rhamnose	β-D-glucose	H	8.43	885.4721

Figure 1. Structures of saponin glycosides in *A. racemosus* root with their molecular formula, molecular weights as well as LC/MS retention time and [M-H]⁻

Table 1: Validation parameters of the five saponin glycoside standards using HPLC-Q-TOF-MS/MS

Saponin	[M-H] ⁻	^a Regression equation (n=3)	Linearity (R ²)	Linear range (µg/mL)	^b LOD (ng)	^c LOQ (ng)
Asparacoside	1003.4966	y = 13.20.x-30.43	0.999	5.00 - 30.00	10	50
Shatavarin IX	901.4766	y = 10.13.x-13.75	0.999	2.50 - 30.00	5	25
Shatavarin IV	885.4745	y = 17.79.x-3.16	0.999	2.50 - 30.00	5	25
Asparanin A	739.4162	y = 10.16.x-2.26	0.998	2.50 - 30.00	5	25
Shatavarin V	885.4721	y = 9.47.x-14.95	0.997	2.50 - 30.00	10	25

^ay is the peak area; x standard solution concentrations, non-weighted. R² is the determination coefficient. ^bLOD, limits of detection, S/N = 3. ^cLOQ, limits of quantitation, S/N = 10.

was undetectable in the Indian sample. Other compounds were also erratically distributed (Table 4). Shatavarin IV known to dominate the Indian varieties [9b] was not found in any Thai samples. The Rayong root sample contained most compounds (Table 4, Fig 2b). The reason for this diversity apparently is not genomic because genetic variation in plants harvested at 9 diverse Thai locations [21] and 7 Indian sites [22] showed little variation. This strongly suggests that the growing conditions play a major role in expression of these saponins [23]. It also implies that horticultural conditions can be optimized to produce the most biologically active components.

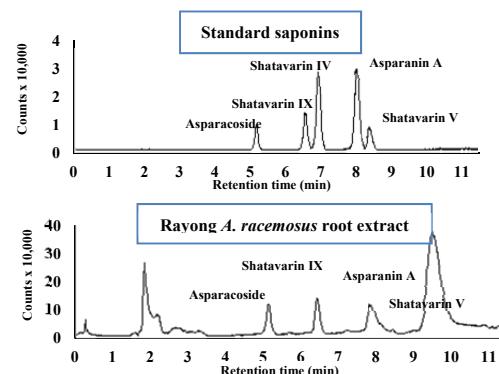


Figure 2. TIC chromatogram of (a) five saponin glycosides at concentration 10 µg/mL (b) *A. racemosus* root from Rayong, Thailand in 70% MeOH at concentration 20 µg/mL obtained from HPLC-Q-TOF-MS/MS analysis in negative mode.

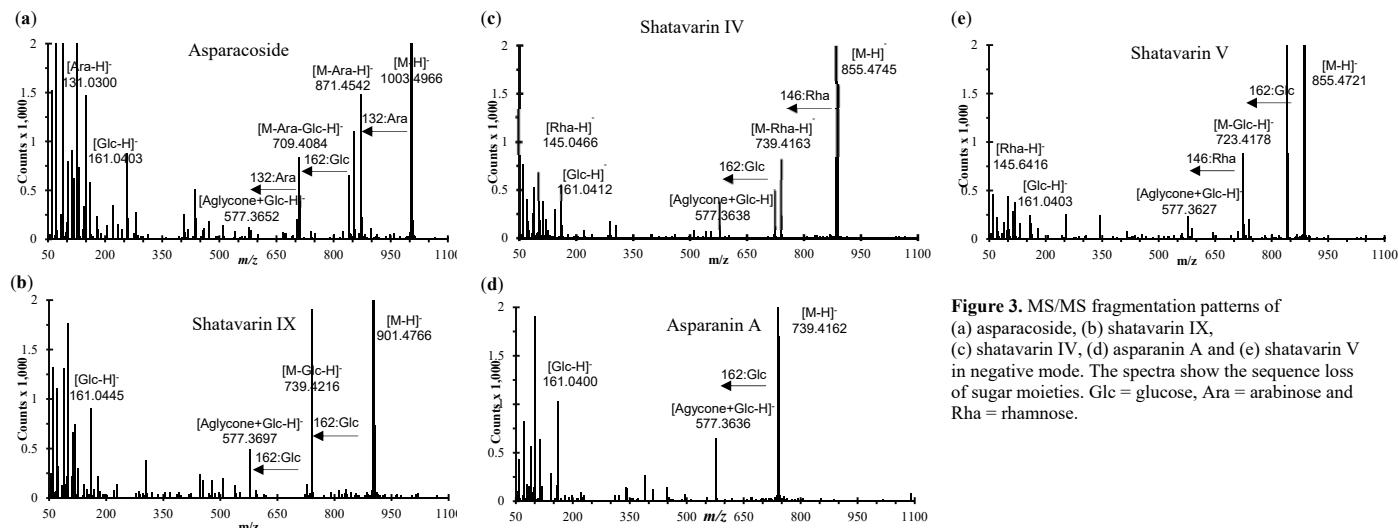


Figure 3. MS/MS fragmentation patterns of (a) asparacoside, (b) shatavarin IX, (c) shatavarin IV, (d) asparanin A and (e) shatavarin V in negative mode. The spectra show the sequence loss of sugar moieties. Glc = glucose, Ara = arabinose and Rha = rhamnose.

Table 2: Intra- and inter-day precisions of the five saponin glycoside standards by HPLC-Q-TOF-MS/MS.

Saponin	Spiked Conc. ($\mu\text{g/mL}$)	Intraday (n=5)			Interday (n=15)		
		Measured Conc. ($\mu\text{g/mL}$)	% Accuracy	% RSD	Measured Conc. ($\mu\text{g/mL}$)	% Accuracy	% RSD
Asparacoside	6	6.3 \pm 0.14	104.1 \pm 2.38	2.3	6.2 \pm 0.14	103.3 \pm 2.26	2.2
	15	15.4 \pm 0.22	102.6 \pm 1.44	1.4	15.6 \pm 0.37	104.2 \pm 2.48	2.4
	25	25.6 \pm 0.12	102.2 \pm 0.46	0.5	26.0 \pm 0.47	104.1 \pm 1.86	1.8
Shatavarin IX	6	6.1 \pm 0.19	102.1 \pm 3.15	3.1	6.1 \pm 0.31	101.3 \pm 5.15	5.1
	15	15.4 \pm 0.32	102.5 \pm 2.10	2.1	15.6 \pm 0.72	104.2 \pm 4.82	4.6
	25	25.7 \pm 1.05	102.8 \pm 4.19	4.1	25.7 \pm 0.66	102.9 \pm 2.63	2.6
Shatavarin IV	6	5.9 \pm 0.14	97.8 \pm 2.31	2.4	6.2 \pm 0.29	103.4 \pm 4.89	4.7
	15	15.5 \pm 0.35	103.2 \pm 2.32	2.3	15.5 \pm 0.50	103.0 \pm 3.35	3.3
	25	26.2 \pm 0.80	104.7 \pm 3.18	3.0	26.2 \pm 0.51	104.8 \pm 2.03	1.9
Asparanin A	6	6.3 \pm 0.17	104.3 \pm 2.81	2.7	6.2 \pm 0.34	102.5 \pm 5.61	5.5
	15	15.6 \pm 0.38	104.1 \pm 2.56	2.5	15.7 \pm 0.47	104.8 \pm 3.16	3.0
	25	25.9 \pm 0.42	103.6 \pm 1.70	1.6	25.9 \pm 0.76	103.5 \pm 3.03	2.9
Shatavarin V	6	6.1 \pm 0.24	101.1 \pm 4.04	4.0	6.0 \pm 0.29	100.7 \pm 4.79	4.8
	15	15.6 \pm 0.52	103.7 \pm 3.45	3.3	15.6 \pm 0.39	104.1 \pm 2.59	2.5
	25	25.4 \pm 0.71	101.6 \pm 2.85	2.8	26.0 \pm 0.79	103.9 \pm 3.15	3.0

Table 3: The accuracy HPLC-Q-TOF-MS/MS method for quantitative analysis of five saponin glycosides when three concentrations of standard saponin glycosides were spiked into the *A. racemosus* root from Bangkok.

Saponin	Measured concentrations of saponins ($\mu\text{g/mL}$) (n=3)		Recovery (%)	RSD (%)
	Spiked	Original	Detected	
Asparacoside	6	21.5 \pm 0.21	26.2 \pm 1.03	95.2 \pm 4.13
	15	24.2 \pm 0.26*	39.2 \pm 0.55*	100.0 \pm 1.21
	25	24.3 \pm 0.23*	52.9 \pm 0.71*	105.0 \pm 0.75
Shatavarin IX	6	-	6.2 \pm 0.14	103.9 \pm 2.30
	15	-	15.6 \pm 0.51	104.3 \pm 3.39
	25	-	26.2 \pm 0.52	104.9 \pm 2.07
Shatavarin IV	6	-	5.9 \pm 0.27	98.7 \pm 4.46
	15	-	15.7 \pm 0.21	104.4 \pm 1.43
	25	-	25.7 \pm 0.32	102.7 \pm 1.27
Asparanin A	6	-	6.2 \pm 0.19	104.0 \pm 3.22
	15	-	15.5 \pm 0.21	103.1 \pm 1.42
	25	-	26.2 \pm 0.38	104.9 \pm 1.54
Shatavarin V	6	-	6.2 \pm 0.13	103.4 \pm 2.09
	15	-	15.1 \pm 0.45	100.5 \pm 3.00
	25	-	24.8 \pm 0.19	99.0 \pm 0.74

* Samples were diluted for 3 times before injected to LC-MS.

The five saponin glycosides have the same aglycone (sarsasapogenin), but differ in monosaccharide moieties. Their bioactive properties require the aglycone but the nature of the monosaccharide residue influences potency as shown by tumor cell susceptibility to cytotoxic actions [24]. Therefore, the saponin profile of *A. racemosus* could dictate its potency and pharmacology and thus have unpredictable therapeutic outcomes. Only the types of methods described here, especially HPLC-Q-TOF-MS/MS, have the flexibility, sensitivity, accuracy, and selectivity for reliable routine quality control.

Table 4: The percentage of five saponin glycosides in roots of *A. racemosus* collected from different locations (n=3).

Sample	Saponin glycosides (mean \pm SD, mg/g)				
	Asparacoside	Shatavarin IX	Shatavarin IV	Asparanin A	Shatavarin V
Bangkok	1.1 \pm 0.01	nd	nd	nd	nd
Phetchabun	12.3 \pm 1.30	< 0.13	nd	nd	nd
Phitsanulok	1.2 \pm 0.06	0.5 \pm 0.05	nd	nd	nd
Rayong	2.6 \pm 0.09	2.5 \pm 0.13	nd	0.3 \pm 0.02	0.3 \pm 0.02
India	nd	0.2 \pm 0.01	3.1 \pm 0.09	nd	0.4 \pm 0.02

nd: not detected

The quantitative analysis of five saponin glycosides in *A. racemosus* was investigated by HPLC-Q-TOF-MS/MS for the first time. The extremely diverse expression of the compounds from plants taken from different location, highlights the importance of quality control when applied to clinical treatments. Only analyses as described here have the required attributes to provide the all-important quality control of *A. racemosus* formulations for such applications.

Experimental

Chemicals and reagents: MeOH, acetonitrile and water used for LC-MS/MS were LC-MS grades (RCI Labscan, Thailand). Sterile water was purchased from A.N.B. Laboratories Co., Ltd., Thailand. Analytical grade formic acid was purchased from Fisher Scientific (UK), standard grade shatavarin IV from Natural Remedies Pvt. Ltd. (India). Asparacoside, shatavarin IX, asparanin A and shatavarin V were isolated from *A. racemosus* extract as described previously [15] and their structures and purity were confirmed by NMR and MS.

Plant material and sample preparation: *A. racemosus* collected from Bangkok, Phetchabun, Phitsanulok and Rayong Provinces in Thailand and identified by Prof. Wongsatit Chuakul, Faculty of Pharmacy, University of Mahidol, Bangkok, Thailand. The plants were separated into leaves, stems and roots, cut and each dried separately at 45 °C for 3 days before milling into a fine powder. Indian *A. racemosus* dried root powder was purchased from Classic Ayurveda Herbal Products, Aurangabad, India. To extract the plant material, 1 mL of 70% (v/v) MeOH in water was added to 20 mg of each of the fine dried powders of leaf, stem or root samples, and sonicated in a TranssonicTP690 (Elma, Germany) for 15 min at maximum power, filtered through a 0.45 μm syringe filter and injected (10 μL) into the LC-MS system. All samples were prepared in triplicate.

Standard solution preparation: Asparacoside, shatavarin IX, shatavarin IV, asparanin A and shatavarin V were used as saponin glycosides standards in this study. Stock solutions of 30 $\mu\text{g/mL}$ in 70% (v/v) MeOH were appropriately diluted with 70% MeOH to construct standard calibration curves, limits of detection (LOD), and limits of quantification (LOQ) for the five saponins. All solutions were stored at 4 °C until analysis, sample injection volumes 10 μL .

Saponin analysis by HPLC-Q-TOF-MS/MS: Separation used a Phenomenex Luna C8 column (100 \AA , 5 μm , 150 x 4.6 mm, Phenomenex USA) connected with a SecurityGuardTM C8 guard column (100 \AA , 5 μm , 4 x 3 mm, Phenomenex, USA) and the column temperature was held at 35 °C. The mobile phase was composed of 0.1% formic acid in water (eluent A) and 0.1% formic acid in acetonitrile (eluent B). The gradient program was: 50:50% eluents A/B at the beginning, then 40:60% A:B for 5 min, then linearly decreased to 10:90% (A/B) over the next 4 min, and held for 3 min. The flow rates were 0.6 mL/min. The liquid chromatograph was coupled to a Q-TOF mass spectrometer with a dual electrospray ion (ESI) source (Agilent Technologies G6540B model, Agilent Technologies, Singapore). The MS parameters were: acquisition-mass range set at 100-1200 m/z with a 250 ms/spectrum, the drying gas was N₂ at 350 °C, flowing at 10 L/min, nebulizer pressure 20 psi, capillary voltage 3500 V, the fragmentor voltage 250 V, skimmer 65 V, octopole RFV 750 V. The target MS/MS was set up with collision energy at 10, 20 and 40 V. All the acquisition and analyses of data were controlled by Agilent LC-MS-Q-TOF MassHunter Data Acquisition Software version B.05.01 and Agilent MassHunter Qualitative Analysis Software B 06.0, respectively (Agilent Technologies, USA). Determinations were in both negative and positive ion modes but the negative mode gave higher peaks and was used to quantitate saponin glycosides.

Method validation: The established HPLC-Q-TOF-MS/MS method was evaluated by linearity, LOD, LOQ, intraday and interday precision, and accuracy tests. The calibration curves were constructed with five concentrations of saponin glycoside standards in duplicate. The LOD and LOQ were measured under the LC analytical conditions at a signal-to-noise (S/N) ratio of 3 and 10,

respectively. The repeatability (intraday precision), used three concentrations, (6, 15 and 25 µg/mL) measured 5-times within one day and interday precision by repeating measurements 5x per day for three consecutive days. Matrix effects and accuracy were determined using the *A. racemosus* root from Bangkok by spiking 20 mg of powders with 1 mL of 70% MeOH containing appropriate concentrations of standards (Table 3), and repeated in triplicate. After 15 min of sonication, filtrates were extracted and quantified as above.

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New Alkaloids and Anti-inflammatory Constituents from the Leaves of *Antidesma ghaesembilla*

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Using various chromatographic methods, two new alkaloids, antidesoic acids A (1) and B (2) along with fourteen known compounds (3-16) were isolated from the leaves of *Antidesma ghaesembilla* Gaertn. Their chemical structures were elucidated by physical and chemical methods. All the isolated compounds were evaluated for their inhibitory activity on LPS-stimulated nitric oxide (NO) production in BV2 cells and RAW 264.7 macrophages. Bisflavone **8** significantly inhibited LPS-stimulated NO production in BV2 cells and RAW 264.7 macrophages with IC₅₀ values of 5.4 and 8.0 μM, respectively. Compounds **1-3**, **7**, **10**, **12**, **14**, and **16** showed moderate inhibitory activities with IC₅₀ values ranging from 11.7 to 77.4 μM.

Keywords: *Antidesma ghaesembilla*, Euphorbiaceae, Alkaloid, Anti-inflammation, Antidesoic acid A, Antidesoic acid B.

Nitric oxide (NO) produced in large amounts by inducible nitric oxide synthase (iNOS) is known to be responsible for the vasodilation and hypotension observed during septic shock and inflammation [1]. NO mediates a variety of biological actions ranging from vasodilatation, neurotransmission, inhibition of platelet adherence and aggregation, as well as macrophage and neutrophil-mediated killing of pathogens [2].

Microglia are the resident macrophage-like cells of the central nervous system with a broad role in the brain's innate immunity and in inflammatory neuropathologies. Macrophages are also considered to play essential roles in inflammation. Although inflammation is a general immune system to protect the body from infection or tissue injury, overexpression of inflammatory mediators such as NO can cause cell damage and inflammatory diseases including rheumatoid arthritis [3], diabetes [4], and chronic hepatitis [5].

Antidesma, a genus of tropical plants belonging to the Euphorbiaceae family, comprises about 100 species, 29 of which are found in Vietnam. The leaves of *A. ghaesembilla* Gaertn. have been used in traditional medicine for the treatment of skin diseases and headache [6]. In addition, its fruits have been used to treat sore throat and lung diseases [6]. However, there are few reports about chemical constituents and biological activities of this plant [7].

In a drug discovery program for Vietnamese traditional medicinal plants for finding anti-inflammatory compounds, the methanol extract of *A. ghaesembilla* leaves was found to inhibit NO productions in LPS-stimulated BV2 cells and RAW264.7 macrophages with IC₅₀ values of 133.0 and 152.4 μg/mL, respectively. We report herein the isolation, structural elucidation, and evaluation of the inhibitory activity on NO production of two new and fourteen known compounds from the leaves of *A. ghaesembilla*.

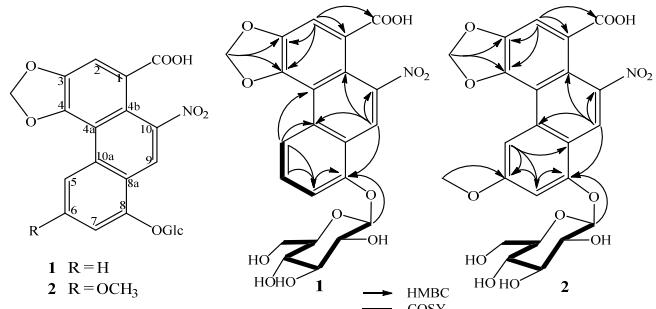


Figure 1: The chemical structures and key HMBC correlations of compounds **1** and **2**

Compound **1** was obtained as a yellow amorphous powder and its molecular formula was determined as C₂₂H₁₉O₁₂N from the HR-ESI-MS pseudo-molecular ion peak at *m/z* 488.0829 [M - H]⁻ (calcd. for C₂₂H₁₈O₁₂N, 488.0834). The ¹H NMR spectrum of **1** showed the presence of five aromatic protons at δ_H 7.45 (d, *J* = 8.5 Hz, 1H), 7.67 (t, *J* = 8.5 Hz, 1H), 7.69 (s, 1H), 8.75 (s, 1H), and 8.80 (d, *J* = 8.5 Hz, 1H); two methylenedioxy protons at δ_H 6.34 (s, 2H); and one anomeric proton at δ_H 5.19 (d, *J* = 7.5 Hz, 1H). The ¹³C NMR and DEPT spectra of **1** revealed signals of 22 carbons, including one carboxylic (δ_C 175.0), nine non-protonated carbons (δ_C 118.1, 119.0, 122.0, 131.9, 134.5, 145.4, 147.5, 149.1, and 155.7), ten methines (δ_C 71.3, 74.5, 78.1, 78.3, 102.5, 112.6, 113.6, 119.6, 122.1, and 131.0), and two methylene carbons (δ_C 62.5 and 103.4), suggesting the presence of a phenanthrene glycoside. The ¹H- and ¹³C-NMR data of **1** (Table 1) were very similar to those of aristolochic acid IIIa-6-O-β-D-glucopyranoside [8], except for the position of the sugar moiety at C-8. The HMBC correlations (Figure 1) between H-2 (δ_H 7.69) and C-1 (δ_C 118.1)/C-3 (δ_C 145.4)/C-4 (δ_C 147.5)/COOH (δ_C 175.0); between OCH₂O (δ_H 6.34) and C-3 (δ_C 145.4)/C-4 (δ_C 147.5) confirmed the presence of a carboxylic group at C-1 and a methylenedioxy group at C-3 and C-4. Acid hydrolysis of **1** afforded D-glucose as the sugar component

Table 1: ^1H - and ^{13}C -NMR data for compounds **1** and **2**.

C	1		2	
	$\delta_{\text{C}}^{\text{a},\text{b}}$	$\delta_{\text{H}}^{\text{a},\text{c}}$ (mult., J in Hz)	$\delta_{\text{C}}^{\text{a},\text{b}}$	$\delta_{\text{H}}^{\text{a},\text{c}}$ (mult., J in Hz)
Aglycone				
1	118.1	-	118.8	-
2	112.6	7.69 (s)	112.8	7.67 (s)
3	145.4	-	145.5	-
4	147.5	-	147.1	-
4a	122.0	-	12.0	-
4b	131.9	-	133.2	-
5	122.1	8.80 (d, 8.5)	104.5	8.37 (d, 2.0)
6	131.0	7.67 (t, 8.5)	162.9	-
7	113.6	7.45 (d, 8.5)	103.7	7.13 (d, 2.0)
8	155.7	-	157.0	-
8a	134.5	-	116.4	-
9	119.6	8.75 (s)	112.0	8.68 (s)
10	149.1	-	147.1	-
10a	119.0	-	118.7	-
COOH	175.0	-	175.0	-
3,4-OCH ₂ O	103.4	6.34 (s)	103.4	6.35 (s)
6-OMe			56.2	3.99 (s)
8-OGlc				
1'	102.5	5.19 (d, 7.5)	102.5	5.18 (d, 7.5)
2'	74.5	3.71 (dd, 7.5, 8.5)	74.9	3.71 (dd, 7.5, 8.0)
3'	78.1	3.57 (dd, 8.5, 9.0)	78.1	3.57 (dd, 8.0, 9.0)
4'	71.3	3.49 (t, 9.0)	71.4	3.48 (t, 9.0)
5'	78.3	3.57 (m)	78.4	3.57 (m)
6'	62.5	3.76 (dd, 5.5, 12.0)	62.5	3.76 (dd, 5.0, 12.0)
		3.94 (dd, 2.0, 12.0)		3.95 (dd, 2.0, 12.0)

^aMeasured in methanol-*d*₄, ^b125 MHz, ^c500 MHz, Assignments were made from HSQC, HMBC, and COSY experiments.

Table 2: IC₅₀ values of compounds **1–16** on LPS-stimulated NO production in BV2 cells and RAW 264.7 macrophages.

Compounds	IC ₅₀ (μM) BV2	IC ₅₀ (μM) RAW264.7
1	37.3 ± 8.7	77.4 ± 9.1
2	23.8 ± 3.1	20.3 ± 2.4
3	36.3 ± 3.8	53.2 ± 12.5
4	9.5 ± 1.3	>80
5	32.4 ± 9.9	>80
6	62.4 ± 11.2	>80
7	56.6 ± 5.7	54.5 ± 7.8
8	5.4 ± 0.6	8.0 ± 1.0
9	48.3 ± 7.3	>80
10	50.2 ± 5.4	79.1 ± 12.8
11	72.7 ± 20.9	>80
12	21.4 ± 4.4	11.7 ± 1.8
13	44.3 ± 8.9	>80
14	67.9 ± 26.0	76.7 ± 8.4
15	>80	>80
16	48.2 ± 6.8	64.4 ± 10.7

(identified as TMS derivative by GC) [9]. In addition, the HMBC correlation from glc H-1' (δ_{H} 5.19) to C-8 (δ_{C} 155.7); COSY correlations between H-5 (δ_{H} 8.80)/ H-6 (δ_{H} 7.67)/ H-7 (δ_{H} 7.45) were observed. These results indicated that the glucopyranosyl moiety was linked at C-8 of the aglycone. Finally, the position of the nitrogen dioxide group at C-10 was confirmed by HMBC correlation between H-9 (δ_{H} 8.75) and C-10 (δ_{C} 149.1), as well as by the HR-ESI-MS of **1**. Based on the above evidence, the structure of **1** was elucidated as aristolochic acid Ia-8-*O*- β -D-glucopyranoside.

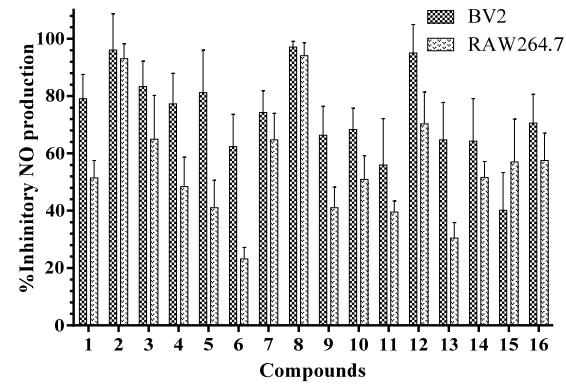
Compound **2** was obtained as a yellow amorphous powder and its molecular formula was determined as C₂₃H₂₁O₁₃N, by the HR-ESI-MS pseudo-molecular ion peak at *m/z* 542.0910 [M + Na]⁺ (calcd for C₂₃H₂₁O₁₃NNa, 542.0911). The ^1H NMR spectrum of **2** showed the presence of four aromatic protons at δ_{H} 7.13 (d, J = 2.0 Hz 1H), 7.67 (s, 1H), 8.37 (d, J = 2.0 Hz, 1H), and 8.68 (s, 1H); two methylenedioxy protons at δ_{H} 6.35 (s, 2H); one anomeric proton at δ_{H} 5.18 (d, J = 7.5 Hz, 1H), and one methoxy group at δ_{H} 3.99 (s, 3H). Two *meta*-coupled proton signals at δ_{H} 7.13 (d, J = 2.0 Hz)

and 8.37 (d, J = 2.0 Hz) in the ^1H NMR spectrum of **2** in comparison with **1** suggested a 1,2,3,5-tetra-substituted group in the third benzene ring of **2**. The ^{13}C NMR and DEPT spectra of **2** revealed signals of 23 carbons, including one carboxylic, ten non-protonated carbons, nine methines, two methylenes, and one methoxy carbon. The ^1H - and ^{13}C -NMR data of **2** (Table 1) suggested that the structure of **2** was similar to that of **1** except for the addition of a methoxy group at C-6. This was further confirmed by the HMBC correlation between the methoxy protons (δ_{H} 3.99) and C-6 (δ_{C} 162.9) (Figure 1). Similar to **1**, acid hydrolysis of **2** afforded D-glucose as the sugar component (identified as TMS derivative by GC). Furthermore, the sugar was linked at C-8, confirmed by the observation of an HMBC correlation between glc H-1' (δ_{H} 5.18) and C-8 (δ_{C} 157.0). Thus, the structure of **2** was elucidated as aristolochic acid IVb-8-*O*- β -D-glucopyranoside.

Compounds **1** and **2** were expected to be metabolic products of the related aristolochic acid uptake by the butterfly *Battus polydamas*, using HPLC-MS analysis [10]. However, to the best of our knowledge, this is the first report in which the chemical structure and NMR data of **1** and **2** are clearly provided. Thus, compounds **1** and **2** are considered to be new compounds, named as antidesoic acid A and antidesoic acid B, respectively.

Compounds **3–16** were identified as vitexin (**3**) [11], orientin (**4**) [11], isovitexin (**5**) [12], homoorientin (**6**) [11], luteolin-4'-*O*- β -D-glucopyranoside (**7**) [13], amentoflavone (**8**) [14], vanillyl alcohol 4-*O*- β -D-glucopyranoside (**9**) [15], 4-hydroxy-3,5-dimethoxybenzyl-*O*- β -D-glucopyranoside (**10**), 5-hydroxy-3,4-dimethoxyphenyl-*O*- β -D-glucopyranoside (**11**) [16], 3,4,5-trimethoxyphenyl-*O*- β -D-glucopyranoside (**12**) [17], sinapyl alcohol 4-*O*- β -D-glucopyranoside (**13**) [18], lyoniresinol (**14**) [19], syringaresinol (**15**) [20], and alangioside (**16**) [21] by comparison of their NMR and MS data with the reported values in the literature. This is the first report of compounds **3–6**, **9–14**, and **16** from the genus *Antidesma*.

All the isolated compounds were tested for their inhibitory activity on LPS-induced NO production in BV2 cells and RAW 264.7 macrophages. First, the compounds were examined at the concentration of 80 μM to screen their cytotoxicity and inhibitory activity on NO production. None of the tested compound showed significant cytotoxic activity (data not shown).

**Figure 2:** Inhibitory activity of compounds **1–16** on LPS-stimulated NO production in BV2 cells and RAW 264.7 macrophages at the concentration of 80 μM .

Nitrite concentrations were measured in the supernatant of BV2 cells and RAW 264.7 macrophages by the Griess reaction [22]. At the concentration of 80 μM , compounds **2** and **8** significantly decreased NO production in both BV2 cells and RAW264.7

macrophages with inhibitory percentages > 80%; compounds **1**, **3**, **5**, **7**, **10**, **12**, **14**, and **16** moderately decreased NO production with inhibitory percentages from 50.8% to 79.0% (Figure 2).

In order to obtain IC₅₀ values for the potent inhibitors, dose dependent responses were performed. Bisflavone **8** showed the most potent inhibitory activity on NO production in LPS-stimulated BV2 cells and RAW 264.7 macrophages with IC₅₀ values of 5.4 and 8.0 μM, respectively (Table 2). Compounds **1-3**, **7**, **10**, **12**, **14**, and **16** showed moderate inhibitory activities with IC₅₀ values ranging from 11.7 to 77.4 μM. Thus, it is possible to demonstrate that bisflavone **8** might be an important anti-inflammatory constituent of this plant.

Experimental

General: Optical rotation, Jasco DIP-370 automatic polarimeter; HR-ESI-MS, Varian 910 FT-ICR-MS 7 tesla; 1D and 2D NMR, Bruker AM500 FT-NMR spectrometer; Column chromatography was performed using either silica gel (Kieselgel 60, 70 – 230 mesh and 230 – 400 mesh, Merck, Whitehouse Station, NJ) or RP-18 resins (150 μm, Fuji Silysia Chemical Ltd.), and thin layer chromatography (TLC) using pre-coated silica-gel 60 F₂₅₄ (0.25 mm, Merck) and RP-18 F_{254S} plates (0.25 mm, Merck).

Plant material: The leaves of *A. ghaesembilla* were collected in Dak Lak province, Vietnam, in March 2013, and authenticated by Dr Nguyen Quoc Binh, Vietnam National Museum of Nature, VAST, Vietnam. A voucher specimen was deposited at the Institute of Marine Biochemistry, VAST.

Extraction and isolation: The dried leaves of *A. ghaesembilla* (2.5 kg) were sonicated in MeOH 3 times to yield 115.0 g of a dark solid extract, which was then suspended in water and successively partitioned with dichloromethane and ethyl acetate (EtOAc) to obtain dichloromethane (AG1, 30.0 g), ethyl acetate (AG2, 20.0 g), and water layers (AG3, 65.0 g) after removal of solvent *in vacuo*. The AG1 layer was chromatographed on a silica gel column with gradient mixtures of *n*-hexane/acetone (100/1 → 0/1, v/v) to obtain 5 fractions, AG1A-AG1E. Three smaller fractions (AG1E1-AG1E3) were obtained from the AG1E fraction by silica gel CC using dichloromethane/MeOH (18/1, v/v) as eluent. Compounds **14** (5.0 mg) and **15** (12.0 mg) was obtained from AG1E3 and AG1E1, respectively, using a RP-18 column and acetone/water (1/1.5, v/v). The EtOAc layer (AG2, 20.0 g) was chromatographed on a silica gel column eluting with dichloromethane/MeOH/water (5/1/0.1, v/v/v) to give 3 fractions, AG2A-AG2C. The AG2A fraction was chromatographed on a RP-18 column with MeOH/water as eluent (2/1, v/v) to yield compound **8** (20.0 mg). The AG2B fraction was chromatographed on a RP-18 column eluting with MeOH/water (1/1.5, v/v) to yield compound **3** (15.0 mg). The AG2C fraction was chromatographed using a RP-18 column then a silica gel column to yield compound **7** (7.0 mg). Compounds **1** (10.0 mg) and **2** (12.0 mg) were obtained from a RP-18 column using methanol/water (1/4.5, v/v), then a silica gel column using dichloromethane/methanol/water (2/1/0.1, v/v/v). The water layer (AG3, 65.0 g) was chromatographed on a Diaion HP-20 column eluting with water to remove sugar, then with an increased concentration of methanol in water (25, 50, 75, and 100 %) to obtain 4 fractions, AG3A-AG3D. The AG3A fraction was chromatographed on a silica gel column eluting with EtOAc/MeOH/water (5/1/0.5, v/v/v) to give 3 fractions, AG3A1-AG3A3. The AG3A1 fraction was chromatographed on a RP-18 column eluting with MeOH/water (1/2, v/v) to yield compounds **9** (6.0 mg) and **11** (5.0 mg). The AG3A3 fraction was chromatographed on a RP-18 column eluting with MeOH/water

(1/2, v/v) to yield compound **10** (5.0 mg). The AG3D fraction was chromatographed on a silica gel column eluting with CH₂Cl₂/MeOH/water (5/1/0.1, v/v/v) to give 4 fractions, AG3D1-AG3D4. Compounds **12** (7.0 mg) and **13** (5.0 mg) were obtained from the AG3D2 fraction using a RP-18 column and MeOH/water (1/2.5, v/v) as eluent. The AG3D3 fraction was applied on a silica gel column then a RP-18 column to yield compounds **4** (7.0 mg), **5** (8.0 mg), **6** (5.0 mg), and **16** (8.0 mg).

Antidesoic acid A (1)

Yellow amorphous powder.

[α]_D²⁵: -45.0 (c 0.1, MeOH).

HR-ESI-MS *m/z*: 488.0829 [M - H]⁻ (calcd for C₂₂H₁₈O₁₂N, 488.0834).

¹H- and ¹³C-NMR (CD₃OD): Table 2.

Antidesoic acid B (2)

Yellow amorphous powder.

[α]_D²⁵: -63.0 (c 0.1, MeOH).

HR-ESI-MS *m/z*: 542.0910 [M + Na]⁺ (calcd for C₂₃H₂₁O₁₃NNa, 542.0911).

¹H- and ¹³C-NMR (CD₃OD): Table 2.

Acid hydrolysis: Each compound (**1-2**, 2.0 mg) was separately dissolved in 1.0 N HCl (dioxane - H₂O, 1:1, v/v, 1.0 mL) and heated to 80°C in a water bath for 3 h. The solvent in acidic solution was removed under a N₂ stream. After extraction with CHCl₃, the aqueous layer was concentrated to dryness using N₂. The residue was dissolved in dry pyridine (0.1 mL), followed by addition of L-cysteine methyl ester hydrochloride in pyridine (0.06 M, 0.1 mL). The reaction mixture was heated at 60°C for 2 h. Trimethylsilylimidazole solution (0.1 mL) was then added, followed by heating at 60°C for 1.5 h. The dried product was partitioned with *n*-hexane and H₂O (0.1 mL each), and the organic layer was analyzed by gas chromatography (GC): column DB-5 (0.32 mm ID × 30 m length), detector FID, column temp 210°C, injector temp 270°C, detector temp 300°C, carrier gas He (2 mL/min). Under these conditions, the standard sugars gave peaks at *t*_R (min) 14.11 and 14.26 for D- and L-glucose, respectively. Peaks at *t*_R (min) 14.11 of D-glucose for **1** and **2** were observed.

Cell culture: BV2 microglial cells and RAW 264.7 macrophages were maintained at 37°C in a humidified atmosphere containing 5% CO₂ and 95% air at 5 × 10⁵ cells/mL in DMEM medium supplemented with 10% heat-inactivated FBS, penicillin G (100 units/mL), streptomycin (100 mg/mL), and L-glutamine (2 mM).

Nitrite production determination: BV2 microglial cells and RAW 264.7 macrophages were pretreated for 3 h with the compounds, and treated for 24 h with LPS (1 μg/mL). The nitrite concentration in the medium was measured as an indicator of NO production by the Griess reaction. In brief, an aliquot (100 μL) of each supernatant was mixed with an equal volume of Griess reagent (0.1%, w/v, *N*-(1-naphthyl)-ethylenediamine and 1%, w/v, sulfanilamide in 5%, v/v, phosphoric acid) for 10 min at room temperature. The absorbance of the final reactant at 525 nm was measured using an ELISA plate reader, and the nitrite concentration was determined using a standard curve of sodium nitrite prepared in DMEM without phenol red. The inhibitory effect on the NO production was estimated by calculating the concentration ratio of nitrite produced in the cells treated with LPS and with/without compounds.

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Chemical Characterization of *Narcissus poeticus* from Sirente –Velino (Apennines - Italy): Galantamine Accumulation and Distribution of Allergenic Compounds in the Flower

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Species of *Narcissus* (family *Amaryllidaceae*) are a potential source for large-scale extraction of alkaloids and fragrances. The bulbs typically accumulate a large number of alkaloids, including galantamine, a benzazepine alkaloid proven to be a cholinesterase inhibitor and which is used in the treatment of Alzheimer's disease. The presence of galantamine in *N. poeticus* L. collected in Abruzzo (Italy) was assessed and several levels of alkaloid were found in all parts of the plant (flower, stem, bulb and root) and not only in the bulb. The amount of galantamine obtained was tested by using two different extraction solvents. Extraction of *N. poeticus* absolute from the flowers was also performed, as this product is an important floral note in perfumery, and the distribution of allergenic compounds in the coronas and in the tepals was assessed. Moreover, the *in vitro* propagation of *N. poeticus* was tested as it may be a valuable resource from which to produce biomolecules, as an alternative to chemical synthetic processes.

Keywords: *Narcissus poeticus*, *Amaryllidaceae*, Absolute, Allergenic compounds, Bulbs, Fragrance, Galantamine, *in vitro* Propagation.

Narcissus poeticus L. is a wild monocotyledonous plant belonging to the *Amaryllidaceae* family. It is a native plant of the Mediterranean basin, widespread in the alpine areas of southern Europe, in Spain and Italy; it grows mainly at medium-altitude and on high mountains, in cool, moist and semi-shaded habitats [1]. It is a wild, bulbous, herbaceous plant, from 20 to 30 cm high; the leaves are radical, green, long and narrow (Figure 1a). Geophyta is the plant life form of the *Narcissus* genus; the storage organ contains reserves of carbohydrates and water and helps the plant to survive adverse environmental conditions such as cold or drought. It blooms in April-May, presenting an extremely fragrant flower per stem, characterized by a corona and a perianth tube which is pure white with six tepals fused at the base onto a green floral tube (5 mm diameter and 3 cm long). The corona (1.3 cm in diameter and 2-4 mm long) is light yellow, with a distinct reddish edge as shown in Figure 1b. Geophytes have economic value due to their use as a drug and as a fragrance for the perfume industry [2,3]. In recent years, the absolute fragrance obtained from flowers has been widely used in sophisticated modern perfumes due to its strong, green, woody and deep floral scent, which blends well with many of the floral absolutes, including clove bud carnation, jasmine, neroli, ylang ylang, and mimosa, as well as Karo-karounde. Ferri and co-workers have characterized the essential oil of *N. poeticus* flowers, which contained a great variety of components [4]. However, many of these are allergenic, as stated by Directive 2003/15 /EC, implemented in Italy by Legislative Decree n° 193, 3 February 2005 [5].

N. poeticus is also a valuable source of galantamine [6,7] a tertiary alkaloid, which is one of the therapeutic agents for the symptomatic treatment of Alzheimer's disease, the fourth leading cause of death in developed countries. Recent studies [8,9] supported the cholinergic hypothesis which postulates that memory impairment in patients with Alzheimer's disease results from a deficit of the

cholinergic function in the brain. As a lipophilic molecule able to go through the blood-brain barrier (BBB), the action of galantamine is to restore the level of acetylcholine by inhibiting the action of acetylcholinesterase enzymes [10].

Commercial drugs contain galanthamine bromohydrate as the active molecule, which is obtained by chemical synthesis through an expensive and complex process. The high costs and the growing demand for galantamine have driven scientists to obtain the active ingredient from plant matrices. Moreover, plant extracts often have a more effective and durable pharmacological action than monotherapies, as they contain different active substances that may have a synergistic action.

Previous work on the *in vivo* and *in vitro* propagation of plants containing compounds of particular interest both from Central Apennines (Italy) [11,12] and other sites [13], allowed us to gain knowledge about widely tested technologies. On these premises, in order to exploit the industrial potentialities of *N. poeticus* from Sirente-Velino (Abruzzo region, Italy), the aims of this investigation were: 1) to verify the presence and the amount of allergenic substances in the tepals and corona of the flowers; 2) to investigate the accumulation of galantamine in the different organs of plants; 3) to determine the galantamine content in the bulbs obtained by *in vitro* culture of *N. poeticus*.

Total yield and percentage of absolute of *n*-hexane extraction confirm our previous data [4]. The total yield (concrete) was found to be 0.45% of the weight of fresh flowers, that is higher than the values reported in the literature (0.20 and 0.26%) [1,4]. The percentage of absolute in the *n*-hexane extract was 30.5% of the weight of the concrete, in agreement with the literature [14]. The distribution of allergenic compounds in the tepals and corona are shown in Table 1. Highly allergenic substances, such as cinnamyl

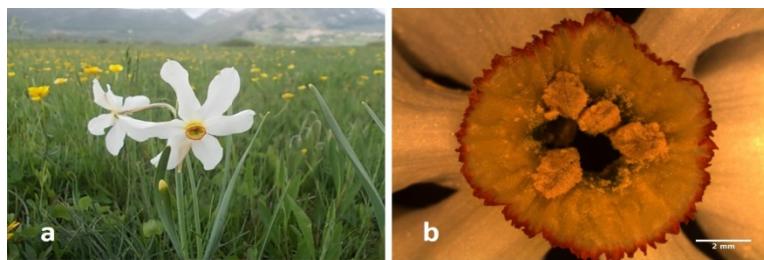


Figure 1: Flowers of *Narcissus poeticus* in Altopiano delle Rocche (a), detail of corona (b).

alcohol, which is the most abundant compound (30.2 µg/g), and cinnamyl aldehyde, are found only in the tepals. Less allergenic substances, such as benzyl benzoate and limonene, are mainly contained in the tepals (70 and 75% respectively), while 70% of isoeugenol, another highly allergenic compound, and 75% of eugenol, a less allergenic substance, are contained in the coronas. The tepals showed a greater amount (total mean value 40 µg/g) of more allergenic compounds compared with the corona (total mean value 15 µg/g). These data are also confirmed for the less allergic components (25 µg/g detected in tepals and 12 µg/g in corona).

Table 1: Percentage distribution of compounds in tepal and corona of *Narcissus poeticus* flowers after *n*-hexane extraction.

Compound	µg/g fresh flower	Tepals %	Coronas %
Cinnamyl alcohol **	30.2	100.0	-
Benzyl benzoate *	28.5	70.0	30.0
Isoeugenol methylether	28.1	40.3	59.7
cis Ocimene	25.2	21.6	77.4
Isoeugenol **	22.9	36.4	63.6
Eugenol methyl ether	20.3	35.5	64.5
α-Terpineol	19.9	1.3	98.7
Eugenol *	16.0	25.0	75.0
Benzyl alcohol	8.8	59.7	40.3
Methyl benzoate	5.8	64.8	35.2
Phenyl propyl alcohol	5.0	30.7	69.3
p-Cresol	3.9	75.0	25.0
Nonaldehyde	3.2	45.8	54.2
Benzyl salicylate	3.0	34.7	65.3
Cinnamyl aldehyde **	1.4	100.0	-
p-Methyl anisole	1.2	53.2	46.8
p-Propenyl anisole	1.2	60.0	40.0
Cinnamyl acetate	0.9	39.9	60.1
Limonene*	0.8	75.0	24.0

* Less allergenic compound; ** More allergenic compound.

As the extraction of absolute from tepals and coronas separately shows that the allergenic substances are mostly found in tepals, thus in the preparation of perfumes a safer absolute can be obtained from the corona only. This is due to the fact that despite preserving its distinctive fragrance it contains a lower quantity of allergenic components. Figure 2 shows the trend of galantamine in roots, bulbs, stems and leaves, and flowers in three phenological stages over four consecutive years. The means show that all parts of the plant contain galantamine, although the greatest amount was found in the underground organs. However, our experimental data showed an accumulation of the alkaloid in stems, leaves and flowers. At blooming stage the galantamine content was highest in both the bulbs and roots (respectively 12 mg /100 g dry weight and 13 mg / 100 g dry weight). The aerial parts, including flowers, stem and leaves (unique value), contain about 9 mg / 100 g dry weight; thus the amount of galantamine in the plant was about 34 mg / 100 g dry weight.

The highest content of galantamine was found in the bulb and root. Table 2 shows the differences observed in the various years at blooming time [15].

The data show slight differences for 2012, 2013 and 2014. In 2015 the lowest contents of alkaloid in both the underground organs

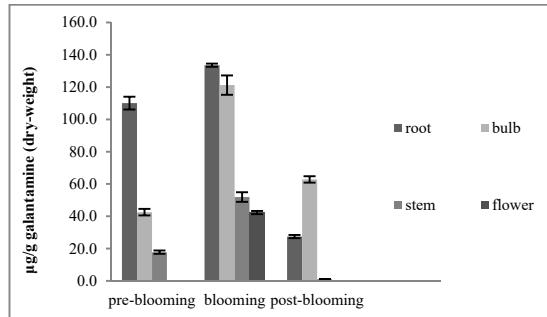


Figure 2: Galantamine distribution at different development stages of *N. poeticus*.

Table 2: Galantamine distribution (µg/g dry weight) in underground organs of *Narcissus poeticus* in different seasons.

Season	Root		Bulb	
	2012	2013	2014	2015
2012	122.0 ± 2.2		110.2 ± 10.5	
2013		137.5 ± 1.8	134.5 ± 3.2	
2014			133.6 ± 1.2	121.3 ± 5.7
2015				77.3 ± 3.1
				63.8 ± 2.9

might be due to a sudden rise in temperatures with subsequent drought during the spring (2015 was registered as the warmest year of the last 100 years; <http://www.noaa.gov/>). A further galantamine extraction both with methanol and ethanol was carried out on one additional sample in order to verify the use of a less-toxic solvent. Bulbs, roots, flowers, stems and leaves, before, during and after blooming were gathered and processed according to the procedures described in the Experimental section: Galantamine extraction.

Figure 3 shows the differences in the efficiency of extraction when ethanol and methanol were used. The distribution of values are plotted separately per group of cases (30) defined by values of categorical (grouping) variables (bulbs, flowers, roots, stems and leaves together). During pre- and post-blooming the content of galantamine extracted from the various organs of the plant was similar using either EtOH or MeOH (in the first case a greater variability among samples was detected). In the blooming stage, when the concentration of alkaloid is greater, the use of MeOH provides the best results in terms of increased efficiency of extraction from the different tissues of the plant, in particular from its roots. From the results of the statistical tests (Supplementary Data: 1-3) we can conclude that the differences between the two samples are statistically significant ($p < 0.00005$). Although the yield obtained by ethanol is good, methanol is a more efficient solvent to extract galantamine from *N. poeticus*.

Primary sterile explants of bulbs of *N. poeticus* were obtained in a medium contained benzylaminopurine (BAP), and naphthalene acetic acid (NAA). Such tests were carried out in order to grow parts of plants *in vitro* (Figure 4) on different media (G) (Supplementary Data: 4).

Bulbs obtained by *in vitro* propagation (Figure 4 b) were extracted with methanol and processed according to the procedure described in the Experimental section. The resulting galantamine content was

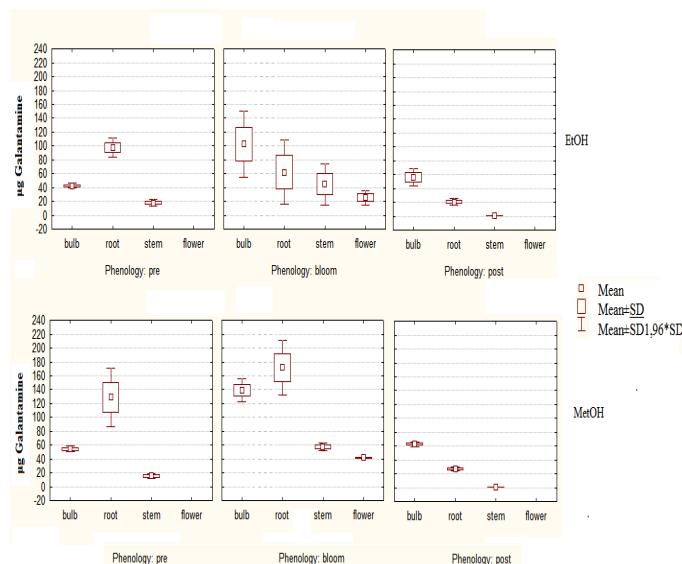


Figure 3: Distribution of galantamine extracted by ethanol (EtOH) and methanol (MeOH) in the three different phenological phases.

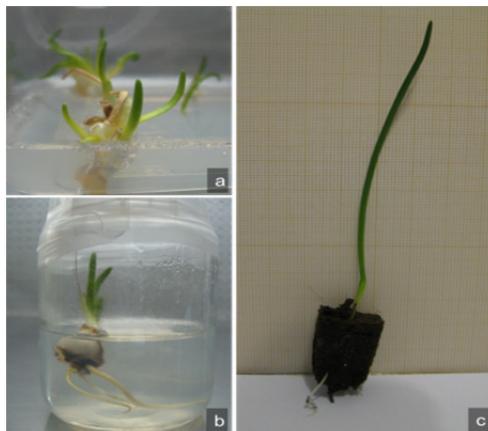


Figure 4: *In vitro* propagation of *Narcissus poeticus*
a) Shoots produced during the multiplication stage.
b) Rooted bulblet exhibiting three leaves.
c) Plants obtained from *in vitro* propagation transplanted to soil

2.5 mg ± 0.9 / 100 g dry weight, five times less than the amount obtained from wild bulbs. This preliminary result encourages us to continue the development of *in vitro* culture of *N. poeticus* bulbs. The organs, including bulbs, microtubers, corms, somatic embryos and shoots are perfect for mass propagation by using bioreactor techniques for large-scale propagation of geophytes, as described for *Gladiolus*, *Narcissus* and *Crocus* [16].

In conclusion, we have demonstrated that all parts of *N. poeticus* contain galantamine, confirming the presence in the species of this important alkaloid. The highest amounts are found in the underground organs (roots and bulbs). The accumulation of alkaloid is also documented in *in vitro* conditions opening new perspectives to biotechnological applications (i.e. bioreactors for mass propagation of adventitious roots), with the advantages to reduce time and costs. Moreover, with the goal to enhance the exploitation of the species for the local economy, the biochemical characterization of the different parts of the flowers pave the way towards a target product for the perfume industry. In summary, *N. poeticus* offers potential opportunities to increase the economy of local mountain communities to the development of marginal areas.

Experimental

Chemicals: Solvent: *N*-hexane HPLC grade, methanol HPLC grade, ethanol HPLC grade, dichloromethane; pure standards, agar, benzylaminopurine, naphthalene acetic acid, starch, sucrose and Murashige and Skoog medium were supplied by Sigma-Aldrich Chemie (Steinheim, Germany).

Plant materials: Wild plants of *Narcissus poeticus* were collected before blooming, during blooming and after blooming in a 300 sq m area of the plateau of Rocca di Mezzo-Sirente Velino Regional Park (L'Aquila, Italy), located at 13°31'35.9''E-42°13'16.3''N-1327 m asl, in 2012, 2013, 2014 and 2015. About 30 plants at each plant stage were collected in April, May and July. Plants were separated into bulbs, roots, flowers and stems. The collected materials were frozen in liquid nitrogen and then powdered using a mortar. The ground materials were freeze-dried for 48 h at -50°C and at about 0 KPa by an EDWARDS 5 Pascal LIO-5PDGT freeze-dryer and then stored at -20°C.

About 2000-2500 fully blooming flowers with 20 cm stems (500 – 650 g) were picked in the early morning during the May campaigns. The samples were preserved at 4°C in closed plastic tanks in a nitrogen atmosphere and extractions were performed the day after collection. Before each extraction the fresh flowers were manually reduced to small pieces of about 0.2 cm size.

Essential oil extraction: With the aim of evaluating the distribution of the allergenic compounds in different parts of the flower, tepals were separated from coronas (Figure 1). About 1 g of fresh flowers, corona and tepals were extracted separately in 10 mL of *n*-hexane for 3 h at room temperature under agitation. Then the solvent was evaporated under reduced pressure using a rotary evaporator and each extract was transferred with *n*-hexane to a volumetric flask, diluted to volume and analysed. To determine the yield of essential oils, 85 g of fresh flowers were extracted with 850 mL *n*-hexane and the concrete obtained (absolute and waxes). The hexane was removed by evaporation and the concrete was treated with ethanol cooled at -18°C to precipitate waxes. In order to improve the separation of waxes from the absolute, the mixture was centrifuged at 1500 x g for 30 min using a refrigerated centrifuge cooled at -15°C. The difference in weight between the concrete and waxes determined the absolute.

Galantamine extraction: Fifty mg of bulbs, roots, flowers and stems were transferred to a 1.5 mL Eppendorf tube with 1 mL of methanol adjusted to pH 4 with 25% ammonia solution. Each sample (3 samples were prepared for each vegetable matrix) was macerated with methanol for 24 h by sonication for 30 min every 8 h in an ultrasonic bath at room temperature. After maceration, extracts were centrifuged at 7800 x g for 10 min., and supernatants analysed [17]. The same extraction procedure was applied using ethanol instead of methanol, to assess the use of a less toxic solvent. The supernatant was added to 10 mL of NaHCO₃ 0.1M, and then galantamine was extracted with 2 mL CH₂Cl₂. The organic phase was dehydrated using Na₂SO₄ and concentrated under a gentle stream of nitrogen to 0.5 mL.

GC-MS analyses: Analyses were performed with a GC-MS-Thermo TraceMS spectrometer using a 60 m long capillary column with an internal diameter of 0.25 mm; the bonded phase was methyl-5% phenylsilane of thickness equal to 0.25 mm (column DB-5 MS, J & W Scientific). The detector was a selection quadrupole mass with electron impact ionization 70 eV, used in scanning (550-50). The operating conditions were: carrier gas helium N55, flow 1 mL min⁻¹; volume injected 1 mL. Injector: splitless (60 sec) at 250°C;

transfer line: 320°C; temperature program: 60°C for 1 min., ramp of 30°C min⁻¹ up to 100°C held for 1 min., ramp of 3°C min⁻¹ up to 300°C. The instrument was equipped with a NIST software library for the recognition of the mass spectra of the considered analytes. All experiments were carried out in triplicate and on at least 2 separate occasions.

In vitro culture: Bulbs of *N. poeticus* were gathered from flowering plants during the spring. The samples without papery scales and roots were washed with soap, rinsed with tap water and then disinfected according to the following scheme: Tween 20 for 15 min then washed several times with sterilized distilled water; 70% ethanol (3 min), followed by 10% commercial bleach (30 min), and finally rinsed several times with sterile distilled water. Bulbs were then vertically cut through two-thirds of their height (leaving one-third of its basal part intact). From each bulb two-thirds of the apical portion was removed, as well as the two external scales; a thin layer of the basal plate was also discarded. The remaining portion of the bulb was longitudinally divided to obtain explants formed by segments of twin-scales and a thick segment (2–3 mm) of the basal plate tissue. Twin-scales were used as primary explants cultured on a modified Murashige and Skoog medium with 2% sucrose supplemented with different growth regulators {benzylaminopurine (BAP) and naphthalene acetic acid (NAA)} at different

concentrations [18–20]. The media were solidified with 0.6% agar at pH 5.7 and autoclaved at 121°C (0.1 MPa) (Supplementary data 4). The explants were placed vertically into the culture media with the basal plate tissue inserted into the medium [21–23]. For each culture medium 80 explants were used. About 42 bulbs were used for shoot initiation; from each bulb an average of 8–10 explants were obtained, which were divided into the media. All cultures were incubated in a growth chamber at 22°C with neon Osram L30/41 lamps and a photoperiod of 14 h.

Statistical analysis: The STATISTICA 8 package (Statsoft) was used. Two samples of galantamine were assessed: one sample extracted with methanol and the other with ethanol. Each sample was achieved from plant organs (bulbs, stems plus leaves, roots, flowers) in 3 phenological stages (before blooming, during blooming and after blooming) for a total of 30 cases. Means and standard deviations of data from the 2 samples are shown in box plots (supplementary data 1). In order to evaluate differences between the 2 samples the *t*-test (Supplementary data 2) was used, followed by homoscedasticity and normality of the distributions (Shapiro-Wilk test, Supplementary data 2). Finally, the differences between the means were assessed by the Wilcoxon non-parametric test (Supplementary data 3).

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A New Imidazole from the Sponge *Dercitus (Halinastra) japonensis*Hiromi Hirade^a, Takahiro Haruyama^b, Nobuyuki Kobayashi^b, Nicole J. de Voogd^c and Junichi Tanaka^{a,*}^aDepartment of Chemistry, Biology and Marine Science, University of the Ryukyus, Nishihara, Okinawa 903-2013, Japan^bCentral Research Center, AVSS Corporation, Nagasaki 852-8137, Japan^cNaturalis Biodiversity Center, P.O. Box 9517, 2300 RA Leiden, The Netherlands

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A new imidazole sulfate (**1**) and three known compounds (**2-4**) were isolated from the sponge *Dercitus (Halinastra) japonensis*. The structure of compound **1** was elucidated by spectroscopic means. Compound **2** was confirmed to show anti-HIV activity, whereas compounds **1**, **3** and **4** were inactive.

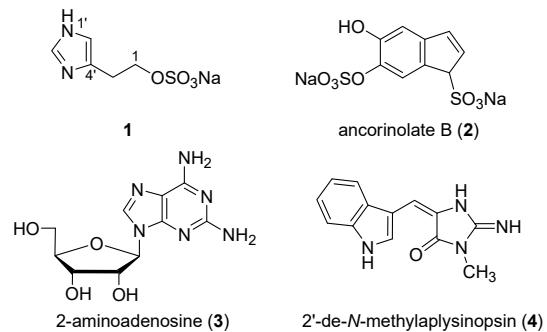
Keywords: Anti-HIV, Ancorinolate B, 2-Aminoadenosine, 2'-De-N-methylaplysinopsin, *Dercitus (Halinastra) japonensis*.

Human immunodeficiency virus (HIV) was discovered in the middle 1980s as a causative agent of acquired immune deficiency syndrome (AIDS). Current therapy, named HAART (Highly Active Anti-Retroviral Therapy), against the disease by administering “cocktails” of drugs prevents the patients from development of the sickness, but they have to keep taking the medication as HAART cannot eradicate the virus [1]. However, there exists a demand for newer drugs for a complete cure.

In our quest for new antiviral drug leads not only for HIV, but for several viruses, we screened a library of 800 extracts of coral reef organisms collected along the islands of Okinawa. Chromatographic study of an anti-HIV extract led to four molecules (**1-4**) being obtained. Although the antiviral activity was ascribed to compound **2**, a new compound **1**, separated together with **2**, is the subject of this note.

As the MeOH soluble extract of the sponge *Dercitus japonensis* showed anti-HIV activity, a portion of it was separated by gel filtration and reversed phase HPLC resulting in the isolation of compound **1** with three known compounds: ancorinolate B (**2**) [2], 2-aminoadenosine (**3**) [3], and 2'-de-N-methylaplysinopsin (**4**) [4]. Each known compound was identified by comparing its spectroscopic data with those published.

Compound **1** was obtained as a glassy material and its negative HR-ESIMS showed pseudomolecular ions as $C_5H_7N_2O_4^{32}S$ ($[M-Na]^-$, $\Delta -0.27$ mmu) and $C_5H_7N_2O_4^{34}S$ ($[M-Na]^-$, $\Delta -1.06$ mmu) indicating three degrees of unsaturation without the valence of a sulfur atom. The 1H NMR data contained characteristic low-field signals at δ 7.41 (1H, brs) and 8.87 (1H, d, $J = 1.1$ Hz), while the ^{13}C NMR spectrum showed three sp^2 signals at δ 116.4 (CH), 131.0 (C), and 133.6 (CH) suggesting the presence of a heteroaromatic moiety. With the presence of two nitrogen atoms in the molecule, it is likely to have a monosubstituted imidazole or pyrazole ring. Another structural feature is two consecutive methylenes at δ 2.91 (t, $J = 6.4$ Hz) and 3.99 (t, $J = 6.4$ Hz) showing COSY cross peaks with each other and also ^{13}C NMR signals at δ 24.8 and 63.8. Judging from the chemical shifts, one of the methylenes is oxygenated. The remaining structural unit was assigned as a sulfate (1223, 1062 cm^{-1}) attached to the oxygenated methylene.

Figure 1: Structures of isolated compounds **1-4**.

The sulfate was confirmed as a sodium salt by atomic absorption spectroscopy (AAS). Therefore, the structure of compound **1** was elucidated as an imidazole or a pyrazole with a sulfated ethanol moiety.

HMBC correlations between the aromatic portion and the ethyl group supported the two possible structures, however, it was not conclusive to determine either a pyrazole or imidazole derivative. Since aromatic proton and carbon signals of compound **1** showed close similarity to those (δ 7.39 s, 8.91 d, $J = 1.4$ Hz; δ 116.1, 131.5, 133.2) of 2-(1*H*-imidazol-4-yl)-ethanol (**5**) and also to those (δ 7.53 s, 9.04 d, $J = 1.3$ Hz; δ 116.7, 128.9, 133.7) of histamine dihydrochloride (**6**), an imidazole moiety was proved in **1**. By treating compound **5** with sulfuric acid in the presence of dicyclohexylcarbodiimide (DCC) [5], a trace amount of compound **1** was formed, giving a pseudomolecular ion ($\Delta -1.71$ mmu). Thus, we concluded that compound **1** is sodium 2-(1*H*-imidazol-4-yl)-ethyl sulfate.

Figure 2: Structures of reference compounds **5** and **6**.Table 1: Anti-HIV activity (IC_{50}) and cytotoxicity (CC_{50}) of compounds **1-5**.

Compounds	IC ₅₀ , μM	CC ₅₀ , μM
1	n.d.	> 93
2	109	> 2.84×10 ²
3	> 2.6×10 ²	65.2
4	> 4.2×10 ²	3.8×10 ²
5	n.d.	> 1.8×10 ²
AZT (nM)	9.0	> 100.0

n.d.: not determined

The anti-HIV activity of compounds **1–5** were evaluated as shown in Table 1. Compound **2** showed moderate activity, but the other compounds did not show any activity.

Experimental

General: HPLC separation was carried out on either a Nacalai Cosmoseil C18 (10×250 mm) or HILIC column (4.6×250 mm, 10×250 mm) with an HPLC unit composed of RI and UV detectors. NMR spectra were recorded on a Bruker AVANCE III spectrometer either in dimethylsulfoxide (DMSO)-*d*₆ or D₂O. ESIMS data were recorded on a Jeol JMR-T-100LP mass spectrometer. FTIR and UV spectra were taken on Jasco FT/IR-6100 and Jasco V-660 spectrophotometers, while atomic absorption was recorded on a Hitachi Z-2010 instrument. All solvents used were reagent grade (Wako). Compounds **5** (Synthonix) and **6** (TCI) were purchased.

Sponge: A specimen of the sponge *Dercitus (Halinastra) japonensis* van Soest, Beglinger & de Voogd (2010) was collected by hand with scuba at a reef of Yonaguni Island, Okinawa, September 1992. The specimen was brought back to the laboratory and kept frozen at -20°C until extraction. The sponge was identified by NJdV and a voucher specimen coded RMNH POR 8684 is deposited at Naturalis Biodiversity Center.

Extraction and isolation: The specimen (0.70 kg, wet weight) was soaked in acetone (1.0 L) 3 times at room temperature. The acetone solution was filtered and concentrated under vacuum, and the resulting material was partitioned between water and ethyl acetate (EtOAc). The water layer was concentrated and the residue washed with methanol (MeOH). The MeOH solution was concentrated to give a crude extract (49.8 g). As this extract showed inhibition against HIV, part of it (505.3 mg) was separated on a Sephadex LH-20 (Pharmacia) column (20 x 300 mm) using aqueous MeOH to give 8 fractions. The sixth fraction was found to contain compound **3** (16.7 mg). While waiting for anti-HIV assay results, we separated the last fraction to obtain compound **4** (11.6 mg) by C18 HPLC (0.01% TFA in CH₃CN). As the fourth fraction showed anti-HIV activity, it was further separated by HILIC HPLC (CH₃CN:H₂O, 9:1) to give pure compound **2** (15.8 mg) and impure compound **1** (2.5 mg). To supply an additional amount of compound **1**, another portion of the extract (303.3 mg) was successively separated by Sephadex LH-20 (aqueous MeOH), C18 HPLC (water), and HILIC HPLC (CH₃CN:H₂O, 9:1) to give pure compound **1** (3.6 mg).

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Compound (1)

Colorless glass.

IR (film): 3154, 1628, 1223, 1062, 998, 770, 626, 437 cm⁻¹.

UV/Vis λ_{max} (MeOH): 210 nm (log ϵ) 3.69.

¹H NMR (500 MHz, DMSO-*d*₆): δ 8.87 (1H, d, *J* = 1.1 Hz, H-2'), 7.41 (1H, brs, H-5'), 3.99 (2H, t, *J* = 6.4 Hz, H-1), 2.91 (2H, t, *J* = 6.4 Hz, H-2).

¹³C NMR (125 MHz, DMSO-*d*₆): δ 133.6 (CH, C-2'), 131.0 (C, C-4'), 116.4 (CH, C-5'), 63.8 (CH₂, C-1), 24.8 (CH₂, C-2).

COSY: H-1/H-2.

HMBC: H-1/C-2,4', H-2/C-1,4', H-2'/C-4',5', H-5'/C-2',4'.

HRESIMS: *m/z* 191.01238 [M-Na]⁺; calcd for C₅H₇N₂O₄³²S 191.01265; *m/z* 193.00739 [M-Na]⁺; calcd for C₅H₇N₂O₄³⁴S 193.00845.

AAS: c_{Na} = 0.23 ppm.

Sulfation of 5: Compound **5** (1.0 mg) in dimethylformamide (DMF, 150 μL) was mixed with DCC (63.0 mg) in DMF (300 μL) and sulfuric acid (3.2 μL) in DMF (150 μL). The reaction suspension was stirred overnight at rt. After adding water, the mixture was filtered and the filtrate partitioned against CH₂Cl₂. The water layer was concentrated to give a crude product (56.1 mg), which was separated by HILIC HPLC (CH₃CN:H₂O, 9:1) to give a fraction (4.7 mg) containing compound **1**.

HRESIMS: *m/z* 191.01094 [M-Na]⁺; calcd for C₅H₇N₂O₄³²S 191.01265.

Anti-HIV assay: For evaluation of anti-HIV activities, human T-cell line MT-4 cells were seeded in a 96-well plate at a density of 5 × 10³ cells/mL in 100 μL /wells of Roswell Park Memorial Institute (RPMI) 1640 medium supplemented with 10% fetal bovine serum and then cells were infected with HIV-1 strain IIIB at a viral multiplicity of infection of 0.001 in the presence of various concentrations of each compound. Moreover, to confirm the cytotoxic effects, MT-4 cells were inoculated with each compound at various concentrations without virus infection. After incubation at 37°C in an atmosphere of 5% CO₂ for 5 days, cell viability was determined by WST-1 assay (Dojindo Molecular Technologies, Inc.) that measured absorbance at 450 and 650 nm using the sodium salt of 4-[3-(4-iodophenyl)-2-(4-nitrophenyl)-2H-5-tetrazolio]-1,3-benzene disulfonate, which produces a highly water-soluble formazan dye. The calculation of the 50% virus inhibitory concentration (IC₅₀) and the 50% cytotoxic concentration (CC₅₀) of each compound were performed by statistical analysis software, GraphPad Prism (GraphPad Software, Inc.).

Acknowledgments - We thank Dr Chiaki Tanaka, Kyushu University, for the database search. This research was supported by a grant from Okinawa Intellectual Cluster Programs of Okinawa Science and Technology Center.

Acetyl-cholinesterase Inhibitory Activity of Methoxyflavones Isolated from *Kaempferia parviflora*

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MeOH extracts of *Kaempferia parviflora* Wall. ex. Baker, family Zingiberaceae, were consecutively partitioned with CHCl₃, EtOAc, and *n*-BuOH. The CHCl₃ fractions were diluted in distilled water with *n*-hexane-CH₂Cl₂ and three methoxyflavones were isolated from the CHCl₃ extract. Based on spectral analysis and comparison of the spectral data with literature values, the compounds were identified as 3,5,7,3',4'-pentamethoxyflavone (KP1), 5,7-dimethoxyflavone (KP2), and 5,7,4'-trimethoxyflavone (KP3). In relation to their possible effectiveness against Alzheimer's disease, these compounds were tested for their ability to inhibit acetylcholinesterase activity and neurite outgrowth in the PC12 cell line. Of the three compounds, KP1 was the only one to inhibit significantly the acetylcholinesterase activity in a dose-dependent manner.

Keywords: *Kaempferia parviflora*, Zingiberaceae, Acetylcholinesterase inhibitory, Methoxyflavone,

Kaempferia parviflora (KP; family Zingiberaceae) is endemic in Thailand and can be found cultured in the wild in Pitsanulok and Phetchabun. Locally, it is named Krachai Dum [1]. The KP plant grows low to the ground and has short, fleshy, and tuberous dark purple rhizomes and small violet flowers [2]. The black or purple color of KP rhizomes is affected by the conditions of the soil environment. The climatic and soil conditions of South Korea are not ideal for cultivation of this plant.

KP ethanol or methanol extracts are effective for treating human disorders such as visceral fat accumulation, hyperinsulinemia, glucose intolerance, hypertension, diabetes, obesity, peripheral neuropathy, gastric ulcer, and Alzheimer's disease (AD) [3]. Compounds isolated from KP include polymethoxyflavonoids and the phenolic glycoside kaempferioside [4]. Moreover, phytochemical screening of KP extracts has produced positive tests for alkaloids, anthrones, and coumarins [5]. 5,7,4'-Trimethoxyflavone, 5,7,3'4'-tetramethoxyflavone, and 3,5,7,4'-methoxyflavones have exhibited significant inhibitory activity against various biological processes [4].

AD is one of the principal causes of disease and death among the elderly [6]. The disease has numerous important confirmed facets, including the hallmark β -amyloid plaques [3]. There have been many different ways and attempts at curing and preventing AD. In recent years, cholinesterases have received increasing attention for the role they play in the formation from β -amyloids during the early stages of AD [7]. Acetylcholinesterase (AChE) serves as the key enzyme in AD, as it hydrolyzes the ester bond (at thiocholine and acetic acid) of the neurotransmitter acetylcholine in the cholinergic synapse [6].

The current study describes the isolation of three methoxyflavones from *K. parviflora*. The molecular structures were determined through ¹H NMR spectroscopic, mass spectrometric and literature data. Compounds KP1, KP2, and KP3 were dissolved in DMSO. KP1, at the low concentration of 40 μ M, exhibited strong inhibitory activity (80%) against AChE. KP2 and KP3, on the other hand, exhibited weak AChE-inhibitory activity (20%) at a concentration

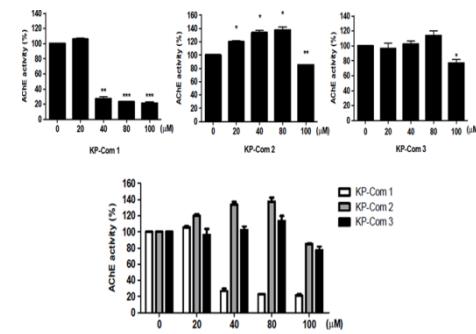


Figure 1: Acetylcholinesterase (AChE) activity in the presence of KP1, KP2 and KP3.

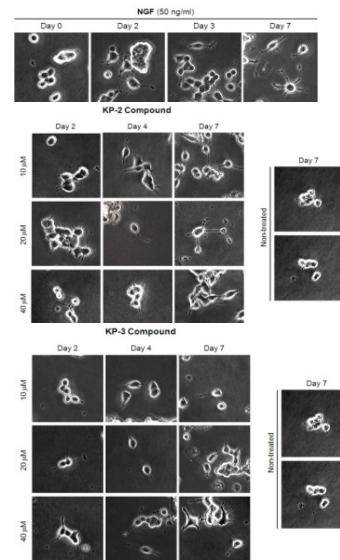


Figure 2: Neurite outgrowth of PC12 cells in the presence of KP2 and KP3. NGF, nerve growth factor.

of 100 μ M (Figure 1). In fact, the AChE activity increased to 140% in the presence of 40 and 80 μ M of KP2 and KP3, confirming that these two compounds did not have high AChE-inhibitory activity.

Although the AChE activity was increased in the presence of 20 μ M KP1, the ginger plant compound at 40, 80, and 100 μ M concentrations significantly reduced the enzyme activity. It was previously reported that K2 and K3 had high AChE-inhibitory activity, although much lower than that of K1. PC12 cells were treated with the three compounds at the same concentrations to compare their effects on the changes in cell morphology. The cytotoxicity of the compounds in order of strength was KP1 > KP2 > KP3. In the case of KP1, serious cell damage occurred at 40 μ M, whereas at the same concentration, KP2 caused only slight cell damage and KP3 did not cause any cell damage at all. PC12 cells were treated with the three compounds dissolved in DMSO to confirm their effects on neurite outgrowth. Almost no neurite outgrowth was observed in the presence of KP1, which was likely an influence of the cell damage caused by the cytotoxicity of this compound. In the cases of KP2 and KP3, neurite outgrowth was similar to or more than that observed in the presence of nerve growth factor (Figure 2), indicating that KP2 and KP3 have the ability to induce neurite outgrowth in PC12 cells. The isolated compounds were identified as 3,5,7,3',4'-pentamethoxyflavone (KP1), 5,7-dimethoxyflavone (KP2), and 5,7,4'-trimethoxyflavone (KP3). KP2 and KP3 significantly inhibited AChE activity in a dose-dependent manner.

Experimental

General experimental procedures: 1 H NMR spectra were recorded with an Agilent MR 400 DD2 400MHz spectrometer. Column chromatography (CC) was conducted using silica gel 60 (40-63 and 63-200 μ m particle size) and RP-18 (40-63 μ m particle size), both obtained from Merck III.

Plant material: *Kaempferia parviflora* rhizomes were kindly provided by a local farmer from Jang heung, Jeon Nam, South Korea. The rhizomes were identified by Prof. Hyung-In Moon. A voucher specimen (No. 2012-0405) has been deposited in the Herbarium of the Dong-A University (Busan, South Korea).

Extraction and fractionation: The rhizomes of *K. parviflora* (5 kg) were sliced and extracted under reflux with 9 L of MeOH, 3 times at 5°C. The MeOH extracts (198 g) were combined and concentrated *in vacuo* at 40°C. The MeOH extract was diluted with distilled water (0.9 L) and then partitioned with CHCl₃ (0.6 L×3). The CHCl₃-soluble fraction was diluted with distilled water and partitioned with *n*-hexane (0.9 L×3), CH₂Cl₂ (0.7 L×3), EtOAc (0.8 L×3) and *n*-BuOH (0.8 L). The CH₂Cl₂ extract (49 g) was separated by silica gel CC by gradient elution with CHCl₃ : MeOH (from 30:1 to 0:1 v/v); 12 fractions were obtained (K1-K12). Fractions K1 to

K8 were further separated by CC using CHCl₃ : MeOH (30:1, v/v). All fractions were recrystallized at -20°C from MeOH; KP1 (5.6 g) was obtained from fraction K5, KP2 (3g) from fraction K6 and KP3 (2.7g) from fraction K4.

Acetylcholinesterase inhibitory activity assay: PC-12 cells were obtained from American Type Culture Collection (ATCC; USA). Cells were cultured in a collagen coated culture dish in Roswell Park Memorial Institute 1640 (RPMI1640; GIBCO, Korea) medium supplemented with 10%, v/v, heat-inactivated fetal bovine serum (FBS; WelGENE, Korea), 5%, v/v, heat-inactivated horse serum (HS; WelGENE, Korea), 100 Units/mL penicillin, and 100 mg/mL streptomycin at 37°C in a humidified atmosphere of 5% CO₂. For differentiation, the medium was replaced with RPMI1640 supplemented with 2% FBS and 1% HS. PC-12 cells were treated with 50 ng/mL nerve growth factor (NGF; Sigma) for 7 days. For microscopic observation at 0, 2, 4, and 7 days of neurite outgrowth, PC-12 cells were treated with 10, 20, 40 μ M of KP 1-3. Microscopic observations were made after 2, 4, and 7 days of neurite outgrowth.

KP 1-3 were measured for acetylcholinesterase inhibitory activity using the chromogenic method, with slight modification. Acetylcholinesterase from human erythrocytes (10 μ g/mL, 860 U/mg, Sigma) was dissolved in 100 mM phosphate buffer (PB, pH 7.0) containing dithiobis (2-nitrobenzoic acid) (DTNB, Sigma), 10 mM (3.96 mg/ mL). Acetylthiocholine iodide, 75 mM (21.67 mg/mL) in the same buffer (pH 7.0), was used as a substrate solution. The enzyme solution (90 μ L) and each concentration (0, 20, 40, 80, 100 μ M) of the KP compounds (18 μ L) were mixed in a well of a microtiter plate and measured (415 nm) at time zero with a microplate reader. After incubation for 15 min at room temperature the substrate solution (90 μ L) was added and the reaction left for 10 min at 37°C. The increase in absorbance from zero time was measured.

Statistical analysis: The results were expressed as the mean \pm SE. Each value represents the mean of at least 3 independent experiments in each group. The statistical significance of the difference between two cell populations was determined using the two-tailed Student's t-test (Origin software; OriginLab). P values equal to or less than 0.05 were considered significant. A difference was considered to be significant at $P<0.05$.

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Antimutagenic Effects of Polymethoxy Flavonoids of *Citrus unshiu*

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Citrus fruits have been used as edible fruit and traditional medicine for various diseases such as cancer. In the course of our study to find antimutagens, we have found that the ethanolic extract of the peel of *Citrus unshiu* Marc showed antimutagenic effects against several mutagens in the Ames test using *Salmonella typhimurium* TA98 strain. Three polymethoxy flavonoids, nobiletin, 3,5,6,7,8,3',4'-heptamethoxyflavone, and tangeretin, were identified in the extract as major constituents. These three polymethoxy flavonoids showed antimutagenic effects in the Ames test *in vitro* and in the micronucleus test *in vivo*.

Keywords: *Citrus unshiu*, Polymethoxy flavonoids, Antimutagenic effect.

Citrus unshiu Marc., known as cold hardy mandarin, satsuma mandarin, satsuma orange, Christmas orange, and tangerine [1a], is cultivated in the southern regions of Japan and China. In addition, its peel has been traditionally used in East Asia as a drug for the treatment of vomiting and dyspepsia. Reports about the peel show anti-inflammation [1b], hepatoprotective, neuroprotective [2a], and anti-microbial activities [2b].

Polymethoxy flavonoids such as nobiletin (**1**), 3,5,6,7,8,3',4'-heptamethoxyflavone (HMF, **2**), and tangeretin (**3**) are known as the major constituents of *Kaempferia parviflora* and *Citrus* plants. Recent study showed that nobiletin (**1**) has chemopreventive effects against colon carcinogenesis, partly through regulation of leptin levels [3a]. The antimutagenic effects of nobiletin (**1**) were reported against the mutagens 3-amino-1,4-dimethyl-5H-pyrido[4,3-*b*]indole (Trp-P-1) [3b], 3,4-dimethyl-3H-imidazo[4,5-*f*]quinolin-2-amine (MeIQ) [4a] and benzo[*a*]pyrene (BaP) [4b] *in vitro* and against 4-(methyl-nitrosamino)-1-(3-pyridyl)-1-butanone (NNK) *in vivo*.

In the course of our study to find antimutagens, we found antimutagenic effects of the EtOH extract of *C. unshiu* in the *Salmonella typhimurium* mutagenicity assay against several mutagens. This short-term assay, namely the Ames test, has been extensively used to survey a great variety of environmental substances for mutagenic activity. In this study we evaluated the antimutagenic effects of the ethanolic extract of *C. unshiu*, nobiletin (**1**), HMF (**2**), and tangeretin (**3**) in the Ames test *in vitro* and the micronucleus test *in vivo*.

The EtOH extract of *C. unshiu* was partitioned into an EtOAc-MeOH-H₂O (13:3:16, v/v/v) mixture to give an organic layer and an aqueous layer. The antimutagenic effects of the EtOH extract, the organic layer, and H₂O layer against 1-nitropyrene (1-NP), 2-amino-1-methyl-6-phenylimidazo[4,5-*b*]pyridine (PhIP), Trp-P-1, 3-amino-1-methyl-5H-pyrido[4,3-*b*]indole (Trp-P-2), and BaP, were evaluated by the Ames test using *S. typhimurium* TA98 strain. 1-NP and BaP are representative environmental mutagens/carcinogens produced by internal combustion engines, such as diesel engines and emitted into the air. PhIP, Trp-P-1, and Trp-P-2 are mutagenic/carcinogenic heterocyclic amines formed by cooking meat. The EtOH extract showed antimutagenic effects against 1-NP, PhIP, Trp-P-1, and Trp-P-2 (Table 1). The organic

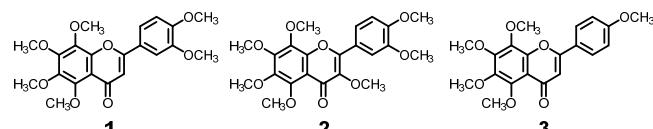


Figure 1: Structures of polymethoxy flavonoids **1–3**.

layer showed antimutagenic effects at lower concentrations [IC_{50} against 1-NP: 0.32 mg/plate, against PhIP: 0.02 mg/plate, against Trp-P-1: 0.02 mg/plate, against Trp-P-2: 0.08 mg/plate] than that of the H₂O layer [IC_{50} against PhIP: 0.21 mg/plate, against Trp-P-1: 0.42 mg/plate, against Trp-P-2: 1.97 mg/plate]. This result suggests that the active constituents were concentrated in the organic layer. On the other hand, the extract did not show an antimutagenic effect against BaP at 0.5 mg/plate.

Because polymethoxy flavonoids are reported to be major constituents of Citrus plants, we analyzed the EtOH extract of peels of *C. unshiu* by UV-HPLC to examine whether polymethoxy flavonoids existed in the extract. Peaks of polymethoxy flavonoids **1–3** were identified in the extract by comparing retention times and UV spectra. The amounts of **1–3** in 1 g of EtOH extract from peels of *C. unshiu* were as follows: nobiletin (**1**), 2.64 mg; HMF (**2**), 7.90 mg; and tangeretin (**3**), 2.42 mg.

To clarify the antimutagenic effects of nobiletin (**1**), HMF (**2**), and tangeretin (**3**), their antimutagenic activities against 1-NP, PhIP, Trp-P-1, and Trp-P-2 were examined by the Ames test using *S. typhimurium* TA98 strain. Compounds **1–3** showed antimutagenic effects against PhIP, Trp-P-1, and Trp-P-2 (**1** [IC_{50} against PhIP: 15.1 nmol/plate; against Trp-P-1: 37.0 nmol/plate, against Trp-P-2: 63.2 nmol/plate], **2** [IC_{50} against PhIP: 12.7 nmol/plate; against Trp-P-1: 22.4 nmol/plate, against Trp-P-2: 92.3 nmol/plate], and **3** [IC_{50} against PhIP: 2.9 nmol/plate; against Trp-P-1: 2.6 mol/plate, against Trp-P-2: 28.8 nmol/plate]) (Table 2). Among the tested samples, tangeretin (**3**) showed antimutagenic effects at the lowest concentration against PhIP, Trp-P-1, and Trp-P-2. On the other hand, **1–3** did not show significant antimutagenic effects against 1-NP and BaP. The IC_{50} value of **1** against Trp-P-1 in this report was identical to that reported [IC_{50} : between 30 and 60 nmol/plate] [3b].

To examine antimutagenic effects of extract of *C. unshiu*, and polymethoxy flavonoids **1–3** *in vivo*, we conducted a micronucleus

Table 1: Antimutagenic effects of EtOH extract, organic layer, and H₂O layer from *C. unshiu*.

Mutagen	With/Without S9 mix	Sample	Dose (mg/plate)	Revertants (number/plates)	Inhibition (%)
1-NP	without	<i>C. unshiu</i> ext.	0	1597	0
			0.1	1435	10
			0.2	1224	23
			0.4	791	51
			0.8	552	65
	Organic layer		0	1597	0
			0.05	1387	13
			0.1	1185	26
			0.2	985	38
			0.4	682	57
	H ₂ O layer		0	1597	0
			0.5	1495	6
			1	1458	9
			2	1466	8
			4	1206	25
PhIP	with	<i>C. unshiu</i> ext.	0	1115	0
			0.025	701	37
			0.05	512	54
			0.1	296	74
			0.2	220	80
	Organic layer		0	1031	0
			0.0125	564	45
			0.025	336	67
			0.05	261	75
			0.1	176	83
	H ₂ O layer		0	1031	0
			0.125	772	25
			0.25	415	60
			0.5	340	67
			1	241	77
Tr-P-1	with	<i>C. unshiu</i> extract	0	1198	0
			0.025	820	32
			0.05	586	51
			0.1	378	69
			0.2	234	81
	Organic layer		0	945	0
			0.0125	528	44
			0.025	433	54
			0.05	295	69
			0.1	225	76
	H ₂ O layer		0	945	0
			0.125	882	7
			0.25	556	41
			0.5	432	54
			1	286	70
Trp-P-2	with	<i>C. unshiu</i> ext.	0	1235	0
			0.125	1043	16
			0.25	748	40
			0.5	558	55
			1	344	72
	Organic layer		0	2075	0
			0.05	1300	37
			0.1	915	56
			0.2	663	68
			0.4	607	71
	H ₂ O layer		0	2075	0
			0.5	1656	20
			1	1389	33
			2	1026	51
			4	695	67
BaP	with	<i>C. unshiu</i> ext.	0	495	0
			0.0625	469	5
			0.125	529	ND
			0.25	495	0
			0.5	374	24
	Organic layer		0	495	0
			0.05	473	5
			0.1	508	ND
			0.2	493	0
			0.4	375	24
	H ₂ O layer		0	495	0
			0.5	446	10
			1	423	15
			2	460	7
			4	514	ND

test using peripheral blood of male ICR mice. The micronucleus test is a test to detect chromosomal damage induced by genotoxic/carcinogenic compounds, and has been used to evaluate the antimutagenic agents *in vivo* [3b]. In this study, we evaluate the *in vivo* antimutagenic effects of the EtOH extract of *C. unshiu* and

Table 2: Antimutagenic effects of 1–3.

Mutagen	With/Without S9 mix	Sample	Dose (nmol/plate)	Revertants (number/plates)	Inhibition (%)
1-NP	without	Nobiletin	0	1191	0
			5	940	21
			10	956	20
			20	957	20
			40	968	19
	HMF		0	1191	0
			100	1092	8
			200	944	21
			400	976	18
			800	1309	ND
PhIP	with	Tangeretin	0	1191	0
			50	985	17
			100	950	20
			200	926	22
			400	944	21
	Nobiletin		0	818	0
			2.5	730	11
			5	537	35
			10	460	44
			20	360	56
Trp-P-1	with	HMF	0	1031	0
			1.25	914	11
			2.5	984	5
			5	867	16
			10	549	47
			20	427	59
	Tangeretin		0	786	0
			0.47	678	14
			0.9	623	21
			1.86	468	40
			3.72	295	62
Trp-P-2	with	Nobiletin	0	973	0
			10	934	4
			20	596	39
			40	468	52
			80	426	56
	HMF		0	1543	0
			6.25	1317	15
			12.5	998	35
			25	712	54
			50	580	62
Tangeretin	with	Tangeretin	0	1179	0
			0.625	879	25
			1.25	800	32
			2.5	593	50
			5	441	63
	Nobiletin		0	1086	0
			20	692	36
			40	649	40
			80	467	57
			160	414	62
BaP	with	HMF	0	2075	0
			12.5	1948	6
			25	1718	17
			50	1593	23
			100	937	55
	Tangeretin		0	1853	0
			6.25	1522	18
			12.5	1214	35
			25	966	48
			50	709	62

polymethoxy flavonoids against PhIP and mitomycin C (MMC). We gave normal feed or sample feed that included the EtOH extract or polymethoxy flavonoids 1–3 at low or high dose (5% or 10% for the EtOH extract, and 0.01% or 0.02% for polymethoxy flavonoids). Mouse tail vein blood (5 µL) was taken prior to administration of either PhIP or MMC and after the administration (24, 48 and 72 hours).

In the micronucleus test, the EtOH extract of *C. unshiu* significantly decreased the frequency of micronucleated reticulocytes (MNRTs) treated with PhIP and MMC at 24 and 48 hours after the administration (Figures 2 and 3). The antimutagenic effects of the extract of *C. unshiu* were dose-dependently increased at 5% and 10%. The foods containing 0.02% nobiletin (1) produced inhibition significantly at 24 h and 48 hours after administration (Figure 4). The feed containing HMF (2) at 0.01 and 0.02% showed significant

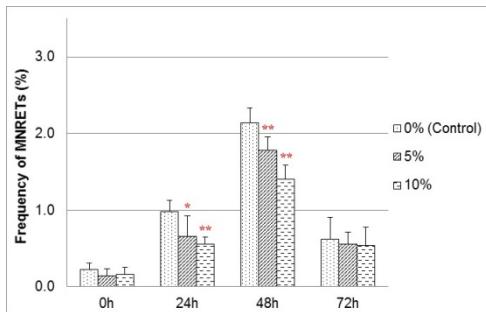


Figure 2: Frequency of micronucleated reticulocytes (MNRETs) from peripheral blood of mice treated with PhIP (50 mg/kg bw) and EtOH extract of *C. unshiu*. Each point represents the mean and standard deviation of five mice. Significant difference: * $P<0.05$; ** $P<0.01$ (Student's t test).

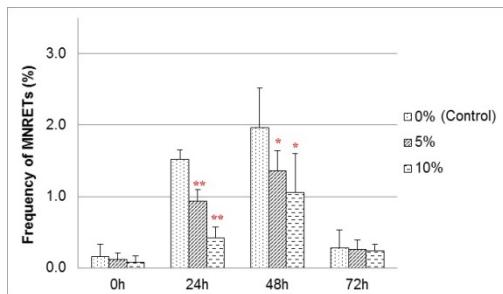


Figure 3: Frequency of MNRETs from peripheral blood of mice treated with MMC (1 mg/kg bw) and EtOH extract of *C. unshiu*. Each point represents the mean and standard deviation of five mice. Significant difference: * $P<0.05$; ** $P<0.01$ (Student's t test).

inhibition at 24 and 48 hours after administration (Figure 5), whereas the feed containing tangeretin (3) at 0.01 and 0.02% showed significant inhibition at 48 hours after administration (Figure 6). Polymethoxy flavonoids are known inhibitors of cytochrome P450 CYP1 enzymes [5]. Among the cytochrome P450s, CYP1A2 is mainly involved in the bioactivation of heterocyclic amines such as PhIP and Trp-P-1 [6]. Therefore, antimutagenic effects of polymethoxy flavonoids against heterocyclic amines found *in vitro* and *in vivo* tests in this research were mainly induced by inhibitory effects of CYP1A2. On the other hand, MMC has been recognized as a classical DNA damaging agent, on account of its monofunctional and bifunctional DNA alkylating activity [7]. The antimutagenic effects of the EtOH extract of *C. unshiu* against MMC in the micronucleus test may be caused by inhibition of the alkylating activity.

In conclusion, the antimutagenic effects of *C. unshiu* extract were evaluated by the Ames test *in vitro*. The inhibitory effects of polymethoxy flavonoids against chromosomal damage induced by genotoxic/carcinogenic compounds were also evaluated by the micronucleus test *in vivo*. We identified the polymethoxy flavonoids 1–3 in the extract of peels of *C. unshiu*. These compounds also showed antimutagenic effects *in vitro* and *in vivo*. A previous report described nobiletin (2) as having an inhibitory effect on PhIP-induced colon carcinogenesis in F344 rats [8a]. In this paper, we report for the first time that the polymethoxy flavonoids 1–3 show antimutagenic effects against PhIP in an *in vivo* micronucleus test. These results suggest that the peels of *C. unshiu* and polymethoxy flavonoids have a potential to prevent cancer produced by several mutagens.

Experimental

General: For HPLC, a Shimadzu SPD-M10Ayp was used with UV-VIS detectors (Kyoto, Japan). Nobiletin and HMF were purchased

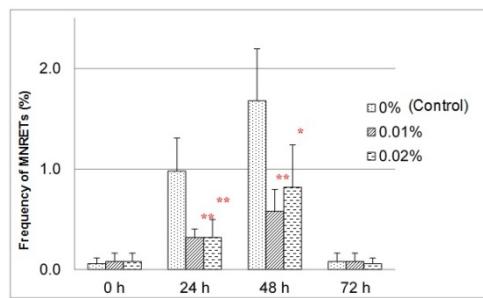


Figure 4: Frequency of MNRETs from peripheral blood of mice treated with PhIP (50 mg/kg bw) and nobiletin (1). Each point represents the mean and standard deviation of five mice. Significant difference: ** $P<0.01$ (Student's t test).

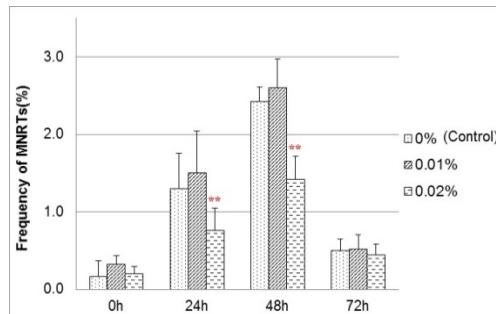


Figure 5: Frequency of MNRETs from peripheral blood of mice treated with PhIP (50 mg/kg bw) and HMF (2). Each point represents the mean and standard deviation of five mice. Significant difference: * $P<0.05$; ** $P<0.01$ (Student's t test).

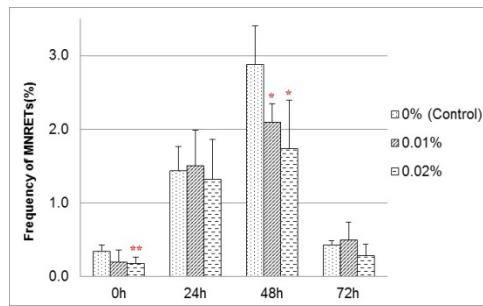


Figure 6: Frequency of MNRETs from peripheral blood of mice treated with PhIP (50 mg/kg bw) and tangeretin (3). Each point represents the mean and standard deviation of five mice. Significant difference: * $P<0.05$; ** $P<0.01$ (Student's t test).

from Wako Pure Chemical Industries (Osaka, Japan), and tangeretin from INDOFINE Chemical Company. Trp-P-1 (CAS no. 62450-06-0), Trp-P-2, PhIP, and BaP were purchased from Wako Pure Chemical Industries, 1-NP from Sigma-Aldrich (St. Louis, MO, USA), phenobarbital sodium salt from Tokyo Chemical Industry Co (Tokyo, Japan) and β -naphthoflavone from Nacalai Tesqu.

Animals: Male SD rats (7 weeks old, 200 g) and male ICR mice (7 weeks old, 32–34 g) were purchased from Japan SLC (Shizuoka, Japan). The experiments were conducted according to the “Guidelines for Animal Experiments in the Kyoto Pharmaceutical University”.

Extraction and partition: The EtOH extract from 1 g of *C. unshiu* peel was partitioned into a EtOAc-MeOH-H₂O (13:3:16, v/v/v) mixture to give organic (167.7 mg) and aqueous layers (766.9 mg).

Identification and quantitative analysis of polymethoxy flavonoids: Quantitative analysis was carried out by the absolute

calibration curve method. The separation was performed on a COSMOSIL 5C18-MS-II (250×4.6 mm) column (Nacalai Tesqu, Kyoto, Japan). The mobile phase consisted of 0.1% formic acid-water/acetonitrile (6:4, v/v) at a flow rate of 0.7 mL/min. The injection volume, column temperature and UV detection wavelength were set at 2 µL, 40°C and 210, 450 nm. Identification of each polymethoxy flavonoid in the test solution was compared with the retention time and UV-visible absorption spectrum of a standard.

In vitro antimutagenic assay: Mutagenicity was examined by the preincubation method [8b] using *Salmonella typhimurium* TA98 with and without S9 mix. The S9 mix (0.5 mL) contained 0.025 mL of S9. The S9 was prepared from the liver of male Sprague-Dawley rats treated with phenobarbital sodium salt and β-naphthoflavone. Dimethyl sulfoxide (DMSO) was used as a negative control. 1-NP (5 µg), PhIP (1 µg), Trp-P-1 (0.04 µg), Trp-P-2 (0.01 µg), and BaP (2.5 µg) were used as positive controls. The antimutagenic effect was examined of several doses of *C. unshiu* extract, and the organic and H₂O layers of the extract. Antimutagenicity was evaluated by the reduction of revertant colonies induced by a mutagen in the presence of samples. The IC₅₀ was calculated from the dose responses of the samples. Mutagenicity of 1-NP was examined without S9 mix and that of PhIP, Trp-P-1 and Trp-P-2 with it.

In vivo micronucleus assay: The micronucleus test was carried out according to the method published previously [9]. Micronucleus induction was examined with 5 ICR mice per dose. During the period of taming and experiment, the animals were provided with food (CE-2 pel-let diet; CLEA Japan, Inc., Tokyo, Japan) and tap water *ad libitum*. PhIP (50 mg/kg bw) and MMC (1 mg/kg bw) were administered as the positive control group. The feed containing a test sample was ingested from 2 days before the experiment. *C. unshiu* EtOH extract (5% or 10%, w/w) containing feed, nobiletin (0.01% or 0.02%, w/w) containing feed, tangeretin (0.01% or 0.02%, w/w) containing feed, and HMF (0.01% or 0.02%, w/w) containing feed were used as prepared feeds. As a positive control, CE-2 was provided instead of the feed containing a test sample. Mouse tail vein blood (5 µL) was taken prior to administration of a positive control and after the administration (24, 48 and 72 h). The observation, excitation filter - absorbing filter (Nikon fluorescence block B-2A, Nikon, Ltd.) epi-equipped with a fluorescence microscope [power supply HB-10101AF, OPTIPHOT-2 lens (400 times), Nikon, Ltd.] was used. One thousand reticulocytes were observed per slide, and those including micronuclei were counted. The data were analyzed for significance using Student's t-test. The significance levels of the test were 5 and 1%.

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Application of Mixture Analysis to Crude Materials from Natural Resources (V)^[1]: Discrimination of *Glycyrrhiza uralensis* and *G. glabra* by EI mass spectrometry

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The roots and stolons of some *Glycyrrhiza* species are used worldwide for traditional folk medicines and commercial pharmaceuticals. Phenolic constituents such as flavonoids and coumarins are medicinal and vary according to species. Therefore, species identification is important for quality analysis. In order to identify *Glycyrrhiza* species by chemical fingerprinting, methanol extracts of the root bark of *Glycyrrhiza uralensis* Fischer and *G. glabra* Linné were analyzed using EI-MS. Differences in kinds and quantity of components are reflected in complex EI-MS data and determining characteristic peaks for each species is straightforward. The characteristic peaks were determined statistically by volcano plot, a multivariate analysis method. EI-MS data of *G. uralensis* and *G. glabra* showed differential patterns, and the notable peaks in each pattern were identified. Peaks at *m/z* 153 and 221 are signature peaks of *G. uralensis*, and at *m/z* 173, 309, and 324 are those of *G. glabra*. In conclusion, we found species-specific patterns by EI-MS that distinguish *G. uralensis* and *G. glabra*. This method based on chemical constituent patterns can be applied to identify other *Glycyrrhiza* species and similar natural products.

Keywords: EI-MS, *Glycyrrhiza*, Volcano plot, Metabolomics.

Natural products such as medicinal plants, herbs, and crude drugs vary greatly in quality and composition due to their origin, place of production, and seasonal variation. Therefore, quality control is needed for each natural product.

Multivariate analysis has been applied to natural products. Metabolic profiling of plant extracts provides information about their chemical composition [2]. Many examples of general metabolic profiling have been developed. In general, LC-MS, GC-MS, and NMR are methods used for such analyses.

As part of a series of metabolomics studies, we have reported application of mixture analysis to crude materials from natural products [1a-d]. One of these applications is identification of *Glycyrrhiza* species [1a, d].

The roots and stolons of some *Glycyrrhiza* species are used worldwide in traditional folk medicines and commercial pharmaceuticals. Phenolic constituents such as flavonoids and coumarins are medicinal and vary according to species [3]. These medicinal properties include antimicrobial, anti-inflammatory, and antispasmodic activity [4-7]. Therefore, species identification by constituent pattern has potential for quality analysis.

In a previous study, the bark of the roots and stolons of *G. uralensis* Fischer and *G. glabra* Linné were analyzed using DART (Direct Analysis in Real Time)-MS. The characteristic peaks of each species were determined statistically by volcano plot. DART-MS provides $[M+H]^+$ cation adducts in positive mode, resulting in fewer cleavage reactions. The goal of this study was to demonstrate another methodology for chemical fingerprinting of *Glycyrrhiza*

species using electron ionization mass spectrometry (EI-MS). EI-MS is a popular mass spectrometric method that has been applied to discrimination of flavonoids, saponin aglycones, and others. As it provides many fragment peaks in addition to molecular ion peaks, spectral data become complex. The root bark of *G. uralensis* and *G. glabra* contains many compounds. The difference of kinds and quantity of these components are reflected in complex EI-MS data, making it straightforward to determine characteristic peaks of each species with high reproducibility. Once a spectral pattern model is constructed for each species, the chemical fingerprint can be widely shared.

Three kinds of *G. uralensis* (from China, Russia, and Mongolia) and two kinds of *G. glabra* (from Kazakhstan and Uzbekistan) were examined. A methanol extract of the root bark of each *Glycyrrhiza* sample was prepared. Each extract was dissolved in dimethyl sulfoxide (DMSO) at a concentration of 5 mg/mL, and then subjected to EI-MS analysis. Each extract was measured in triplicate.

In the range of *m/z* 50–494, the characteristic peaks of each species were determined statistically by volcano plot. The spectral data when ion contents were at a maximum in total ion chromatography was used for statistical analysis.

Figure 1 shows spectral data for a representative sample of each species. The other samples displayed very similar patterns. Each spectrum is shown in Supplementary data S1-S15. The spectral data for each species were dissimilar, and notable peaks of each pattern were identified. The highest abundance peak was used as standard, and the intensities of the other peaks were calculated relative to the

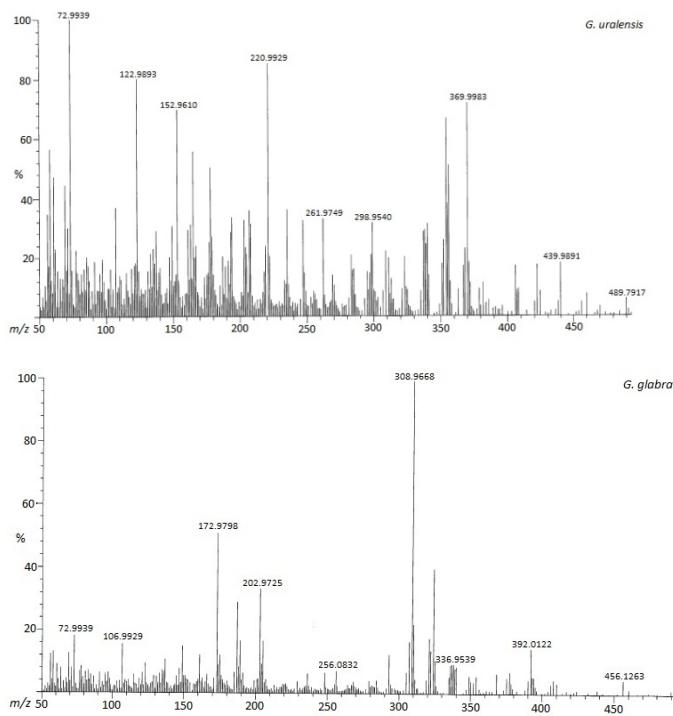


Figure 1: EI-MS data of methanol extracts of root and stolon bark of each *Glycyrrhiza* species.

standard by dividing by the standard deviation (SD) in the sample of the same individual ($SD = 1$).

The volcano plot summarizes the relationship between the p -values of a statistical test and the magnitude of the difference values of the samples in the groups. p -Values were calculated by Student's t -test. The difference values calculated as $[(\text{average of } G. \text{ glabra}) - (\text{average of } G. \text{ uralensis})]$ are plotted on the x-axis. The $-\log_{10} p$ values are plotted on the y-axis.

A volcano plot of the EI-MS data of the *Glycyrrhiza* species is shown in Figure 2. Peaks landing in the upper left region and the upper right region have a small p -value with a large absolute difference. In this case, peaks that had p values < 0.05 in the t -test and absolute differences of ≥ 4 were especially notable.

The volcano plot showed that peaks for m/z 153 and 221 are signature peaks of *G. uralensis*, and m/z 173, 309 and 324 are those of *G. glabra*. The two *Glycyrrhiza* species can be distinguished using these peaks in the EI-MS data (Figure 3).

Moreover, the relative intensities of the above particular peaks of the EI-MS data of the other root barks of *G. uralensis* and *G. glabra*, when applied to the constructed model, verified the process and corresponded with the preliminary identification (Figure 4).

Both species contain specific compounds, for example glycy coumarin for *G. uralensis* and glabridin for *G. glabra* [3].

Kusano et al. reported EI-MS data of glabridin; m/z 324 (M^+), 309, 187, and 173 [8]. From the view point of this, glabridin has a part role in the signature peaks of *G. glabra*. On the other hand, Demizu et al. reported MS data of glycy coumarin: m/z 368 (M^+) and 313 [5], which have little relevance to the above signature peaks of *G. uralensis*.

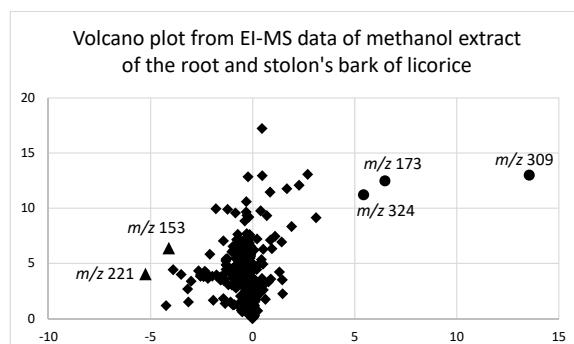


Figure 2: The volcano plot from the EI-MS data of methanol extracts of the root and stolon's bark of licorice. The x-axis is the difference in the average of both groups, and the y-axis is the computed p -value in $-\log_{10} p$ scale. Two peaks from the upper left region (characteristic for *G. uralensis*; represented as \blacktriangle) and three peaks from the upper right region (characteristic for *G. glabra*; represented as \bullet) were selected based on the following cutoff thresholds: absolute values of differences ≥ 4 and a p -value of 0.05.

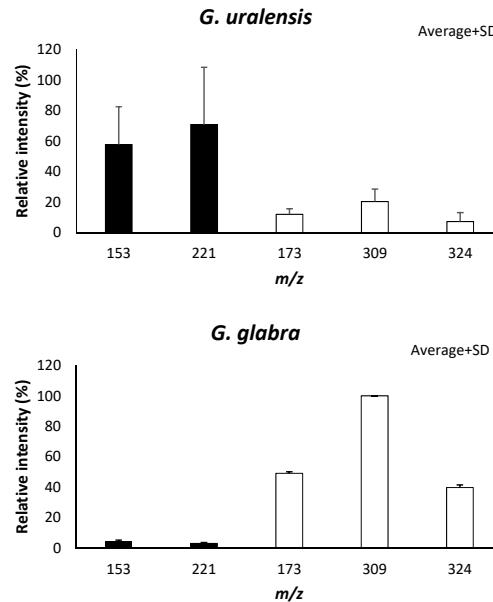


Figure 3: The average selected peak pattern of *G. uralensis* and *G. glabra*. The bars represent the original relative signal intensity. Black bars are signature peaks of *G. uralensis*. White bars are signature peaks of *G. glabra*.

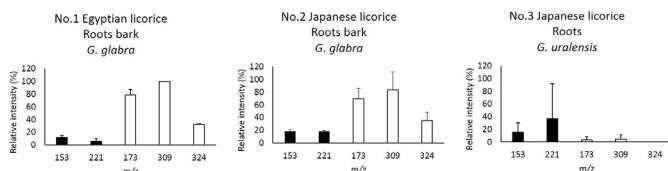


Figure 4: The average selected peak pattern of additional *G. glabra* (from Egypt and Japan) and *G. uralensis* (from Japan). The bars represent the original relative signal intensity. Black bars are signature peaks of *G. uralensis*. White bars are signature peaks of *G. glabra*.

The issue of identification of compounds generating species specific peaks is not irrelevant, but an important point is that the relative intensity of particular peaks in the EI-MS data led to the discrimination of the two species without identification of any index compounds.

In the same way, the search for potential markers using EI-MS data with multivariate analysis is very useful for quality evaluation of

natural resources, regardless of the existence of an index component.

In conclusion, we found species-specific patterns that distinguish *G. uralensis* and *G. glabra* by EI-MS. This method, based on chemical constituent patterns, can be applied to other *Glycyrrhiza* species and other morphologically similar natural products.

Experimental

Plant material: The roots and stolons of *G. uralensis* Fischer and *G. glabra* Linné were kindly supplied by Mr Fujio Kanai, Kanai Tokichi Shoten Co., Ltd. Voucher specimens were deposited at the Dept. of Natural Medicine and Phytochemistry, Meiji Pharmaceutical University, Tokyo, Japan (No.11G0001, No. 11G0002, No. 11G0003, No.11G0004, and No.11G0005, respectively).

The roots and stolons of additional *G. glabra* were purchased from Tachibana Japan (Tokyo) and identified by Dr Genjiro Kusano, Osaka University of Pharmaceutical Science, and Dr Tohru Okuyama, Meiji Pharmaceutical University. Voucher specimens were deposited at the Dept. of Natural Medicine and Phytochemistry, Meiji Pharmaceutical University, Tokyo, Japan (No. G0001).

The roots and stolons of additional *G. uralensis* (Hokkaido, Japan) and *G. glabra* (Hokkaido, Japan) were kindly supplied and identified by Dr Toshiro Shibata, Research Center for Medicinal Plant Resources in Nayoro, Japan. Voucher specimens were deposited at the Dept. of Natural Medicine and Phytochemistry, Meiji Pharmaceutical University, Tokyo, Japan (No.07G0003, No.07G0002, respectively).

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Species identification by HPLC: All samples were identified by HPLC analysis under similar conditions as those described by Shibano *et al.* [9]. A 5C18-AR-II (i.d. 4.6 x 150 mm, Nacalai Tesque Inc.) column was used. HPLC analysis was carried out on D-7000 type software; column oven: L-7300, Diode Array Detector; L-7450H, pump; L-7100 (HITACHI). The presence of species-specific substances was confirmed: glycyrrhetic acid for *G. uralensis* and glabridin for *G. glabra* [3].

Preparation of Glycyrrhiza extracts for EI-MS analysis: Each root bark specimen of 1 g *Glycyrrhiza* was extracted with 30 mL methanol for 1 h under reflux. The methanolic extract was evaporated under vacuum. After evaporation, DMSO was added to a final residue concentration of 5 mg/mL. One µL of each sample was analyzed using EI-MS.

Chemicals: All chemicals were analytical reagent grade. Glabridin was purchased from Wako Pure Chemical Industries Ltd. (Osaka, Japan), and glycyrrhetic acid was kindly provided by Dr Makio Shibano (Osaka University of Pharmaceutical Science, Japan).

EI mass spectrometry: All experiments were performed using a JMS-GCmate II (JEOL). Accelerating Volts, 2500 V; Mass Scan (Scan Range), *m/z* 50-494; Scan Speed, 0.3 sec/scan; Ion source, 70 eV.

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Comparison of the Qualitative Chemical Composition of Extracts from *Ageratina havanensis* Collected in Two Different Phenological Stages by FIA-ESI-IT-MSⁿ and UPLC/ESI-MSⁿ: Antiviral Activity

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The flowers and leaves of *Ageratina havanensis* (Kunth) R. M. King & H. Robinson are traditionally used as a tea to cure several diseases. The production of active secondary metabolites can be affected by several environmental factors such as climate, altitude, rainfall, phenological stage and other conditions that may influence the growth of plants. In this sense, the development of a methodology to compare the chemical composition of plant extracts is needed. The qualitative chemical composition of the ethyl acetate extracts of flowers and leaves, collected in both reproductive and non-reproductive season, was determined by flow injection analysis-electrospray ionization-ion trap tandem mass spectrometry (FIA-ESI-IT-MSⁿ) and ultra-high-performance liquid chromatography coupled to electrospray negative ionization mass spectrometry (UPLC/ESI-MSⁿ). The qualitative chemical composition of the ethyl acetate extracts of flowers and leaves was very similar in all cases. Also the antiviral activity of flowers against *human herpes simplex viruses* type 1 and 2 (HSV-1, HSV-2) (*Herpesviridae*) was analyzed. Three glucoside flavonoids were isolated from the ethyl acetate extract of the leaves of *A. havanensis* collected in flowering season using chromatographic methods and their structures were elucidated by physical and spectroscopic data measurements, and by comparing the obtained data with previously published values. The compounds were identified as 3-O-β-D-glucosyl-7-methoxyaromadendrin (5), 7-O-β-D-glucosyl-4'-dihydroxy-5-methoxyflavanone (6) and 5-O-β-D-glucosylsakuranetin (7); this is the first report of the isolation of these compounds in the Asteraceae family. Since the qualitative composition of the extracts of *A. havanensis* was similar in all cases, it can be expected that the ethyl acetate extract of the leaves collected in the non-reproductive season has anti-herpetic activity similar to that obtained in the reproductive season.

Keywords: *Ageratina havanensis*, Flavonoids, Tandem mass spectrometry, Anti-herpetic activity.

Ageratina havanensis (Kunth) R. M. King & H. Robinson (common name: Havana snakeroot) is a species of flowering shrub in the family Asteraceae, native of the Caribbean and Texas [1]; and used by humans populations as an antidiarrheal, antirheumatic, aromatic, cholagogue, febrifuge, tonic, and antiparasitic agent [2,3]. This plant is also used to cure diseases of kidneys and bladder [3].

In 1973, Dominguez and de la Fuente isolated sakuranetin and pucherryl acetate from *A. havanensis* grown in Texas [4]. Yu *et al.*, in 1987, isolated flavones, flavonols, dihydroflavonols, chalcones and glycosides from dichloromethane and ethyl acetate extracts [5].

In a previous work, our group reported for the first time the antiviral activity of extracts obtained from the leaves and stems of Cuban *A. havanensis* collected in the flowering season. The major metabolites from the most active extract (ethyl acetate extract of the leaves) were sakuranetin (1), 7-methoxyaromadendrin (2), 4'-O-β-D-glucosyl-7-methoxy-eriodictyol (3) and 4'-O-β-D-glucosyl-sakuranetin (4) [6]. However, there are no literature data concerning the chemical composition and pharmacological properties of the flowers and the plant in the non-reproductive season.

A number of environmental factors, such as climate, altitude, rainfall, phenological stage and other conditions may affect the growth of plants, which in turn affects the production of secondary metabolites present in the species, even when produced at one site

[7]. The purpose of the present work was to develop a relatively fast, precise and accurate method for an extensive profile of flavonoids in *A. havanensis* suitable for a comparative metabolic profiling of this class of phytochemicals allowing standardization of the plant material, and evaluating its seasonality.

Liquid chromatography-electrospray ionization-tandem mass spectrometry (LC/ESI-MSⁿ) is considered as a powerful tool for structural elucidation, providing structure specific data for the selective and sensitive identification of either known or unknown molecular structures [8]. Accordingly, an analytical method based on ultrahigh-performance liquid chromatography coupled to electrospray positive ionization mass spectrometry (UPLC/ESI-MS) and flow injection analysis-electrospray ionization-ion trap tandem mass spectrometry (FIA-ESI-IT-MSⁿ) was implemented for a quick identification of the flavonoids occurring in leaves and flowers of *A. havanensis*.

In order to obtain qualitative information about the chemical composition, ethyl acetate extracts (EtOAc) of the leaves of *A. havanensis* collected in both reproductive (flowering) and non-reproductive seasons, were directly injected into the ESI source of an ion trap spectrometer. Figure 1 shows the ESI-IT full scan mass spectra indicating the [M-H]⁻ ions of major flavonoids present in the EtOAc extracts of leaves.

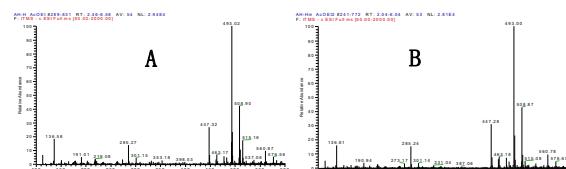


Figure 1: Flow injection analysis ESI-IT-MS fingerprints spectra (enlargement) obtained in negative ion mode of EtOAc extracts of leaves of *Ageratina havanensis* in reproductive season (A) and non-reproductive season (B).

The direct flow injection ESI-IT-MS fingerprint spectra obtained of the EtOAc extracts were very similar. The ion at m/z 285 was assigned to deprotonated sakuranetin (**1**), whose fragmentation (MS^2) led to the base peak at m/z 165 ($[M-120-H]^-$). This peak derives from the C-ring cleavage through a retro Diels-Alder (RDA) mechanism [9, 10]. The deprotonated molecule at m/z 301 suggested the presence of 7-methoxyaromadendrin (**2**). The product ion spectrum of m/z 301 (MS^2) showed the RDA ion at m/z 165 ($[M-136-H]^-$). Fragmentation of the deprotonated molecule at m/z 447 produced a product ion at m/z 285 ($[M-162-H]^-$, loss of a hexose moiety), which was tentatively assigned to the fragmentation of either $4'-O-\beta-D$ -glucosylsakuranetin (**4**) or another isomer of **4**. Similarly, the deprotonated molecule at m/z 463 suggested the presence of either $4'-O-\beta-D$ -glucosyl-7-methoxyeriodictyol (**3**) or a glycoside of 7-methoxyaromadendrin. The product ion spectrum of m/z 463 showed a $[M-162-H]^-$ ion at m/z 301, due to the elimination of a hexose moiety.

A Thermo UPLC C18 analytical column was chosen for the separation of the components present in extracts of the leaves of *A. havanensis* collected in reproductive and non-reproductive seasons. The major compounds were detected in the chromatograms of the EtOAc extract (flowering season) under the UPLC and tandem mass spectrometry conditions (Figure 2).

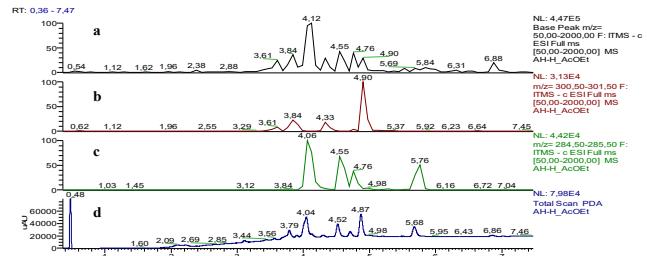


Figure 2: Chromatograms of EtOAc extract of leaves in flowering season. Total ion chromatogram (a), XIC relative to the ions at m/z 301 (b), 285 (c) and PDA total scan (d).

The peaks at 5.76 and 4.90 min were assigned to compounds **1** (sakuranetin, m/z 285 [$M-H]^-$]) and **2** (7-methoxyaromadendrin, m/z 301 [$M-H]^-$), respectively. Analysis of the mass spectral data of extracted ion chromatograms (XIC) at m/z 285 (Figure 2c) revealed the presence of three components eluting at 4.06, 4.55 and 4.76 min. The MS of these chromatographic peaks showed a predominant fragment ion at m/z 447, which was assigned to $4'-O-\beta-D$ -glucosylsakuranetin (**4**) and other isomers of **4**. The mass spectral data of XIC at m/z 301 showed three components eluting at 3.61, 3.84 and 4.33 min. According to the MS, the presence of compound **3** ($4'-O-\beta-D$ -glucosyl-7-methoxyeriodictyol) and another glycoside of either eriodictyol or 7-methoxyaromadendrin are suggested. The behavior of the EtOAc extract from leaves in the non-reproductive season (Figure 3) was very similar to the one of the extract in the flowering season, suggesting that the qualitative chemical composition of both extracts is the same.

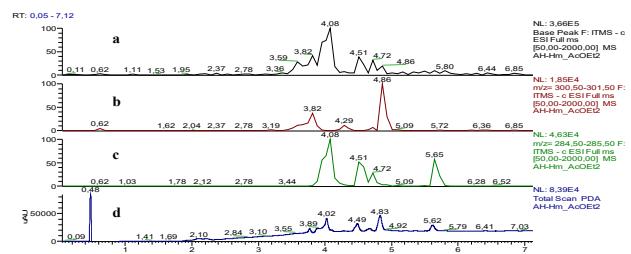


Figure 3: Chromatograms of EtOAc extract of leaves in non-reproductive season. Total ion chromatogram (a), XIC relative to the ions at m/z 301 (b), 285 (c) and PDA total scan (d).

To isolate and characterize the glycosides observed in the EtOAc extracts of the leaves of *A. havanensis*, a combination of column chromatographic techniques was used. The 1H - and ^{13}C -NMR spectroscopic data obtained for these compounds (not shown) are consistent with previously reported data. Compound **5** was identified as $3-O-\beta-D$ -glucosyl-7-methoxyaromadendrin. Compound **6** corresponds to $7-O-\beta-D$ -glucosyl-4'-dihydroxy-5-methoxyflavanone, isolated for the first time from *Prunus cerasoides* (Rosaceae) by Bahuguna *et al.* in 1987 [11]. Compound **7** data matched those for $5-O-\beta-D$ -glucosylsakuranetin. This flavonoid occurs commonly in the barks of plants belonging to the *Cerasus* subgenus of the Rosaceae family [12]. Nevertheless, this is the first report of compounds **5**, **6** and **7** in the Asteraceae.

Since compounds **6** and **7** were obtained by HPLC-UV with very close retention times, the peaks observed in the UPLC/ESI-MS² analysis at 4.55 and 4.76 min (figure 2) were assigned to compounds **6** and **7**, respectively. Therefore, the component eluting at 4.06 min corresponds to compound **4**.

FIA-ESI-IT-MSⁿ analysis of flowers was performed under the same conditions as those used for leaves. Compounds **1-7** were identified in the EtOAc extract, so it is concluded that the qualitative chemical composition of leaves and flowers is very similar.

For the determination of kaempferol and quercetin, the ethyl acetate extracts of leaves collected in both reproductive and non-reproductive seasons and flowers were analyzed using an UPLC system with a triple quadrupole tandem mass spectrometer equipped with an ESI source, and Probe Zspray operated in multiple reaction monitoring in negative ion electrospray mode. The chromatograms obtained showed the presence of quercetin (transition 301>151) in the extracts.

Man commonly uses leaf and flower infusions to heal several diseases. As far as is known, there is no report about the antiviral activity of the EtOAc extract of flowers of *A. havanensis*. Taking into account this fact, as well as considering that extracts from leaves and flowers have similar qualitative chemical composition, it was decided to evaluate the antiviral activity of the EtOAc extract of the flowers against *human herpes simplex viruses* types 1 and 2 (HSV-1, HSV-2) (*Herpesviridae*). The antiviral activity of the extract at non-cytotoxic concentrations was assessed by measuring its protective effects on infected Vero cells [13-16]. SI values greater than 2 were considered as an indication of antiviral activity [8]. The EtOAc extract of the flowers showed activity against *human herpes simplex viruses* types 1 and 2 (HSV-1, HSV-2) since in both cases the SI values were greater than 2 (Table 1). The anti-herpetic activity of this extract remains to be investigated, since the maximum concentration tested against HSV-2 did not reach the EC₅₀ value (SI< 6.21).

Table 1: Cytotoxicity and antiviral activity of EtOAc extract of *Ageratina havanensis* flowers.

Extract	Cytotoxicity		Antiviral Activity		
	CC ₅₀ ($\mu\text{g/mL}$)	CE ₅₀ ($\mu\text{g/mL}$)	HSV-1	CE ₅₀ ($\mu\text{g/mL}$)	HSV-2
AcOEt	1242.8 \pm 37.2	463.4 \pm 12.5	2.7	> 200	< 6.21

Based on the anti-herpetic activity observed in the extract and considering that the qualitative compositions of the extracts from the leaves and flowers are similar, it is suggested that the difference in the anti-herpetic activity is related to the quantitative composition of flavonoids previously identified, and that the presence of these flavonoids has a synergistic effect on the antiviral activity.

Experimental

General: Analytical-grade *n*-hexane, ethyl acetate, *n*-butanol and ethanol were used in this work. For FIA-ESI-IT-MSⁿ and UPLC/ESI-MS², HPLC-grade methanol was purchased from J.T. Baker (Phillipsburg, NJ, USA). HPLC-grade water was prepared using a Milli-Q purification system (Millipore, Bedford, MA, USA). The RP18 cartridge was a Phenomenex Strata C18-E, 55 μm , 70 Å, 500 mg · 3 mL⁻¹. The filter membrane (0.22 μm) was of nylon. Sephadex LH-20 and Silica gel 0.06-0.2 mm (70-230 Mesh, Merck) were used for column chromatography and TLC was performed on 0.2 mm-thick Kieselgel 60 F254 layers (Merck). The melting points (m.p.) were determined on a Reichert-Thermovar apparatus. IR spectra were recorded on a Mattson Genesis spectrophotometer, series FT-IR. ¹H- and ¹³C-NMR measurements were obtained on a Varian Inova 500 MHz NMR spectrometer with TMS as internal reference. HPLC-UV separations were performed on an Agilent 1100 series liquid chromatograph, equipped with a G-1312A binary pump, a G-1328 B rheodyne injector with a 0.2 mL sample loop, and a G-1365 B multiple wavelength detector.

Plant material: *Ageratina havanensis* (Kunth) R. M. King & H. Rob., Asteraceae (leaves and flowers), was collected in the eastern region of Havana (Alamar neighborhood) in November 2009 (flowering season) and March 2010 (non-reproductive season). Plant identification and collection were made by Prof. Iraly Ventosa from the Institute of Ecology and Systematics of Plants (Havana, Cuba), where a voucher specimen was deposited (Ref. HAC-42498). The fresh leaves and flowers were separated, oven-dried at 40 °C, and ground to fine powder.

Extraction: The powdered oven-dried plant sample (leaves from both seasons and flowers) was soaked in *n*-hexane and extracted with ethanol (EtOH). The EtOH extracts were concentrated under reduced pressure to a syrup, further dissolved in water and extracted with ethyl acetate (EtOAc) and *n*-butanol (*n*-BuOH) in a separation funnel. The resulting extracts were concentrated under reduced pressure to obtain the final EtOAc extract.

Sample preparation for mass spectrometric analysis: For mass spectrometric analysis, 2 mL of a MeOH/H₂O 8:2 (v/v) solution of each extract (1 mg mL⁻¹) was subjected to solid-phase extraction using a RP18 cartridge, eluted with MeOH/H₂O 8:2 (v/v). After drying, 1 mg was dissolved in 1 mL of MeOH/H₂O 8:2 (v/v) solution (solution A) and an aliquot (10 μL) was diluted with MeOH/H₂O 8:2 (v/v) up to a final volume of 1 mL, and was filtered through a 0.22 μm membrane of a nylon filter. The solution (150 μL) was diluted with MeOH/H₂O 8:2 (v/v) up to a final concentration of 1 ppm and then introduced by flow injection at 5 $\mu\text{L min}^{-1}$ directly into the ESI source using a syringe pump. For UPLC/ESI-MS analysis, an aliquot (50 μL) of solution A was

diluted with MeOH/H₂O 8:2 (v/v) up to a final concentration of 50 ppm.

Mass spectrometric analysis

Flow injection analysis (FIA): Analyses were performed using a Thermo Finnigan LCQ Deca ion trap mass spectrometer (San Jose, CA, USA) equipped with an ESI interface. Mass spectra were obtained both in positive and negative mode. The data were obtained at full scan (range of *m/z* 50–2000) and tandem mass scanning modes. For MSⁿ analyses, collision energies chosen for each fragmentation was 35%. The optimized instrumental parameters were: capillary temperature 300°C, capillary voltage 13 V, spray voltage 5 kV, sheath gas flow rate 35 (nitrogen, arbitrary units) and auxiliary gas flow rate 10 (arbitrary units).

UPLC-ESI-IT-MS analysis of the extracts: UPLC separations were performed on an Accela UHPLC System, Thermo Fisher Scientific including Accela PDA detector and Thermo Finnigan LCQ Deca ion trap mass spectrometer (San Jose, CA, USA) equipped with Xcalibur 2.2 software. The sample was injected onto a Thermo UPLC column C18 (2.1 x 50 mm x 1.7 μm) with a linear gradient of water (solvent A) and methanol (solvent B), both containing 0.1%, v/v, formic acid, from 0 to 100% B. Elution was performed at flow rate of 400 $\mu\text{L min}^{-1}$ and the volume of the injection was 20 μL . Detection by diode array was performed simultaneously at 3 different wavelengths: 250, 280 and 365 nm. The UV spectra were achieved within a 200–600 nm range. The mass analyses were performed with an ESI interface in negative mode. The data were obtained at full scan (range of *m/z* 50–2000) and tandem mass scanning modes. For tandem mass scanning mode, the percentage of collision energy range was 35%. The optimized instrumental parameters were: capillary temperature 300°C, capillary voltage 13 V, spray voltage 5 kV, sheath gas flow rate 35 (nitrogen, arbitrary units) and auxiliary gas flow rate 10 (arbitrary units).

UPLC-ESI-MS in multiple reaction monitoring mode analysis of the extracts: Analysis was performed on an ACQUITYTM UPLC system (Waters Corp., Milford, MA, USA) with a conditioned auto-sampler. The analytical column used was an ACQUITYTM UPLC Xbridge C18 (2.1 x 50 mm, 2.5 μm column; Waters Corp., Milford, MA, USA). Analysis was carried out with an elution gradient of water containing 0.1% formic acid (A) and methanol (B) at a flow rate of 400 $\mu\text{L/min}$ (10–100% B). The injection volume was 10 μL . Mass spectrometric detection was performed using a Waters ACQUITYTM Xevo TQD triple quadrupole tandem mass spectrometer (Waters Corp., Manchester, UK) equipped with an ESI source, Probe Zspray, operated in multiple reaction monitoring, negative ion electrospray mode. The monitored transitions included the following: sakuranetin (285>165), kaempherol (285>151), 7-methoxyaromadendrin (301>165) and querectin (301>151). The MS conditions were as follows: capillary temperature 200°C, capillary voltage 3.5 kV, cone voltage 39 kV, desolvation temperature 200°C, source gas flow: desolvation 400 L/h. All data were obtained and processed using MasslynxTM V4.1 software.

Fractionation by gel filtration and adsorption: The EtOAc extract obtained from the leaves of *Ageratina havanensis* collected in flowering season (1.0 g) was dissolved in a *n*-hexane/ chloroform/ methanol (2:1:1) mixture and separated using the same mixture as the mobile phase through a Sephadex LH-20 packed column to give 11 fractions (A-K). Compounds **1**, **2**, **3** and **4** were isolated from fractions I, J, E and G, respectively [6]. Fraction H (328.1mg) was separated on a silica gel packed column with *n*-hexane/ chloroform/ methanol (1:3:1) to give 6 fractions.

HPLC-UV analysis: Fractions 20-24 (silica gel column) from the leaves Sephadex fraction H (89 mg) were analyzed by HPLC-DAD on a C18 reversed-phase column (Phenomenex Luna (2), 250 mm x 4.6 mm i.d, 5 mm), using H₂O+TFA 0.1 % (solvent A) and MeOH+TFA 0.1 % (solvent B) as mobile phases, at a flow rate of 4.7 mL/min. Elution was performed in 100 min under an isocratic gradient (35% B). The chromatographic separation yielded 3 compounds, which were identified on the basis of IR and NMR experiments, as compounds **5** (4.7 mg, tR = 14.71 min, 206-207°C), **6** (20.9 mg, tR = 24.01 min, 200-202 °C), and **7** (29.0 mg, tR = 27.66 min, 200-202°C). Compounds **5**, **6** and **7** were monitored at 254 nm.

Flower antiviral activity

Cytotoxic assays: Vero cells (ATCC, CCL81) were seeded in 96-well plates at a density of 2×104 cells/ well, and incubated at 37°C in a 5% CO₂ atmosphere for 48-72 h, until 90% or greater confluence of the monolayers was reached. The cells were then incubated with increasing concentrations of the extracts at 37°C in a 5% CO₂ atmosphere for 72 h. Afterwards, a MTT solution (0.5 mg/mL) was added and the plates were further incubated for 4 h to allow formazan production. The solid purple precipitate was dissolved in dimethylsulfoxide (Sigma) and the absorbance at 570 nm was measured using a μQUANT Spectrophotometer (Bio-Tek Instruments) with a reference wavelength of 620 nm. The concentration reducing cell viability by 50% (cytotoxic mean concentration, CC₅₀) was calculated by regression analysis [17].

Antiviral assays: Ninety-six well plates containing confluent cell monolayers were pre-incubated for 1 h with increasing non-cytotoxic concentrations of the plant extracts, with a replicate number of 6. Subsequently, the cells were infected with HSV-1 and HSV-2 (10 TCID50). The plates were incubated at 37°C in a 5% CO₂ atmosphere and daily observed for viral-induced cytopathic effect (CPE) using a light microscope. When CPE was observed in all virus control wells, the percentage of wells with CPE was determined for each treatment concentration, as previously reported [16]. Acyclovir (ACV) at concentrations varying from 0.1-10 µg/mL served as positive control during HSV evaluations.

Statistics: Statistical analyses were performed using Statistic 6.1 software. Differences among antiviral activity of extracts were determined using Student unpaired t-tests. Values of *p*<0.05 were considered indicative of statistical differences. The concentration reducing cell viability by 50% (mean cytotoxic concentration, CC₅₀) and that reducing viral-induced CPE by 50% (antiviral effective mean concentration, EC₅₀) were calculated by regression analyses using the dose-response curves (not shown) generated from the experimental data. A selectivity index (SI) was calculated for each test simple by dividing its CC₅₀ by the corresponding EC₅₀ value.

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Characterization of the Polyphenolic Profiles of Peel, Flesh and Leaves of *Malus domestica* Cultivars Using UHPLC-DAD-HESI-MSⁿ

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Qualitative and quantitative analyses of polyphenols extracted from 21 *Malus domestica* cultivars using ultra high performance liquid chromatography with diode array detection coupled to heated electrospray ionization mass spectrometric detection was performed for separation of 27 phenolic compounds on a reversed phase UHPLC column with an optimized gradient consisting of 1% formic acid in water and 1% formic acid in methanol within 20 minutes. According to retention times, UV maxima and mass spectra of the peaks in the chromatograms obtained from extracts of apple peel, flesh and leaves, the polyphenolic compounds were identified and quantified. Based on fragmentation patterns, 6 phenolic acids, 5 flavan-3-ols, 5 dihydrochalcones, 8 flavonols and 3 flavone derivatives were characterized in the studied samples. The method was then employed for analysis of the polyphenolic pattern of 21 apple cultivars, both commercial and autochthonous for the Macedonian region, as well as for monitoring the influence of long term storage on the polyphenolic content and composition of apple fruits and for comparison of polyphenolic profiles of apple cultivars during two years of harvesting. The obtained results revealed minor differences in the quality and major variation in the content of phenolic compounds in the flesh, peel and leaves in the studied apple cultivars that is attributed mainly to cultivar differences and meteorological factors.

Keywords: *Malus domestica*, Polyphenols, HPLC-DAD-HESI-MSⁿ, Polyphenolic acids, Flavonoids, Procyandins, Dihydrochalcones.

Interest in the polyphenolic composition and content of different apple varieties and apple products originates from their confirmed contribution to human health [1–3]. It has been demonstrated that qualitative and quantitative differences in the phenolic content of apples can be attributed to many factors such as the variety, environmental conditions, maturity, storage conditions and time [4].

Many studies on polyphenols in apple fruits have been reported [5–11], some of them comparing the polyphenol content of peel and flesh, which is considered important since apples are consumed with and without peel. However, data on the polyphenolic profile of the leaves of apple trees are limited even though it has been shown that their composition might be used as a variety marker as well as an indicator of the health condition of the apple tree [12, 13].

Five major groups of polyphenolic compounds have been characterized in apple fruits: monomeric flavan-3-ols (catechins), polymeric flavan-3-ols (procyandins), dihydrochalcones (phloretin glycosides), flavonols (quercetin glycosides), hydroxycinnamic acid derivatives and anthocyanins (mainly cyanidin glycosides) present in the red varieties [8, 14]. Apple leaves contain mostly flavonols, few procyandins, hydroxycinnamic acids, and dihydrochalcones such as phloridzin and phloretin and their derivatives [12, 13]. Unlike from the flesh and the peel, apple leaves also contain flavones [13].

Although the major polyphenolic compounds found in apple varieties have already been identified [6–11], there are not many comprehensive studies on the polyphenolic profiles of large groups of *Malus domestica* varieties where peel, flesh and leaves are discussed. Knowledge on the precise polyphenolic profile of the apple cultivars may contribute to a better understanding of their role in the quality and diversity of apples as agricultural goods and their

role in agricultural products derived from them. Additionally, there are not much data on the specific apple varieties grown in southeast Europe, except for recent data for some Croatian [9] and Serbian cultivars [15].

Therefore, in this work, firstly, an efficient UHPLC-DAD-HESI-MSⁿ method for separation, identification and quantification within 20 minutes was developed and then employed on 21 apple cultivars (11 commercial, 8 autochthonous, and 2 domesticated) growing in the Republic of Macedonia. In that way, a comparative systematic study on the qualitative and quantitative composition of phenolic compounds in lyophilized flesh and peel, and air dried leaves from these regional and international *M. domestica* cultivars from two seasons (2014 and 2015) was carried out providing data on their polyphenolic composition for the first time. Additionally, the method was employed for monitoring the effect of storage on the polyphenolic content and composition of two commercial cultivars (Fuji and Granny Smith).

Twenty-seven phenolic components were determined by the proposed HPLC-DAD method by means of an external standard method. Before quantification, peak identities were confirmed by HPLC-ESI-MSⁿ (n = up to 3), using mass spectral and absorption data and comparison with standards and literature data. UV chromatograms of water/methanol extracts of peel, flesh and leaves of the cultivars and a mixture of standards were prepared; peaks were assigned based on retention, UV and mass spectral data given in Table 1, which also shows satisfactory chromatographic separation parameters.

Identification of polyphenolic compounds. The polyphenolic compounds detected in all studied samples were classified into five groups: phenolic acids, flavan-3-ols, flavonols, dihydrochalcones

and flavones. Sixteen of them were unambiguously identified by comparing retention times (t_R), UV and MS data with those of the available standards (peaks 1, 3, 5, 6, 8, 10, 11, 12, 16, 17, 22-27). Identification of another 11 polyphenolic compounds, for which standards were not available, was based on the UV absorption and MS data for the deprotonated molecules and their fragmentation pathways $[M-H]^-$ (data in Table 1). Whenever possible, these data and literature were used to support the identification of the peaks.

Phenolic acids: Six compounds were identified as phenolic acid derivatives. According to their UV and MS, four of them (5, 8, 9 and 11) were identified as hydroxycinnamic acid derivatives. Compounds 1, 5, 8 and 11 were confirmed using standards as gallic, chlorogenic, *p*-coumaric and caffeic acids, respectively. Mass spectra of compound 4 indicated fragments related to phenolic acids but in the first step of the fragmentation pattern, the loss of 180 amu implied the presence of one hexose moiety. In the literature it is stated that such phenolic acid derivatives are characteristic for cider apples and have structures with hydroxycinnamic acid moieties [16]. Compound 4 and compound 9 had deprotonated molecular ions $[M-H]^-$ at m/z 349 and 343, which fragmented in MS^2 to m/z 163, followed by a fragmentation pattern characteristic for gallic (comp. 4) and *p*-coumaric acids. So, they were tentatively identified as gallic and *p*-coumaric acid derivatives.

Flavan-3-ols: Compounds 2 and 7 showed the same deprotonated molecular ions and fragmentation pattern as compound 3, which was confirmed with standard as procyanidin B2. Following their elution order: compound 2 eluted before catechin and epicatechin, whereas compound 7 between catechin and epicatechin, and according to literature data for distinguishing procyanidins by photodiode array detection [17], compounds 2 and 7 were identified as procyanidin B3 and procyanidin B1, respectively.

Dihydrochalcones: Besides phloridzin (12) and phloretin (16), that were confirmed using standards, three other phloretin derivatives were also characterized in all apple cultivars (assigned as peaks 13, 14 and 15). According to their UV spectra, these compounds can be

described as phloretin derivatives, whereas the comparison of MS data with those found in literature [18-21] indicated the presence of phloretin glycosides. They were found in different concentrations in all analyzed extracts from peel, flesh and leaves. Compound 13 had a deprotonated molecular ion $[M-H]^-$ at m/z 583 and $[2M-H]^-$ at 1166. In the MS^2 spectra, ions at m/z 291 [$M+H_2O-pentosylhexoside]^-$ and 273 [phloretin] $^-$ were observed. Because there were no $[M+H-glucose]^-$ fragments it was concluded that the two sugars were not independently attached to the aglycone. The loss of deoxyhexose (-146) indicated that the glucose was directly attached to the aglycone. Thus, peak 13 was tentatively identified as phloretin rhamnosylglucoside (phloretin rutinoside). Compound 14 showed a deprotonated ion $[M-H]^-$ at m/z 451 and fragment ions at m/z 289 [$M+glucose-H]^-$ and 273 [phloretin]. So, it was tentatively identified as either 3-hydroxyphloretin glucoside or 3-hydroxyphloridzin. Compound 15 had a deprotonated ion $[M-H]^-$ at m/z 567, and $[2M-H]^-$ at 1135 and a fragment ion at m/z 273 [$M-H-pentosylhexose]^-$. It was tentatively identified as phloretin xyloglucoside. The last two compounds (14 and 15) have been identified in apple pomace as 3-hydroxyphloretin-2'-glucoside (3-hydroxyphloridzin) and phloretin-2'-xyloglucoside [22]. When their UV spectra were examined, it was found that the above three dihydrochalcones all showed UV maxima at 286 nm, but the hydroxyphloretin derivatives exhibited additional shoulders at 262 and 242 nm; also confirmed by Escarpa [5] and Tsao [7].

Flavonols: Eight flavonols were identified according to their UV absorption and MS data (Table 1). The $[Y_0]^-$ ions at m/z 317, 301 and 285 indicated three aglycones with the corresponding molecular ions: myricetin (27), quercetin (17-22) and kaempferol (23). Compounds 18, 22, 23 and 27 were determined as quercetin 3-*O*-rutinoside (rutin), quercetin, kaempferol and myricetin, respectively by comparison with standards. Compound 17 had a deprotonated molecular ion at m/z 463 and a fragment ion at m/z 301 indicating quercetin hexoside. The particular hexose moieties from these flavonol glycoside structures could not be confirmed based on MS information whereas, according to previous reports, glucose was the main hexose linked to flavonols in apples. Compounds 19, 20 and

Table 1: Retention times (t_R), UV maxima, $[M-H]^-$ and fragments, selectivity factor (α), asymmetry and resolution (Rs) of phenolic compounds found in the peel, flesh and leaf extracts of the analyzed 21 *Malus domestica* cultivars.

Peak No	t_R	λ_{max}	$[M-H]^-$	MS^{2a}	MS^{3a}	Compound	Sample	α	Asymmetry	Rs
1 ^b	0.43	272		169	125	gallic acid	peel, flesh and leaves	1.22	1.13	4.8
4	1.12	316.7		349	181;163; 125	gallic acid derivative	peel, flesh	1.23	1.22	2.11
5 ^b	1.47	235; 325.4		353	191	chlorogenic acid	peel	1.09	1.06	5.44
8 ^b	2.48	209; 310		163	119	<i>p</i> -coumaric acid	peel, leaves	2.00	1.16	3.85
9	3.11	209; 310		343	163; 119	<i>p</i> -coumaric acid derivative	peel	1.98	0.96	4.25
11 ^b	4.16	220.0; 325.3		179	135	caffeic acid	leaves	1.54	1.23	3.84
2	0.52	277.7		577	425 ; 407; 219	procyanidin B3	peel, flesh and leaves	1.02	1.15	2.15
3 ^b	0.92	277.7		577	425 ; 407; 219	procyanidin B2	peel, flesh and leaves	1.11	1.32	4.39
6 ^b	2.11	279.7		289	244 ; 230; 204	catechin	peel, flesh and leaves	1.07	1.15	3.32
7	2.31	278.1		577	425 ; 407; 219	procyanidin B1	peel, flesh and leaves	1.23	1.15	2.11
10 ^a	3.74	279.7		289	244; 230; 204	epicatechin	peel, flesh and leaves	1.52	1.31	10.88
12 ^b	4.76	225; 286		435	297; 273	phloridzin	peel, flesh and leaves	1.23	1.18	6.3
13	4.91	225; 242sh; 262 sh; 286		581	297; 273	phloretin rutinoside	peel, flesh and leaves	2.11	1.15	1.55
14	5.11	225; 242sh; 262sh; 286		451	289 ; 271	3-hydroxyphloridzin	peel, flesh and leaves	1.45	1.43	1.68
15	5.36	225; 286		567	297; 273 ; 167	phloretin pentosylhexoside	peel, flesh and leaves	1.76	1.28	1.98
16 ^b	6.51	225; 282.8; 369		273	167	phloretin	peel, flesh and leaves	1.32	0.92	20.77
17 ^b	7.92	255; 350		609	301; 179; 151	rutin	leaves	1.11	1.55	1.99
18	7.91	375; 268; 256		463	301 ; 285	isoquercetin	peel, flesh and leaves	1.09	1.16	13.39
19	8.11	255.3; 357.6		433	301 ; 283; 273; 257	quercetin pentoside	peel	1.31	1.58	2.01
20	8.54	255.3; 357.6		447	327; 285	kaempferol hexoside	peel	1.25	1.12	1.78
21	8.92	255.3; 357.6		417	285	kaempferol pentoside	peel	1.02	1.09	1.25
22 ^b	9.64	255.3; 357.6		301	301 ; 179	quercetin	peel, flesh and leaves	1.54	1.32	12.17
23 ^b	10.05	366; 266		285	255	kaempferol	leaves	1.00	1.17	2.71
27 ^b	13.01	353		317	179 ; 151	myricetin	leaves	1.27	1.54	12.07
24 ^b	10.5	255; 318; 350; 410		285	267; 257	luteolin ^b	leaves	1.23	0.98	3.36
25 ^b	12.2	275; 318; 401		431	269 ; 224	apigenin-7- <i>O</i> -glucoside ^b	leaves	1.66	1.23	14
26 ^b	12.4	275; 318; 401		269	224; 151	apigenin ^b	leaves	1.98	1.16	1.67

^aMost abundant ions are marked in bold. ^bCompounds confirmed with standards.

21 showed a deprotonated molecular ion at *m/z* 609 and fragment ions at *m/z* 301. These compounds had the same fragmentation pattern with different abundance of the MS² fragments, and considering their retention times, these three compounds eluting after rutin were tentatively attributed to other quercetin rutinoside isomers.

Flavones: Three flavone derivatives were found in the extracts of leaves samples. Their structures were confirmed using standards of luteolin (24), apigenin 7-*O*-glucoside (25) and apigenin (26).

Overall, catechin and epicatechin (flavanols), chlorogenic and caffeic acid (hydroxycinnamic acids), phloridzin (dihydrochalcones) and rutin (flavonols) were identified in both peel and flesh. Some of the varieties analyzed (Srcika (S12), Adam's Pearmain (Parmenka (S13)), Kojce (S14)) showed similar polyphenolic pattern to the ones found for cider apple cultivars reported by Mangas [23]. During sample preparation these cultivars were browning faster, had a bitter taste and showed high content of phenolic acids, especially *p*-coumaric acid.

The same groups of flavonoids present in apple peel and flesh were also found in the leaves, which also had flavones. Polyphenols in apple leaves (caffeic acid, isoquercetin, quercetin, kaempferol, apigenin-7-*O*-glucoside, apigenin, luteolin and myricetin) have been suggested as taxonomic markers for identification of *Malus* L. and *Pyrus* L. genera [12]. Accumulation of polyphenols has been shown as a post infection response to parasites such as scrubs [12, 13].

Quantification of polyphenolic compounds: The validity of the method was examined according to the FDA guidance recommendation [24]. Six standard solutions were prepared for each standard in the concentration range 0.1–200 µg/mL. All polyphenolic compounds showed good linearity ($r=0.9986\text{--}0.9999$) in the investigated concentration range. The limits of detection and quantification were calculated as amounts exhibiting three times the area of the largest noise peak measured with RSD<10% ($n=10$) and ten times the area of the largest noise peak measured with RSD<2% ($n=10$), respectively. Limits of detection ranged from 0.11 µg/mL for phloridzin to 0.59 µg/mL for rutin, and limit of quantification from 0.19 µg/mL for phloretin and phloridzin to 2.56 µg/mL for rutin. Standard deviations of 6 replicates of all analytes from the mixed standard solutions in 3 days were used for confirmation of the accuracy and precision of the method. Retention times showed deviation from 0.2 to 1% and peak area deviations were from 0.5%–2%. The accuracy of the method was tested by the standard additions method and recovery values in the range from 92–102% were found satisfactory. Method robustness was tested by varying column dimensions, flow and mobile phase and all tested parameters did not exceed changes in peak area and retention times higher than 25%. So, the method was found suitable for quantification of the phenolic compounds detected in the extracts of apple peel, flesh and leaves.

Overall, 27 compounds listed in Table 1 were detected in the fruits and leaves of 21 *M. domestica* cultivars. Gallic acid, *p*-coumaric acid, procyanidin B3, procyanidin B2, procyanidin B1, catechin, epicatechin, phloridzin, phloretin, rutin, quercetin and their derivatives were detected in both, apple fruit and apple leaves.

All results for the content of every measured compound in all studied cultivars from two seasons of apple harvesting are given in the supplementary tables 1S–6S. The total phenolic content of the 21 apple cultivars analyzed in the two years is given in the graphs in Figure 1, where the polyphenolic patterns are given as distributed in

the groups of phenolic acids, flavan-3-ols, dihydrochalcones, flavonols and flavones.

The total polyphenolic content values were in the range from 5.59 to 14.7 mg/g lyophilized peel, from 4.52 to 16.3 mg/g lyophilized flesh, and from 8.07 to 15.1 mg/g dried leaves. As can be seen in Figure 1, flavan-3-ols in the range of 1.95–6.09 mg/g lyophilized sample were most abundant in the apple peel followed by phenolic acids from 0.22 to 4.95 mg/g, and dihydrochalcones from 0.95–3.68 mg/g, whereas apple flesh contained slightly higher flavan-3-ol content than the peel (1.89–9.65 mg/g lyophilized sample) and the same concentration range of phenolic acids and dihydrochalcones as the peel (0.26–4.24 mg/g and 0.71–4.63 mg/g). Leaves also contained flavan-3-ols (2.49–6.52 mg/g dried sample) in highest proportion followed by phenolic acids and flavonols, but also flavones (1.45–2.43 mg/g) that were not detected in apple fruits. The results presented in Figure 1 suggest no significant seasonal variations in the content and distribution of the polyphenolic compounds in the leaves in the two different years (2014 and 2015). However, Figure 1 also suggests that total polyphenol content in the peel was higher, and in the flesh lower in the samples from 2014 compared with the corresponding samples from 2015. Data on the temperature for these years reveal that in 2014 the winter months were with higher temperatures and the temperatures from April to October (the vegetation period) were lower than in 2015, whereas overall 2015 had higher average temperature (Supplementary table 8S with data from the State hydrometeorological service, Skopje). This can be expressed also through the annual temperature sum, which was higher for 2014 (5438.5 vs 5292.5°C in 2015) and the vegetation temperature sum that was lower for 2014 (4215.8 vs 4429.8°C for 2015). As for the air humidity and rainfall, 2014 is characterized by significantly higher rainfall, especially in the vegetation period (543.2 compared with 283.4 mm in 2014). Our results would suggest a positive effect on flesh polyphenol content of higher temperature in the vegetation period and higher humidity, and the reverse effect causes higher polyphenol content in the peel. Still there are no established correlations between polyphenolic composition and climate conditions even though such studies are becoming more relevant due to the increased threat of global warming [25, 26]. Furthermore, Hagen *et al.* [27] have reported that the polyphenolic content, especially of the peel of *M. domestica*, is sensitive to UV irradiation; this was studied by artificially subjecting the fruits to postharvest irradiation. The results from the analysis of the polyphenols in leaf extracts in our study, compared with the ones for the fruit, revealed no seasonal variability.

Tsao *et al.* [7] measured total polyphenols in fresh apple fruits in the range 1.02–2.35 mg/g and Red Delicious contained the highest polyphenolic content in the peel compared with another seven studied apple cultivars popular in Ontario. Also, much lower content of polyphenols was measured in the extracts from the apple flesh (0.53 mg/g for Red Delicious flesh). The Macedonian Red Delicious (S2) cultivar studied in our work contained around 10–12 mg polyphenols/g lyophilized apple peel and around 6 mg polyphenols/g flesh, which is comparable having in mind the water content, but still the flesh polyphenolic content found is higher.

In comparison to our previous work [28], where polyphenols in peel and flesh were measured by spectrophotometric methods, the highest concentrations of polyphenols were found in the flesh of Tetovka (S21) 12.55 mg/g lyophilized sample, which is very similar to the result measured in this study (12.7 mg/g). Moreover, total polyphenols in the peel extracts are in good correlation with the maximal obtained concentrations for the peel of Tetovka, as well as the minimal concentrations measured in this work compared with

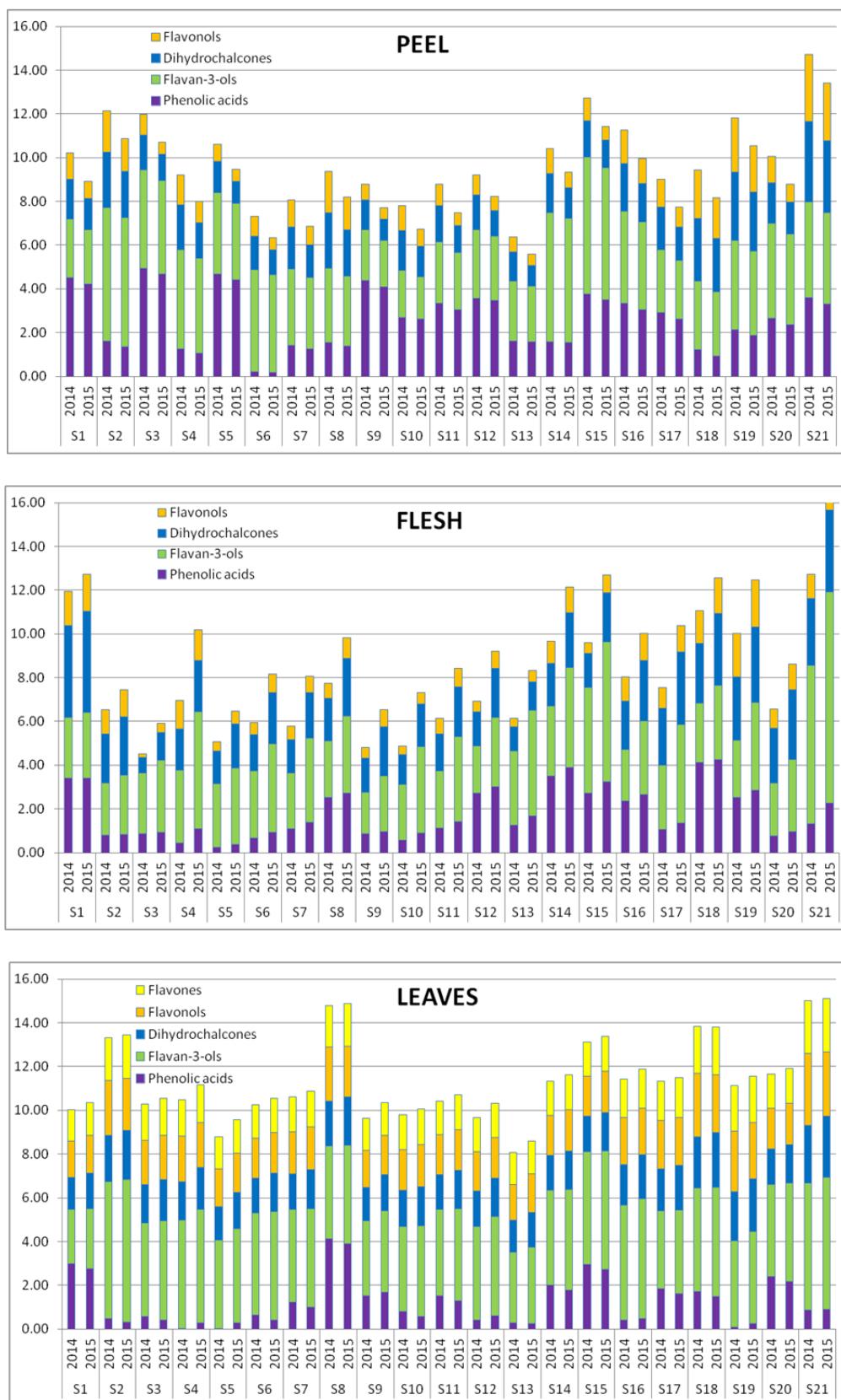


Figure 1: Polyphenolic content (grouped as phenolic acids, flavan-3-ols, dihydrochalcones, flavonols and flavones) in the 21 *Malus domestica* cultivars analyzed in two seasons (2014, 2015) in extracts of leaves, peel and flesh (expressed as mg/g lyophilized peel/flesh and mg/g dried leaves).

the previous work for the peel of Fuji (S10) (about 7 mg/g lyophilized sample). Although the total flavonoid concentration in the flesh was not identical to the results obtained with spectrophotometry, it correlated well to HPLC results with minor differences. Among the five major groups, procyanoindins predominated in both peel and flesh in the total polyphenolic profile. Kojce (S14), Livadarka (S15) and Tetovka (S21) had the highest total concentrations of procyanoindins in the peel and flesh whereas Golden Delicious (S1) had the lowest amounts, which can also cause differences between the test for total catechins and their separate quantification made here.

Similar results from the quantitative analysis of procyanoindins, catechin, epicatechin, caffeoic acid and dihydrochalcones for both flesh and peel measured in Golden and Red Delicious and Granny Smith were reported by Escarpa and Gonzales [5]. Analogous polyphenolic contents as the ones obtained in this work were also observed in the study of Basque cider apples where freeze dried samples were investigated [16]. Even though none of the apple cultivar analyzed by Mangas *et al.* was among the cultivars mentioned here, very similar results for the same detected polyphenolic compounds have been observed in this study [23].

Phenolic acids: The highest total amount of phenolic acids in the peel was 4.95 mg/g measured in 2014 and 4.67 mg/g measured in 2015 for the cultivar Chadel (S3), followed by Golden Delicious (S1), which contained about 44% phenolic acids. The highest percentage of gallic acid was found in Chadel (S3) from 2014 and 2015, 2.33 and 2.29 mg/g, respectively. The phenolic acid derivative with 3.25 mg/g in 2014 and 3.19 mg/g in samples from 2015 was found in Granny Smith (S5). *p*-Coumaric acid and its derivative were also found in Chadel (S3) (in 2014, 1.78 mg/g and in 2015 1.75 mg/g) and Golden Delicious (S1) (in 2014, 2.25 mg/g and in 2015, 2.10 mg/g). Minimal concentrations of polyphenolic acids were noticed in the peel of Idaret (S6), where 0.22 mg/g was measured in 2014 and 2015. The lowest concentration of 0.01 mg/g was found for peak 9 (*p*-coumaric acid derivative) found in the apple peel of the Kojce cultivar (S14). The gallic acid derivative (peak 4) was quantified as the highest measured phenolic acid derivative (2.44 mg/g) in the peel of Sreika (S12). Tsao *et al.* [7] found that about 9.3% of the total peel polyphenols were hydroxycinnamic acids. Our results suggest higher contents of total phenolic acids, up to 40%, due to the contribution of the hydroxybenzoic acids detected.

Maximal concentration of phenolic acids was found in the flesh of Pasalma (S18) for both 2014 and 2015 (4.11 and 4.24 per mg/g lyophilized sample). Slightly lower concentration was measured in the flesh of Golden delicious (S1), 3.42 mg/g in 2014 and 2015. Highest amounts of gallic acid and its derivative were found in the flesh of Pasalma cultivar (S18),

Caffeic acid was detected only in apple leaves in the range from 0.01-2.56 mg/g air dried sample with maximum quantity in the Mutsu cultivar (S8) and lowest in Braeburn (S4) and Granny Smith (S5) from 2014.

Flavan-3-ols: The five flavan-3-ols listed in Table 1 were detected as the major group of all analyzed compounds in peel, flesh and leaves. The highest content of flavan-3-ols in the peel was found in the Livadarka cultivar (S15) in 2014 (6.25 mg/g lyophilized sample), which was slightly higher than that measured in 2015 for the same cultivar (6.04 mg/g). Similar contents of flavan-3-ols were noticed in Red Delicious (S2), 6.09 and 5.88 mg/g, for 2014 and 2015 respectively. The peel from apples harvested in 2014 was

most abundant in epicatechin (Kojce (S14) with 2.20 mg/g). In our previous work [28], the flesh of Kojce was found to contain the highest amount of total catechins: 3.58 mg/g. The peel of Red Delicious (S2) was most abundant in procyanoindin B3 (1.85 mg/g), catechin (1.92 mg/g in 2014 and 1.81 mg/g in 2015), and procyanoindin B1 (1.22 mg/g). Livadarka (S15) contained the highest amounts of procyanoindin B2 (1.85 mg/g), and Kojce the highest amounts of epicatechin in the peel (2.20 mg/g, 2014 and 2.12 mg/g, 2015). These numbers were followed by the contents of flavan-3-ols in Gala (S7), 2.01 mg/g catechin and 0.98 mg/g procyanoindin B3. Fuji cultivar (S10) contained the lowest amounts of flavan-3-ols in the peel (2.16 mg/g in 2014 and 1.95 mg/g in 2015). Table 5S shows that Kojce is right after Red Delicious and Livadarka, containing about 60% flavan-3-ols. The lowest content of flavan-3-ols in the peel was measured in Golden Delicious (S1), 2.49 mg/g in 2014 and 2.76 mg/g in 2015, with the lowest measured content of procyanoindin B1 (0.01 mg/g in 2014 and 0.06 mg/g in 2015).

As for the flesh, the highest content of these compounds was found in the Tetovka cultivar (S21) with 7.27 mg/g for 2014 and 9.65 mg/g for 2015, followed by half of this content in Kojce (S15), 4.83 mg/g for 2014 and 6.40 mg/g for 2015. The lowest flavan-3-ol content was measured in the flesh of Fuji (S10), 2.16 mg/g in 2014 and 1.95 mg/g in 2015. Procyanoindin B3 was found to be the least abundant in Golden Delicious (S1) 0.02 and 0.05 mg/g for 2014 and 2015 and most abundant in the flesh of Tetovka (S21: 0.98 mg/g and 1.01 mg/g in 2014 and 2015). Similar to that, procyanoindin B2 was least abundant in Golden Delicious (0.02 and 0.09 mg/g in 2014 and 2015), but most abundant in Mislimka (S17) 0.74 mg/g and 1.11 mg/g in 2014 and 2015. Highest concentrations of catechin were measured in the flesh of Tetovka (S21) from 2014 (3.11 mg/g) and Gala (S7) from 2015 (1.91 mg/g), whereas epicatechin was most abundant in Kojce (S14), 2.20 and 2.12 mg/g in 2014 and 2015. Procyanoindin B1 was measured at highest level in the flesh of Tetovka with about 1.2 mg/g for the samples from both years.

Tetovka (S21), as the most famous autochthonous cultivar from Macedonia, contained 3.11 and 4.39 mg/g catechin and 2.35 and 1.08 mg/g epicatechin measured in the samples from 2014 and 2015. Similar as the amounts found in the peel of Tetovka, the flesh of Livadarka, Red Delicious and Kojce contained 40-50% flavan-3-ols from the total polyphenolic compounds in the flesh. Analogous to the flesh, epicatechin in the range from 1.32 to 2.31 mg/g and catechin in the range from 1.09-2.08 mg/g were also the most abundant flavan-3-ols in the apple leaves found in all analyzed cultivars.

Dihydrochalcones: Dihydrochalcones and their derivatives were the second major phenolic class in apple fruit and leaf extracts, also known as taxonomical markers for the *Malus* genus. Similar results for polyphenols extracted from apple peel and flesh have been reported [5, 7]. Total maximal dihydrochalcones measured in the apple flesh in the samples from 2014 and 2015 was 4.21 and 4.63 mg/g, respectively. Lower amounts were measured in the apple peel (about 3.68 in 2014 and 3.28 mg/g in 2015). Maximal concentrations of phloretin, phloretin rutinoside and 3-hydroxyphloretin were found in the flesh of Golden Delicious (S1) in both years (Figure 1, Table 3S). The third phloretin derivative assigned as peak 15 (Table 3S) was most abundant in the flesh of Tetovka (S21) and Sareno blago (S20). Highest amount of phloridzin was found in the flesh of Belle de Boskoop cultivar (S19), 1.88 mg/g measured in 2014 and 2015. The peel of Tetovka (S21) contained highest amounts of dihydrochalcones, followed by Sareno blago (S20), Belle de Boskoop (S19) and Pasalma (S18).

Lowest concentration of dihydrochalcones when apple peel, flesh and leaves are compared was measured in the air dried leaves (Tetovka from 2014 contained 2.66 mg/g, and from 2015, 2.81 mg/g). Phloretin, phloridzin and phloretin derivatives were also found in apple leaves. Leaves from other cultivars contained amounts in the range from 1.47 to 2.81 mg/g (Table 1S).

Flavonols: Table 1 shows that out of eight identified flavonols, five were found in the peel, three in the leaves and only two were detected in the apple flesh. Quercetin glycosides were also reported by Tsao *et al.* [7]. Although different classes of flavonols are characteristic for the peel and leaves from the same cultivar, the peel (0.51-3.04 mg/g) and leaves (1.65-3.29 mg/g) contained comparable concentrations of flavonols. Detailed analysis of the peel provided identification of 3 flavonol glycosides, assigned as peak 19, 20 and 21 (Table 1) present with highest amounts in Tetovka cultivar (S21) (0.90, 0.60 and 0.55 mg/g, respectively). The flesh contained flavonols in the range from 0.16 to 2.15 mg/g. The flesh of Belle de Boskoop cultivar (S19) was most abundant in isoquercetin (1.77 mg/g), and that of Golden Delicious (S1) from 2015 had the highest content of quercetin (1.28 mg/g). Kaempferol was most abundant in the leaves of Tetovka (S21) from 2014 (1.15 mg/g), followed by quercetin and myricetin.

Flavones: Flavones are group of compounds found only in apple leaves with concentrations from 1.45-2.43 mg/g air dried sample or 14.5-25 % of the total polyphenols measured in the apple leaves. Luteolin was the most abundant flavone in the leaves of Tetovka (S21) with measured concentrations of 1.20 mg/g in 2014 and 1.14 mg/g in 2015 (air dried sample), followed by the leaves of Pasalma (S18) 1.08 mg/g and 1.02 mg/g in 2014 and 2015) and Karapasa (S16) with about 0.8 mg/g luteolin measured in both years. Lowest amounts of luteolin were found in the leaves of Golden Delicious (S1) (0.73 mg/g and 0.67 mg/g in 2014 and 2015). Apigenin was the second major flavone with highest amounts in the leaves of Tetovka (S21), 0.94 mg/g, Pasalma (S18), 0.88 mg/g and Red Delicious (S2), 0.83 mg/g; all samples from both years contained apigenin in the range from 0.7-0.94 mg/g (Table 1S). Apigenin-7-glucoside was found in lower concentrations, from 0.02-0.26 mg/g and with concentrations similar to the other two flavones, highest in the leaves of Tetovka (S21) and Karapasa (S16), followed by Pasalma (S18), 0.20 mg/g and lowest amounts in the leaves of Golden Delicious (S1), 0.02 mg/g.

Influence of storage on polyphenol content: The effect of storage on the polyphenol profile in the apple cultivars Granny Smith and Fuji was tested during 5 months of storage at 0°C and 90% relative humidity. Figure 2 shows that the content of all polyphenols from the peel and the pulp are decreasing during time (data given in Supplementary table 7S). Moreover, the peel of Fuji, containing lower amount of polyphenols, exhibited more intensive decrease in polyphenols in the first 2 months compared with the peel of Granny Smith that was mainly due to the decrease of phenolic acids content. The major decrease in polyphenols in the flesh was observed in the first month and was mainly due to the decrease of flavan-3-ols content, especially epicatechin. The influence of storage has been investigated by Napolitano *et al.* [4] where analysis of the antioxidant activities showed increases during 3 months of storage and then decreases at 4 months of storage. Golding *et al.* showed that although peel phenolics vary among cultivars, they remain relatively stable during storage in air at 0°C and by treatment with diphenylamine indicating a low peel phenolic metabolism during long-term storage [29]. Small but significant differences were observed by Sluis *et al.* in the catechin content of the whole fruit stored in cold chambers for 4 months for Golden Delicious, Cox's Orange and Elstar apples, [30]. Our results demonstrate that there might be cultivar differences that affect the stability of polyphenols during storage that need further studies with suitable analytical methods such as the method developed in this work.

In conclusion, 27 phenolic compounds have been detected and measured in water/methanol (10:90, v/v) extracts of peel, flesh and leaves of 21 apple cultivars from Republic of Macedonia illustrating a complex profile of phenolic acids, procyandins, dihydrochalcones, flavonols and flavones and their derivatives assayed by UHPLC-DAD-HESI-MS. The methodology developed in this work enabled quantitative determination of the phenolic compounds present not only in the peel and the flesh but also in the leaves from the analysed apple cultivars. The total polyphenolic contents, as well as the major groups and individual compounds, varied significantly among the 21 studied apple cultivars. The autochthonous cultivar Tetovka was the variety with the highest content of polyphenolic compounds in the flesh and the peel, whereas the widespread cultivar Golden Delicious exhibited the lowest polyphenolic content. In all cases apple peel showed higher phenolic content than flesh extracts, except in the case of phloretin, which was found to be more abundant in the apple flesh. The results

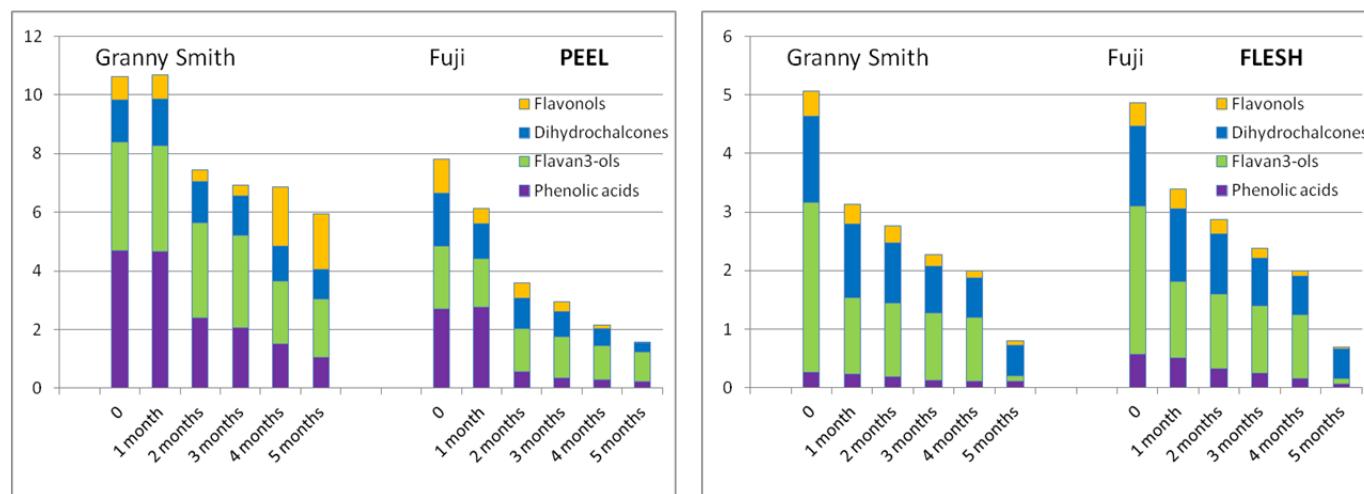


Figure 2: The effect of 5 months of storage on the polyphenolic content and distribution of classes of polyphenolic compounds in the peel and flesh studied on two commercial apple cultivars (Granny Smith and Fuji).

obtained in this study will contribute to the understanding of the polyphenolic composition of apples cultivars, especially for autochthonous cultivars, as well as to their protection.

Experimental

Reagents and standards: Methanol and acetone of HPLC grade were from Merck (Darmstadt, Germany). Water used in all procedures (extraction, separation) was of HPLC quality. Standards of gallic acid, procyanidin B2, chlorogenic acid, catechin, *p*-coumaric acid, epicatechin, caffecic acid, phloridzin, phloretin, rutin, quercetin, kaempferol, luteolin, apigenin-7-*O*-glycoside, apigenin and myricetin were from Sigma Aldrich (Steinheim, Germany).

Table 2: List of the studied apple cultivars.

Code	Cultivar	Type	Region
S1	Golden Delicious	commercial	Resen
S2	Red Delicious	commercial	Resen
S3	Cadel	commercial, local	Resen
S4	Braeburn	commercial	Resen
S5	Granny Smith	commercial	Skopje
S6	Idared	commercial	Resen
S7	Gala	commercial	Resen
S8	Mutsu	commercial	Resen
S9	Jonatan	old variety	Resen
S10	Fuji	commercial	Skopje
S11	Melrose	commercial	Resen
S12	Srcika	autochthonous	Resen
S13	Adam's Pearmain (Parmenka) domesticated	domesticated	Resen
S14	Kojce	autochthonous	Resen
S15	Livadarka	autochthonous	Resen
S16	Karapasa	autochthonous	Resen
S17	Mislinka	autochthonous	Resen
S18	Pasalma	autochthonous	Resen
S19	Belle de Boskoop (Kozara)	domesticated	Resen
S20	Sareno blago	autochthonous	Resen
S21	Tetovka	autochthonous	Resen

Materials: The list of the 21 sample of apple cultivars analyzed in this study is presented in Table 2. Besides Granny Smith and Fuji, which were grown in the region of Skopje (R. Macedonia), all other cultivars used in this study were grown in the region of Resen (R. Macedonia). From each apple cultivar (S1 to S21) peel, flesh and leaves harvested during 2014 and 2015 were separately analyzed.

After achieving full maturity, apple fruits were harvested during September 2014 and 2015. Sampling was randomly made by picking fruits from the top, central and bottom parts of the trees. Leaves from the same apple cultivar were also collected from the apple trees in the same manner during July 2014 and 2015.

Apple fruits of the two cultivars Granny Smith and Fuji were stored in an industrial refrigerator with controlled conditions: temperature 0°C and relative humidity 90%. Samples were taken for analysis of polyphenols every month during 5 months.

Sample preparation: As proposed in our previous work [28], 15-20 apple fruits were collected from each tree. Prior to analysis, apple fruits were washed with distilled water and brushed while washing to avoid contamination. Fruits were then manually skinned; the peel was separated from the flesh and blotted on paper. Averaged samples from peel and flesh of all apples from each cultivar were prepared. Peel and flesh were then lyophilized as fast as possible to avoid browning. Peel and flesh were placed in plastic plates (about

20 g of fresh sample) and were lyophilized for 24 h at -45°C and 0.055 mBar. After lyophilisation, residuals from each apple cultivar were ground with a blender at 400 W. After blending, all samples (leaves, flesh and peel) were kept at -20°C till analysis.

About 50 g of fresh leaves were collected from each apple tree cultivar. After collection, leaves were air dried and then ground in a blender.

Extraction of the polyphenols from the peel, flesh and leaves of the 21 apple cultivars was made with methanol/water (90:10, v/v). About 0.5 g flesh/peel/leaves was extracted with 2 portions (2.5 mL) of the solvent using ultrasound at room temperature for 1 h. After centrifugation (3000 rpm/min) for 10 min the supernatants were combined and made up to a final volume of 5 mL. All samples were prepared in triplicate. The extracts were filtered through 0.45 µm membrane filters (PTFE, Agilent) and then analyzed by spectrophotometry [28], and by HPLC. A balance sensitive to 0.1 mg (Sartorius, Germany), an ultrasonic bath Elmasonic S100H, Freeze-dryer Freezone 2.5 (Labconco, USA), a BOSH blender and a centrifuge Harrie 15/80 MSE were used for sample preparation.

UHPLC-DAD-HESI-MS^a apparatus and conditions: The analysis was performed on a Thermo Scientific Dionex Ultimate 3000 UHPLC-UV-DAD system coupled to a Thermo Scientific LTQ XL linear ion trap mass spectrometer with heated electrospray ionisation (HESI) probe and controlled by Chromeleon 5.0 acquisition and Xcalibur 2.0 data analysis software. Zorbax Eclipse plus C18 column (Agilent Technologies), 2.1 x 50 mm i.d., and 1.8 µm particles were used for separation. Elution was performed at a flow rate of 0.7 mL/min, with a mobile phase consisting of 1%, v/v, formic acid in water (phase A) and 1% formic acid in methanol (phase B) with the following gradient: 0-1.5 min 10% B; 1.5-4.5 min linear gradient from 10 to 35% B; 4.5-5.5 min 35% B; from 5.5 to 8.5 min 40% B; from 8.5-9.0 min gradient to 45% B, from 9.0-10.0 min 45% B; from 10.0-10.5 linear gradient to 80% B; from 10.5-14.5% 80% B. The injection volume was 5 µL, and the column temperature was 40°C, UV detection was carried out at wavelengths of 260, 280, 330, 350 and 520 nm. Peak areas for quantification were taken at 330 nm for all detected polyphenolic compounds since all exhibited absorption at this wavelength. Ion source parameters were as follows: source voltage 4 kV, capillary voltage -40 V, tube lens voltage -80 V, heater temperature 285°C, capillary temperature 275°C, sheet and auxiliary gas flow (N₂) 42 and 11 psi. Double heating of the probe (heater temperature) was employed in order to use higher flow rate of the mobile phase. MS were acquired by full range acquisition covering *m/z* 100-1500. Fragmentation studies, data dependent scans and "in source" fragmentations were performed by developing collision induced dissociation (CID). The normalized collision energy of the collision induced dissociation (CID) was set at 35 eV. Fragmentation mechanisms and characteristic fragments were confirmed using Mass Frontier spectral interpretation software 7.1.

Supplementary data: Supplementary tables: 1S-6S contain quantitative results for all individual polyphenolic compounds; 7S contains data with the content of polyphenolic compounds in apple peel and flesh during 5 months of storage of the cultivars Granny Smith and Fuji; 8S contains meteorological data for 2014 and 2015.

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Preparation and Validated Analysis of Anthocyanin Concentrate from the Calyces of *Hibiscus sabdariffa*

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Extracts of *Hibiscus sabdariffa* calyces have been shown to have various medicinal properties, some of which have been reported to be due to anthocyanins present in the calyces. To study whether these claims are valid, it is necessary to produce an extract with a high anthocyanin content and to have available a method to identify and quantify the individual compounds in the product. A method of extraction and purification has been developed based on a polyamide column chromatographic purification step. Using this method, anthocyanin concentrates were produced containing from 57 to 64% of delphinidin-3-sambubioside plus cyanidin-3-sambubioside. A rapid, efficient and validated HPLC analytical method was developed and used for the analysis of the anthocyanin concentrate.

Keywords: *Hibiscus sabdariffa*, Malvaceae, Calyces, Anthocyanins, Polyamide purification, Validated HPLC analysis.

Hibiscus sabdariffa L. (family Malvaceae) is commonly known in English as roselle, rosella and red sorrel, and in Arabic as karkadeh. Aqueous extracts of the red calyces of the flower are drunk both hot and cold as beverages, but they are also used in folk medicine against many illnesses. The phytochemical, pharmacological and toxicological aspects of *H. sabdariffa* were reviewed by Ali *et al.* in 2005 [1]. Since then, other beneficial effects resulting from the use of the calyx extract have been reported [2, and references quoted therein].

The red color of the calyces is due to the presence of anthocyanins and these compounds have been linked to many of the reported medicinal properties [for example, 2, 3a-e]. The anthocyanins found in *H. sabdariffa* have been shown to vary in different collections. The compounds reported include delphinidinpentoside-glucoside [4a], delphinidin [4b,d], delphinidin-3-sambubioside [4c,d,e,h], delphinidin-3-monoglucoside [4d,e], cyanidin [4b], cyanidin-3-monoglucoside [4d,e], and cyanidin-3-sambubioside [4e,h]. In *H. sabdariffa* var. *altissima*, the anthocyanins reported were cyanidin-3,5-diglucoside and cyanidin-3-glucosylrutinoside [4f]. In the calyces of five strains of *H. sabdariffa* var. *sabdariffa*, cyanidin-3-sambubioside was the major pigment, followed by cyanidin-3-monoglucoside. Delphinidin glycosides were not detected in one strain, but were present in the other four. During calyx growth, the anthocyanin content reached 1.7% to 2.5% of the dry weight in all strains [4g].

We have been studying the biological effects of aqueous extracts of *H. sabdariffa* calyces and trying to determine whether these are due, at least in part, to the anthocyanins present in the extract. For this, it

has been necessary to purify the extracts to obtain products with a high content of anthocyanins. An analytical procedure to determine the compounds present and their contents was also necessary. Here we report the development of the isolation and preparation of a high anthocyanin containing product from the calyces of *H. sabdariffa* and an analytical procedure to identify and quantify the compounds present.

The plant material used was from the same batch as that used in an earlier study. The major anthocyanins were delphinidin-3-sambubioside and cyanidin-3-sambubioside, although these were accompanied by much smaller quantities of other delphinidin and cyanidin glycosides, as well as rutin and chlorogenic acid [4h].

For extraction of the anthocyanins, water was found to be a less suitable solvent than ethanol, because the calyces of *H. sabdariffa* contain mucilage and pectins [5a,b]; these substances increase the solution viscosity and significantly complicate filtration. The ethanolic extract of the calyces was evaporated to dryness and the residue dissolved in water. This aqueous solution was then purified by passage through a column of polyamide resin. This process was chosen so that the final mixture would contain a minimum of cyanidin, delphinidin and their monoglucosides, as these were eluted from the column easily and relatively quickly with water. Both delphinidin- and cyanidin-3-sambubiosides were subsequently eluted by 95% ethanol. After evaporation of the eluate, the residue was re-dissolved in 95% ethanol and the solution mixed with diethyl ether. The resulting precipitate, after removal by filtration and drying was powdered to provide the final product.

Table 1: Composition of anthocyanin concentrate prepared from the calyces of *H. sabdariffa*.

Compound	Content range
Delphinidin-3-sambubioside	46.1 ± 1.1 to 51.2 ± 0.9
Cyanidin-3-sambubioside	11.1 ± 0.8 to 13.5 ± 1.0
Rutin	2.8 ± 0.1 to 3.9 ± 0.3
Chlorogenic acid	0.9 ± 0.07 to 1.14 ± 0.08

*all data were calculated on dry weight of anthocyanin mixture; every analysis was repeated three times.

**three separate samples from each anthocyanin mixture were weighed and each sample was analyzed twice (total 6 analyses of each anthocyanin mixture).

A rapid and efficient HPLC method for separation and analysis of the individual components of the anthocyanin concentrate was developed. All the major components were well separated. The predominant compounds, as before, were delphinidin-3-sambubioside and cyanidin-3-sambubioside, along with small quantities of rutin and chlorogenic acid [4h] (Table 1).

Polyamide is an optimal adsorbent for the separation of water-soluble phenolic and non-phenolic compounds. It was used earlier to separate the high concentration of aliphatic carboxylic acids in the calyxes of *H. sabdariffa* [5c,d]. However, in our work, total removal of these acids did not seem to be effective and during an attempt to increase further the anthocyanin content of the concentrated extract by repeating the column chromatographic purification step, decomposition products were produced resulting in a grey-blue color indicating the negative impact of higher pH. However, both the delphinidin-3-sambubioside (57.4 ± 2.1 to 63.2 ± 2.4) and cyanidin-3-sambubioside (16.1 ± 0.7 to 17.4 ± 1.5) contents were increased, as well as those of rutin (3.2 ± 0.2 to 4.3 ± 0.3) and chlorogenic acid (1.05 ± 0.10 to 1.63 ± 0.09).

The identities of the non-specified components of the anthocyanin concentrate were not established. Some authors report that protocatechuic acid is an important phenolic acid present in *H. sabdariffa* extract [5e, f]; this was isolated from the dried flowers of the plant [5g,h]. The content in *H. sabdariffa* was reported as 1.19 ± 0.12% [5i]. Chlorogenic acid has also been previously recorded (0.27% in the calyxes) [5j], as have rutin (0.21%) and quercetin (0.32%) [5j].

Using the described method, an extract of *H. sabdariffa* calyxes was produced which contained greater than 55% of the major two anthocyanins. This product is the one being used in experiments to elucidate the potential pharmacological role of the anthocyanins of *H. sabdariffa*.

Experimental

Plant material: Dried *Hibiscus sabdariffa* calyxes were imported from Sudan. Their identification was verified by comparison with authentic material. A sample has been lodged in the Herbarium of the Faculty of Pharmacy, Hradec Králové, under the following number CUFPH16130/625.

Adsorbent preparation: Polyamide pearls {Silamid (Nylon N6), Povážske chemické závody, Slovak Republic; 150 g} were dissolved in 1.5 L of glacial acetic acid under reflux (about 1 h). The resulting slightly yellow, viscous solution was poured into a 5 L beaker and allowed to cool for several hours. The polyamide precipitated as a waxy white suspension. Water (400-500 mL) was added to the suspension, which was well stirred, then filtered on a Büchner funnel, and washed with water. The sediment on the filter was suspended in 2 L of water and alkalized carefully by the addition of 5M NH₄OH (pH ~8). After 30 min the suspension was poured into a chromatographic column and washed with distilled water until the eluate was neutral. The adsorbent was then

suspended in 95% ethanol, filtered on a Büchner funnel and the insoluble material dried at room temperature, until the moisture content was 25%. The mass was then passed through a sieve (630 µm) and the material dried again at 60°C to a moisture content of 15%. Final screening was through vibration sieves of 200 µm and 500 µm. For separation of anthocyanins the 200-500 µm fraction was used.

Column preparation: A chromatographic column (average : height 19 : 20 cm) was filled to one third of its volume with water and then, portion-wise, a thick slurry of 600 g of pulverized polyamide (200-500 µm), prepared as above, was slowly added to the column (the adsorbent must be suspended for at least 15 min in the water before pouring).

The column was easily regenerated by washing with boiled water (7 L) at room temperature (air-free). The regenerated column can be used maximally 4 times without change in separation characteristics.

Preparation of anthocyanins concentrate: Milled calyxes (300 g) were added to 2.5 L of boiling 95% ethanol, and the mixture heated, while stirring, for 15 min under reflux. The extract was then decanted and the residue extracted in the same way two more times. The combined extracts were filtered and the filtrate evaporated to dryness. The residue was dissolved in 1 L of water (60°C), with brief sonication, and, after cooling, filtered through a Büchner funnel (filter paper Macherey Nagel MN 640D); the volume of filtrate was made up to 1.5 L. This red solution was filtered through a polyamide column (elution 26-30 mL/min), which was then washed with 6-7 L of water. Then the anthocyanin sambubiosides were eluted by 6-7 L of 95% ethanol.

The red-violet eluate, after evaporation (10-11 g), was dissolved in 100 mL of 95% ethanol and mixed with 750 mL diethyl ether. The precipitate formed was removed by filtration after 15 min, and after a short period of air-drying was dried in a vacuum desiccator (~0.7 kPa, 12 h) over silica gel and then powdered (6-7.6 g).

Table 2: Method validation.

Parameter	Delph-3-samb	Cyan-3-samb	Chlorog acid	Rutin	Criteria
SST					
Repeatability - t _R ^a	0.23	0.19	0.23	0.03	X < 1%
Repeatability - peak areas ^a	0.84	0.57	0.65	0.12	X < 1%
Theoretical plates ^b	2.370	2.768	6.462	-	-
Peak capacity ^b	-	-	-	29	-
Resolution ^b	3.13	4.38	5.94	13.07	R _{ij} > 1.5
Symmetry factor ^b	1.01	0.97	1.15	1.12	0.8 -1.5
VALIDATION					
Precision ^c RSD (%)	1.69	4.86	2.32	4.22	RDS (%) < 5%
Linearity					
Correlation coefficient	0.999	0.999	0.999	0.999	R > 0.999
Calibration curve	10480x + 88472	20301x + 78142	+ 1966x 2576	+ 17053x 14065	-
Calibration	10-120 mg/L	5.0-70 mg/L	2.5-20 mg/L	6.8-72 mg/L	-
Accuracy	99.89	100.69	97.56	103.43	X = 100 ± 5%
recovery ^d (%)	-	-	-	-	-
Accuracy RSD ^d (%)	0.56	1.23	1.50	2.83	X < 5%

Delph-3-samb = delphinidin-3-sambubioside; Cyan-3-samb = cyanidin-3-sambubioside; Chlorog acid = chlorogenic acid.

^a made in six replicates at 50 mg/L delphinidin-3-sambubioside and cyanidin-3-sambubioside, 15 mg/L chlorogenic acid and 40 mg/L for rutin. ^b made in three replicates. ^c six solutions injected three times each. ^d samples spiked with: 20 mg/L delphinidin-3-sambubioside, 20 mg/L cyanidin-3-sambubioside, 5 mg/L chlorogenic acid and 15 mg/L rutin

Chromatographic analysis: HPLC analysis was performed on a Shimadzu LC-2010 C system (Shimadzu, Kyoto, Japan) equipped

with a built-in UV-VIS detector. A Supelco Discovery HS C18 column (250 x 4.6 mm, 5 µm) was used for the analysis of the extract. Isocratic elution continued for 9 min with a mobile phase composed of 10% acetonitrile (ACN) and 90% formic acid (2%); thereafter isocratic elution was changed to gradient mode and the concentration increased within 8 min to 60% ACN followed by a 7 min equilibration to the initial conditions. The mobile phase flow rate was 1 mL/min. The detection was set at 520 nm for anthocyanins and 371 nm for chlorogenic acid and rutin.

Preparation of standard solution: A standard solution was prepared by dissolving 0.5 mg of delphinidin-3-sambubioside chloride (Extrasynthese, Genay, France), 0.5 mg of cyanidin-3-sambubioside chloride (Extrasynthese, Genay, France), 0.15 mg of chlorogenic acid (Sigma-Aldrich/Merck, Czech division, Prague) and 0.4 mg of rutin trihydrate (Sigma-Aldrich/Merck, Czech division, Prague) in 10 mL of methanol.

Sample preparation: The anthocyanin concentrate (0.5 mg) was dissolved in 1 mL of methanol and filtered through a 0.45 µm PTFE filter into an amber vial prior to analysis by HPLC.

Method validation: Method validation was performed to demonstrate the suitability of the chosen method for its intended use. The validation included a system suitability test (SST) and the determination of method precision, accuracy and linearity. The standard solution was injected 6 times to calculate the SST parameters. Relative standard deviations (% RSD) of retention times and area, tailing factor, number of theoretical plates (peak capacity for rutin analyzed in a gradient mode), and resolution, were calculated.

The standard calibration curves were evaluated using 6 calibration levels. Correlation coefficients were > 0.999 for all analytes. The precision was verified on *H. sabdariffa* anthocyanin concentrate. The sample was prepared 6 times and analyzed using HPLC. Relative standard deviations (% RSD) of areas obtained by the recalculation to the same sample amount were evaluated. The accuracy (% of recovery, % of RSD) was verified using standard additions procedure with samples spiked with a reference standard. Method validation results are depicted in Table 2.

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The Influence of Extraction Parameters on Antimicrobial Activity of Propolis Extracts

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The extraction optimization of the poplar-type propolis was performed in order to improve the isolation of flavonoids as well as the corresponding antimicrobial activity of the products obtained. The efficiency of flavonoids extraction depended upon the type of extraction media used, following the rank 80% ethanol > 40% ethanol >> water, regardless of pH value. Ultrasound assisted extraction was as efficient as the maceration procedure, offering additional benefits such as short duration time and low extraction temperature. The antimicrobial efficiency of extracts prepared with 80 and 40% ethanol against the tested microbial stains was comparable, regardless of the extraction technique used, while aqueous extracts mainly showed scarce activity. Observed activity against the yeast *Candida albicans* strongly correlated with flavones and flavonols content in extracts prepared ($r^2=0.8217$), while regression analysis showed that beside flavonoids, some other components which were successfully extracted from the crude propolis contributed to the observed antimicrobial efficiency against *Bacillus subtilis* and *Staphylococcus aureus*.

Keywords: Poplar-type propolis, Flavonoids, Maceration, Ultrasound extraction, *Bacillus subtilis*, *Staphylococcus aureus*, *Candida albicans*.

Propolis, a resinous substance collected and enzymatically modified by honeybees (*Apis mellifera L.*) from exudates and buds of plants, has been used as a traditional remedy in folk medicine since ancient times, especially in Europe and Asia. To date, more than 300 different constituents have been identified in propolis, resulting in a wide spectrum of biological activities. The chemical composition of propolis is highly dependent on the phytographic characteristics of the flora present around the collection site, leading to formulation of several chemical types of propolis, based on their plant source [1-3]. Its documented efficiency in the treatment of cold and upper respiratory tract infections, wound healing, treatment of burns, acne, herpex simplex infection and neurodermitis [2,3], as well as in prevention and treatment of dental diseases [4] puts propolis in the focus of interest of the pharmaceutical industry, especially taking into account that the number of pathogens that are developing resistance to known antibiotics is rising [5].

The antimicrobial efficiency of propolis varies greatly depending on the extraction procedure of the crude material and the solvents used in this process. In this paper, the efficiency of maceration, as a standard propolis extraction technique, was compared with that of ultrasound assisted extraction, in order to select rationally the best extraction technique. In order to optimize additionally the extraction of the bioactive compounds from the crude propolis, a series of extraction media were used, containing variable amounts of ethanol and water at different pH values. Total flavonoid content in the extracts prepared, determined by two complementary colorimetric methods [6], was taken as an indicator of the extraction efficiency. This selection was made taking into account that hydrophobic flavonoid-aglycones are the main constituents of poplar type propolis, which is characteristic for raw propolis samples collected from inland Croatia [7] and are responsible for the majority of the propolis biological activities [1, 8]. The results obtained for extracts prepared are shown in Table 1. Surprisingly, for some extracts prepared by maceration (namely M-80-50-3, M-80-25-7, M-80-25-3 and M-40-25-7), the method with 2,4-dinitrophenylhydrazine resulted in unrealistically high values (data not shown). Since it was

previously shown that flavones, flavonols and isoflavonols with a C2-C3 double bond could not react with 2,4-dinitrophenylhydrazine [6], the observed results were attributed to unknown interferences, but a detailed study of these was not within the scope of this paper.

When analyzing the amount of flavonoids extracted (Table 1), it is obvious that the efficiency of their extraction was mainly dependent upon the ethanol concentration in the extraction media, while pH and temperature of extraction were not critical parameters. In the case of flavones and flavonols, the highest extraction efficiency was obtained when 80% ethanol was used as the extraction medium, while 40% ethanol was somewhat less effective. Water, regardless of the pH used, appeared to be the least efficient extraction medium, resulting in extracts containing only around 0.6%, w/v, of flavones and flavonols (Table 1). Such a result can be related to the rather poor aqueous solubility of those compounds and is in agreement with literature data [2]. When comparing the efficiency of the two extraction methods used, ultrasound assisted extraction resulted in comparable amounts of flavones and flavonols extracted, regardless of the type and pH of the extraction media. However, compared with maceration, the use of ultrasound significantly reduced the extraction time and allowed the preparation of propolis extract at rather low temperature (4°C), thus clearly demonstrating its advantages over maceration. When comparing the amounts of flavanones in the extracts prepared, maceration with 80 and 40% ethanol yielded comparable amounts of flavanones, which were in the range previously found to be characteristic of the propolis from the western part of continental Croatia [7]. Ultrasound assisted extraction using 80% ethanol resulted in comparable amounts of flavanones, while somewhat lower amount of flavanones were extracted when 40% ethanol was used as extraction medium. When water was the extraction medium, both methods resulted in comparable amounts of flavanones extracted, ranging from 6.1 to 6.9% (Table 1).

Finally, when comparing the amounts of flavonoids extracted (Table 1) to their quantity in some commercially available propolis

Table 1: Parameters of the extraction process and flavonoids content in liquid extract prepared.

Sample code	extraction method	solvent	temperature (°C)	pH	Flavonoids (%) ^a	Flavanones ^b
					Flavones and flavonols ^a	Flavanones ^b
M-80-50-7	maceration	ethanol 80%	50	7	22.31	12.51
M-80-50-3		ethanol 80%	50	3	21.88	*
M-80-25-7		ethanol 80%	25	7	21.25	*
M-80-25-3		ethanol 80%	25	3	22.38	*
M-40-50-7		ethanol 40%	50	7	6.94	15.63
M-40-50-3		ethanol 40%	50	3	5.63	10.91
M-40-25-7		ethanol 40%	25	7	3.85	*
M-40-25-3		ethanol 40%	25	3	4.32	13.94
M-W-50-7		water	50	7	0.59	6.57
M-W-50-3		water	50	3	0.59	6.92
M-W-25-7		water	25	7	0.67	6.34
M-W-25-3		water	25	3	0.40	6.43
UZV-80-4-7	ultrasonication	ethanol 80%	4	7	20.63	12.23
UZV-80-4-3		ethanol 80%	4	3	22.13	11.97
UZV-40-4-7		ethanol 40%	4	7	4.73	7.13
UZV-40-4-3		ethanol 40%	4	3	5.06	7.18
UZV-W-4-7		water	4	7	0.51	6.39
UZV-W-4-3		water	4	3	0.44	6.13

^a Expressed as quercetin equivalent; ^b Expressed as naringenin equivalent; * Flavone content determination was hindered by interferences.

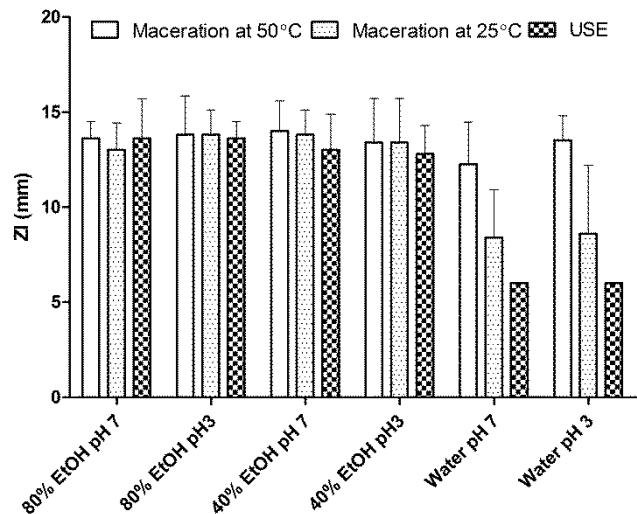


Figure 1: Antibacterial activity of prepared propolis extracts against *Bacillus subtilis* NCTC 8236 expressed as zones of inhibition (ZI) measured in mm (mean±SD, n=6; USE=Ultrasound assisted extraction).

tinctures [9], maceration and ultrasound assisted extraction with hydroethanolic media resulted in 10 to 60 times higher concentrations of flavones and flavonols, while levels of flavanones were comparable. This clearly showed the need for careful adjustment of the extraction parameters and the methodology used in order to extract successfully the poorly soluble compounds from the crude propolis.

The antimicrobial activity of the propolis extracts prepared against *Bacillus subtilis* NCTC 8236, *Staphylococcus aureus* ATCC 29213, and *Candida albicans* ATCC 10231 strains was evaluated by measuring the zones of inhibition using a cylinder-plate diffusion method. The obtained results are shown in Figures 1, 2 and 3. In the diffusion assay, all the media used for propolis extraction showed no antimicrobial action against the microbial species tested (data not shown). On the contrary, propolis extracts prepared with 80 or 40% ethanol, regardless of the pH and extraction technique used for preparation, showed a pronounced antibacterial activity against *B. subtilis* and *S. aureus* (Figures 1 and 2). In fact, *S. aureus* presented

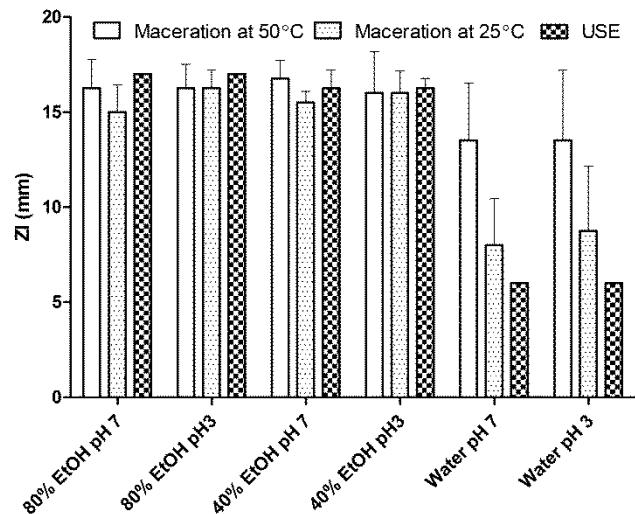


Figure 2: Antibacterial activity of prepared propolis extracts against *Staphylococcus aureus* ATCC 29213 expressed as zones of inhibition (ZI) measured in mm (mean±SD, n=6; USE=Ultrasound assisted extraction).

inhibition zones ranging from 15-17 mm, while *B. subtilis* was somewhat less susceptible to ethanolic extracts of propolis, presenting inhibition zones ranging from 13 to 14 mm. In general, such results are in agreement with previously published data which showed good antimicrobial activity of ethanolic propolis extracts against those microbial species [10].

Analysis of the antibacterial action of propolis extracts against *B. subtilis* and *S. aureus* showed that the ethanol concentration (40 or 80%) and the pH value of the extraction media, as well as the technique used for the extract preparation did not have any statistically significant influence on the observed antimicrobial activity of the extract prepared ($p>0.05$; Figures 1 and 2). This further confirmed the high efficiency of ultrasound assisted extraction in the extraction of antibacterial compounds from the crude propolis, when hydroethanolic extraction media were used. Surprisingly, even the aqueous propolis extracts, prepared by maceration at 50°C (M-W-50-7 and M-W-50-3, respectively), showed comparable antimicrobial efficiency against both microbes

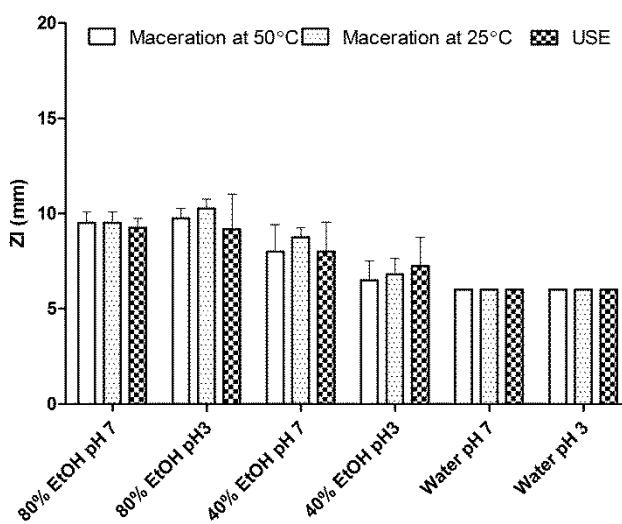


Figure 3: Antifungal activity of prepared propolis extracts against *Candida albicans* ATCC 10231 expressed as zones of inhibition (ZI) measured in mm (mean \pm SD, n=6; USE=Ultrasound assisted extraction).

as observed for ethanolic extracts ($p>0.05$). However, when maceration with water was performed at lower temperatures (25°C), the prepared extracts, namely M-W-25-7 and M-W-25-3, showed approximately 30% lower antimicrobial activity ($p<0.01$), indicating that at this temperature, the extraction of the antibacterial compounds from the crude propolis was less efficient. When aqueous extracts of crude propolis were prepared by ultrasound assisted extraction, the obtained extracts (US-W-4-3 and US-W-4-7) showed no antibacterial effect against *B. subtilis* and *S. aureus* (Figures 2 and 3). Probably, the low temperature at which extraction was performed hampered the successful extraction of the antimicrobial compounds from the raw propolis. Finally, when comparing the antimicrobial efficiency of all extracts prepared, it might be concluded that the pH value of the extraction media is not a critical parameter which would influence the antibacterial activity of the extract prepared.

When the observed antibacterial activity of extracts prepared was correlated with the content of flavones and flavonols, a weak but significant positive correlation was found ($r^2=0.3734$ and 0.3150, for *B. subtilis* and *S. aureus*, respectively). No statistically significant correlation was found between antimicrobial activity of extracts prepared and flavanones content. This indicated that, beside flavones and flavonols, some other components of the propolis with antibacterial action were successfully extracted from the crude material, and contributed to the antimicrobial action of the extracts prepared. Furthermore, such an assumption would also explain the lack of antimicrobial activity of aqueous propolis extracts prepared by ultrasound assisted extraction (US-W-4-7 and US-W-4-3), which contained comparable amount of flavones and flavonols as M-W-50-3 and M-W-50-7 (Table 1), but lacked any antimicrobial activity. The in depth study of their chemistry, especially in the case of aqueous propolis extracts prepared by maceration at 50°C, is the focus of upcoming studies.

In the case of the activity of propolis extracts against the yeast *C. albicans*, somewhat different results were observed (Figure 3). In general, *C. albicans* is less susceptible to the propolis extracts prepared, showing inhibition zones ranging from 9 to 11 mm. Again, extracts prepared with 80% ethanol, regardless of the pH of the extraction media and extraction technique used were the most

active (Figure 3). When 40% ethanol at pH 7 was used as the extraction medium, for both maceration and ultrasound assisted extraction, the obtained extracts (M-40-50-7, M-40-25-7 and US-40-4-7) showed lower action against *C. albicans* ($p<0.001$), with zones of inhibition around 9 mm, while the extracts prepared with 40% ethanol at pH 3 (M-40-50-3, M-40-25-3 and US-40-4-3) were practically inactive (Figure 3). Aqueous propolis extracts, regardless of the preparation method used, showed no antifungal effect against *C. albicans* ($p<0.001$).

When the observed antimicrobial efficiency of extracts against *C. albicans* was correlated with the content of flavones and flavonols, a statistically significant positive correlation was found ($r^2 = 0.8217$), while no statistically significant correlation was found between antimicrobial activity and flavanones content. This indicated that flavones and flavonols have an important role in the activity of the propolis extracts prepared, but they must be present in a sufficient concentration in order to express an anti-*Candida* effect, as was also shown by earlier investigations [9]. Therefore, the extracts of the crude propolis prepared with rather polar extraction media would be less effective. Also, it is possible that some other constituents of propolis could contribute to its anti-*Candida* action. For example, recent studies demonstrated that in the case of ethanolic extracts of Mediterranean propolis, only samples which contained significant amounts of terpenyl esters and hydroxybenzoic acid displayed activity against *C. albicans* *in vitro*, while others were ineffective [11]. All this demonstrates a need of further research in order to determine the individual contribution of each propolis constituent with regard to its anti-*Candida* action and possible synergism, as well as to understand the variability in propolis samples of different geographical origin. Only better knowledge of their chemistry would allow us to develop novel and more efficient extraction procedures, which would lead to standardized and more efficient propolis preparations for use in medicine and in the food industry.

Experimental

Propolis sample: The raw poplar-type propolis was collected from beehives in the outskirts of Zagreb, Croatia.

Preparation of propolis extracts: Prior to extraction, the raw material was frozen at -20°C and then homogenized and pulverized by grinding in a chilled mortar. The extraction was performed by maceration and an ultrasound method, using 1:5 weight ratios between crude propolis and the extraction medium. The extraction solvents used were water and 40 or 80%, v/v, ethanol/water mixtures, with the pH adjusted to 3 or 7 by addition of 0.1M HCl or NaOH solution. The maceration of raw propolis was performed in hermetically sealed Erlenmeyer flasks in a horizontal shaker thermostated at either 25 or 50°C at a shaking frequency of 120 min⁻¹ for 24 h. Ultrasound extraction was performed by introducing an ultrasonic processor probe (Misonix Sonicator® S-4000, Cole-Parmer, USA) operating at 60 W for 30 min directly into the sample, immersed in an ice-bath to prevent sample overheating. After the extraction process, the residual solids were separated from the liquid extracts by centrifugation at 3500 rpm for 10 min, followed by filtration through a Whatman cellulose filter paper No.1. The obtained liquid extracts were stored protected from light at 4°C in airtight containers.

Determination of flavones, flavonols and flavanones: Quantification of flavones and flavonols in extracts prepared was performed by a colorimetric method using aluminum chloride, as described by Woisky and Salatino [12]; results are expressed as quercetin equivalents, while flavanones contents were determined

using a 2,4-dinitrophenylhydrazine colorimetric assay proposed by Nagy and Granca [13]; results are expressed as (\pm)-naringenin equivalent.

In vitro antimicrobial assays: The antimicrobial activity of the prepared propolis extracts against *Bacillus subtilis* NCTC 8236, *Staphylococcus aureus* ATCC 29213 and *Candida albicans* ATCC 10231 was evaluated by a cylinder-plate diffusion method following the procedure described by Kosalec *et al.* [9] and using 60 μ L aliquots of undiluted propolis extracts. Prepared plates were

incubated under aerobic conditions first at +4°C for 1 h and then at +37°C for 18 h in the case of bacteria and 48 h in the case of *C. albicans*. The same procedure was utilized for all extraction media used. The diameter of the inhibition zone around each hole was measured and recorded. All values are expressed as mean \pm SD of 6 separate experiments. Data were compared by one-way ANOVA, followed by Tukey's multiple comparison post-test. The differences were considered statistically significant when $P<0.05$. Calculations were performed using the GraphPad Prism program (GraphPad Software, Inc., San Diego, CA; www.graphpad.com).

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Comparative Evaluation of Phytochemicals, and Antidiabetic and Antioxidant Activities of *Cuscuta reflexa* Grown on Different Hosts in Northern Thailand

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For the first time, total phytochemical profiles of methanol crude extracts of *Cuscuta reflexa* grown on three different hosts, *Coccinia grandis*, *Ficus racemosa* and *Samanea saman*, that were cultivated in northern Thailand were examined, along with their antidiabetic and antioxidant activities. The highest level of total flavonoids (114.4 mg QE/ g extract) and total phenolic content (90.8 mg GAE/ g extract) were observed in the extract of *Cu. reflexa* that was grown on *Co. grandis*. The GC-MS results showed that various types of phenolic compounds, hydrocarbons, saturated fatty acids and methyl ester of fatty acids, unsaturated fatty acids and methyl ester of fatty acids, vitamin E, terpenes and sterol contained in the extracts of *Cu. reflexa* were capable of being grown on three hosts. Moreover, the HPLC results showed the presence of gallic acid, catechin, vanillic acid, rutin and quercetin in all *Cu. reflexa* samples. The extract of *Cu. reflexa* that was grown on *Co. grandis* represented the highest antidiabetic activity with a percent inhibition of 51.2. Moreover, the extract also possessed the greatest antioxidant activity (DPPH; IC₅₀ 168.6 µg/mL, FRAP; 40.5 mg GAE/ g extract).

Keywords: *Cuscuta reflexa*, Phytochemical, Total flavonoid content, Total phenolic content, Antidiabetic activity, Antioxidant activity.

The parasitic plant, *Cuscuta*, is commonly used as an ingredient in functional food and in medicinal tonics as a nutrient. It is also often used in alcoholic beverages [1]. In northern Thailand, one of the most widely distributed *Cuscuta* plants is *Cu. reflexa* Roxb. This can grow on a wide variety of plant hosts including a number of perennial plants and annual crop species. Previous research reported that the chemical constituents of *Cuscuta* depend on the type of host [2–3]. Various types of phenolic compounds and their glycosides, terpenes, sterols, saturated aliphatic hydrocarbon, saturated fatty acids and unsaturated fatty acids were found in *Cu. reflexa* [4]. The study of Perveen *et al.* [5] revealed that the phenolic acid and other phenolic components, such as flavonoids found in *Cu. reflexa*, are often extracted in higher amounts in more polar solvents. Furthermore, previous research studies indicated that the stem of *Cu. reflexa* possessed antibacterial, antiviral, anticancer and antioxidant activities [6–7]. The objectives of this study were to identify the total flavonoid content (TFC) and total phenolic content (TPC), and chemical constituents in the methanol crude extracts of *Cu. reflexa* that were grown on three different hosts, *Coccinia grandis* (L.) Voigt., *Ficus racemosa* Linn. and *Samanea saman* (Jacq.) Merr., as well as to evaluate the antidiabetic and antioxidant activities of the above mentioned crude extracts.

Total flavonoid and total phenolic contents of the extracts of *Cu. reflexa* were evaluated, and the results are shown in Table 1. TFC values ranged from 67.5 to 114.4 mg QE/ g extract, while the TPC values ranged from 84.4 to 90.8 mg GAE/ g extract. The highest value of TFC and TPC was exhibited in the extract of *Cu. reflexa* grown on *Co. grandis*. Anjum *et al.* [1] demonstrated the TFC and TPC in methanol extracts of *Cuscuta* sp. that were grown on 9 different hosts. The results revealed that the TFC ranged from 47.0 to 81.0 mg catechin/ g of the dry weight, while the TPC value was in the range of 39.4 to 66.2 mg/g.

Table 1: Total flavonoid and total phenolic contents of the methanol crude extracts of *Cuscuta reflexa* grown on *Coccinia grandis*, *Ficus racemosa* and *Samanea saman*.

Host plant	Total flavonoid contents (mg QE/ g extract)	Total phenolic contents (mg GAE/ g extract)
<i>Coccinia grandis</i>	114.4±0.7a	90.8±2.9a
<i>Ficus racemosa</i>	79.0±1.8b	86.6±0.5b
<i>Samanea saman</i>	67.5±2.6c	84.4±1.4b

Average ± standard deviation from three replicates.

The different letters in the same column are considered significantly different according to Duncan's multiple comparison test (*P* < 0.05).

The phytochemicals of the all extracts of *Cu. reflexa* were analyzed by gas chromatography and mass-spectrometry (GC-MS). Each compound was identified based on mass spectral matching (≥ 90%) from W8N08 and Wiley7n libraries and is reported in Table 2. The compounds identified in all of the *Cu. reflexa* extracts were (1) phenolic compounds (4-vinylphenol, 4-vinylguaiacol, methyl cinnamate, methyl 4-hydroxycinnamate and methyl-3-(3,5-ditertbutyl-4-hydroxyphenyl) propionate), (2) hydrocarbons (cyclopentadecane, nanocosane and squalene), (3) saturated fatty acids and methyl esters of fatty acids (palmitic acid methyl ester, steric acid methyl ester, palmitic acid amide, arachidic acid methyl ester, stearic acid amide, and lignoceric acid methyl ester), (4) unsaturated fatty acids and methyl esters of fatty acids (linoleic acid methyl ester, linolenic acid methyl ester and oleic acid amide), (5) vitamin E (α- and γ-tocopherol), (6) terpenes (phytol and β-amyrin), and (7) sterols (campesterol, stigmasterol, (24RS)-24, 27-dimethylcholesta-5,26-dien-3β-ol and β-sitosterol). However, methyl ferulate, palmitic acid, pentadecanoic acid methyl ester, palmitoleic acid methyl ester, *E*-9-heptadecenal and baccharan-3β-ol were only found in the extract of *Cu. reflexa* grown on *Co. grandis*. Additionally, docosanoic acid methyl ester was only detected in the extract of *Cu. reflexa* grown on *S. saman*. As in the study of Bais and Kakkar [3], the chemical constituents of *Cu. reflexa* vary with the type hosts. In the same way, linolenic acid, linoleic acid, oleic acid, stearic acid, palmitic acid, β-sitosterol,

stigmasterol, campesterol and β -amyrin, were reported in the list of chemical constituents isolated from species of *Cuscuta* [4, 7]. Moreover, 4-vinylphenol, which is widely used as a flavoring agent [8], was found as the major component in the extract of *Cu. reflexa* grown on *S. saman* (18.5%). On the other hand, unsaturated fatty acids and methyl esters of fatty acids were found to be the major components of the extract of *Cu. reflexa* grown on *F. racemosa* (46.9%), *Co. grandis* (33.1%) and *S. saman* (31.6%), respectively. The essential fatty acids (linoleic acid and linolenic acid) that cannot be synthesized in the human body and must be obtained from the diet were found in the extracts of *Cu. reflexa*; therefore, this plant can be considered a good source of essential fatty acids. Moreover, linolenic acid has been associated with health benefits in the treatment of several diseases such as coronary artery disease and rheumatoid arthritis [9–10].

Table 2: Chemical compositions of the methanol crude extracts of *Cuscuta reflexa* grown on *Coccinia grandis*, *Ficus racemosa* and *Samanea saman*.

Peak no. (min) ^a	RT	Components ^b	Total peak (%)		
			CCG	CFR	CSS
1	7.7	4-Vinylphenol	13.3	14.0	18.5
2	10.0	4-Vinylguaiacol	4.7	3.3	4.4
3	11.8	Methyl cinnamate	0.2	0.3	0.2
4	21.0	Methyl 4-hydroxycinnamate	0.8	0.6	0.8
5	22.0	E-9-Heptadecenal	3.2	ND	ND
6	22.5	Pentadecanoic acid methyl ester	0.5	ND	ND
7	23.1	Methyl ferulate	0.4	ND	ND
8	24.6	Palmitoleic acid methyl ester	0.4	ND	ND
9	24.9	Palmitic acid methyl ester	12.6	10.1	10.7
10	25.1	Methyl-3-(3,5-diterbutyl-4-hydroxyphenyl) propionate	1.1	0.5	1.0
11	25.8	Palmitic acid	0.5	ND	ND
12	28.3	Linoleic acid methyl ester	8.6	6.0	7.5
13	28.5	α -Linoleic acid methyl ester	9.9	7.5	7.3
14	28.7	Phytol	0.4	0.5	0.8
15	29.0	Steric acid methyl ester	3.7	2.7	3.0
16	30.1	Palmitic acid amide	1.8	2.8	1.7
17	32.5	Arachidic acid methyl ester	1.0	0.6	0.6
18	33.3	Oleic acid amide	14.1	33.4	16.8
19	33.6	Stearic acid amide	1.2	1.8	1.1
20	35.2	Cyclopentadecane	1.0	0.5	1.2
21	35.7	Docosanoic acid methyl ester (behenic)	ND	ND	0.4
22	38.6	Lignoceric acid methyl ester	0.9	0.6	1.0
23	39.8	Squalene	3.2	2.6	2.1
24	40.9	Nanocosane	2.1	5.1	2.4
25	43.0	γ -Tocopherol	0.9	1.0	0.9
26	44.1	α -Tocopherol	1.1	0.7	1.0
27	45.8	Campesterol	1.8	0.9	3.2
28	46.3	Stigmasterol	1.1	0.5	1.6
29	47.2	(24RS)-24,27-dimethylcholesta-5,26-dien-3 β -ol	3.3	1.7	4.9
30	47.5	β -Sitosterol	2.1	0.8	4.4
31	48.3	β -Amyrin	1.2	1.3	2.6
32	49.4	Baccharan-3 β -ol	2.5	ND	ND
Phenolic compounds			20.5	18.7	24.9
Hydrocarbons			6.3	8.2	5.8
Saturated fatty acids and methyl ester of fatty acids			22.2	18.7	18.4
Unsaturated fatty acids and methyl ester of fatty acids			33.1	46.9	31.6
Fatty aldehydes			3.2	ND	ND
Vitamin E			2.1	1.6	1.8
Terpenes			1.6	1.8	3.4
Sterols			10.9	3.9	14.0
Total			99.9	99.8	99.9

^aRetention time (minutes). ^bCompounds listed in order of elution from a HP-5 MS column.

ND = not detected

Host plants: CCG = *Cu. reflexa* grown on *Co. grandis*, CFR = *Cu. reflexa* grown on *F. racemosa*, CSS = *Cu. reflexa* grown on *S. saman*

Additionally, as shown in Table 3, 20 phenolic compounds were identified and quantified by high performance liquid chromatography (HPLC). Gallic acid, catechin, vanillic acid, rutin and quercetin could be found in the *Cu. reflexa* extracts grown on *Co. grandis* ranging from 0.34 to 6.96 mg/g extract of which the greatest one was rutin. The extract of *Cu. reflexa* that was grown on *F. racemosa* contained high contents of catechin (5.81 mg/g extract), quercetin (4.53 mg/g extract) and rutin (3.95 mg/g extract). On the other hand, the extract of *Cu. reflexa* that was grown on *S. saman* contained lower amounts of gallic acid, catechin, vanillic acid, rutin and quercetin than when grown on the other hosts, with contents ranging from 0.34 to 1.65 mg/g extract.

Previous studies reported that rutin and quercetin were usually found in extracts of *Cuscuta* species [5, 8, 11].

Table 3: Phenolic compound contents in the methanol crude extracts of *Cuscuta reflexa* grown on *Coccinia grandis*, *Ficus racemosa* and *Samanea saman*.

Compounds	Retention times [*]	Content (mg/g extract)		
		<i>Co. grandis</i>	<i>F. racemosa</i>	<i>S. saman</i>
Gallic acid	11.7	0.90 \pm 0.02b	1.25 \pm 0.01a	0.35 \pm 0.02c
Catechin	23.1	5.97 \pm 0.05a	5.81 \pm 0.06a	1.14 \pm 0.13b
Vanillic acid	24.7	0.34 \pm 0.02c	1.27 \pm 0.01b	1.64 \pm 0.11a
Rutin	32.2	6.96 \pm 0.04a	3.95 \pm 0.03b	1.44 \pm 0.06c
Quercetin	46.6	0.87 \pm 0.06c	4.53 \pm 0.10a	1.45 \pm 0.03b

HPLC conditions: column Eclipse-C₁₈ (150 \times 4.6 mm, i.d.5 μ M), gradient solution (A) 0.1% formic acid in water and (B) 100% acetonitrile (B. 30%, 0-50 min, 50%, 51-60 min, UV detector 280 nm, flow rate 0.8 mL/min and injection volume 20 μ L).

Values are expressed as means \pm S.D.

The different letters in the same column are considered significantly different according to Duncan's multiple comparison test ($P < 0.05$).

The results of the antidiabetic assay of the *Cu. reflexa* extracts (1.0 mg/mL) based on the α -glucosidase inhibition model showed that the extract of *Cu. reflexa* that was grown on *Co. grandis* exhibited the highest inhibition percentage of 51.2%, followed by *F. racemosa* (45.7%) and *S. saman* (31.2%). Previously, Rahmatullah et al. [12] investigated the hypoglycemic effects of the methanol and chloroform extracts of *Cu. reflexa* based on oral glucose tolerance tests. The results showed that both methanol and chloroform extracts of *Cu. reflexa* demonstrated significant oral hypoglycemic activity in glucose-loaded rats. Previous studies have suggested that the phenolic and flavonoid compounds that are present in plant extracts are responsible for the antidiabetic activity [13–14].

The methanol extracts of *Cu. reflexa* grown on three different hosts were also investigated for their possible antioxidant activities by DPPH free radical (DPPH $^{\bullet}$) scavenging and ferric reducing antioxidant power (FRAP) assays, and the results are displayed in Table 4. In the DPPH $^{\bullet}$ scavenging system, the IC₅₀ of *Cu. reflexa* extracts grown on different hosts varied from 279.7 to 168.6 μ g/mL. The highest activity was noticed in *Cu. reflexa* grown on *Co. grandis*, and this specimen displayed potential antioxidant activity with a statistical difference ($P < 0.05$) when compared with the other hosts. In the FRAP system, the extract of *Cu. reflexa* grown on *Co. grandis* still possessed the highest FRAP value of 40.5 mg GAE/g dry extract), which was significantly higher than that of the extracts from the other hosts ($P < 0.05$). Amol et al. [15] evaluated the antioxidant activities of the methanol extract of *Cu. reflexa* using the DPPH $^{\bullet}$ scavenging and reducing power assays. The extract showed antioxidant activity with an IC₅₀ of 359.5 μ g/mL, and also exhibited high reducing capacity in the reducing power system. Based on our results, the high phenolic and flavonoid contents of the extract of *Cu. reflexa* grown on *Co. grandis* could be responsible for its greater antioxidant activities. This finding has also been suggested in many earlier reports [16–17]. Previous research studies have suggested that the antioxidant activity of plants extracts can be partly attributed to some constituents other than fatty acids, such as phenolic compounds and vitamin E [18–19]. According to the GC-MS results of the extract of *Cu. reflexa* grown on *Co. grandis*, the presence of methyl ferulate and vitamin E, which are already known to possess high antioxidant activities [20–21], may be related to the greater antioxidant activities. Moreover, previous research reports have revealed that phenolic compounds can exhibit high antioxidant activity [22–23], especially rutin, catechin and quercetin, which were found in the extracts of *Cu. reflexa* grown on *Co. grandis* and *F. racemosa*. However, the unidentified active phenolic compounds that present in the methanol crude extracts of *Cu. reflexa* may also play a role in antioxidant activity.

Table 4: Antioxidant activities of the methanol crude extracts of *Cuscuta reflexa* grown on *Coccinia grandis*, *Ficus racemosa* and *Samanea saman*.

Host plant	Antioxidant activity	
	DPPH scavenging activity (IC_{50} µg/mL)	Ferric reducing antioxidant power (mg GAE/g extract)
<i>Coccinia grandis</i>	168.7±1.0a	40.5±0.8a
<i>Ficus racemosa</i>	201.6±0.5b	35.0±1.4b
<i>Samanea saman</i>	279.7±6.0c	26.8±0.3c

*Average ± standard deviation from three replicates.

The different letters in the same column are considered significantly different according to Duncan's multiple comparison test ($P < 0.05$).

Experimental

Plant material: *Cu. reflexa* grown on *Co. grandis*, *F. racemosa* and *S. saman* was collected locally in Suthep District, Muang, Chiang Mai, Thailand. The plants were dried at 45°C, ground into powder and stored at room temperature for further extraction.

Preparation of the extracts: Ten g of dried stem material of *Cu. reflexa* was extracted with 100 mL of methanol, and left overnight at room temperature. The extract was sonicated using an ultrasonicator (Crest, USA) for 30 min, filtered through Whatman no. 1 filter paper and evaporated under vacuum at 40°C using a rotary evaporator until dry. The dry extracts were kept at room temperature in the dark until required.

Determination of TFC: TFC was determined using the method of Kaewnarin *et al.* [24], with slight modification. The extract (0.5 mL) was mixed with 2 mL of methanol, followed by the addition 0.15 mL of 50 g/L NaNO₂. After 5 min, 0.15 mL of 100 g/L AlCl₃ was added and the mixture was incubated at room temperature for 15 min before the absorbance was measured at 415 nm. Quercetin solution was used as a standard for the determination and the results were expressed as mg QE/g dry extract.

GC-MS analysis: Analysis of the extracts of *Cu. reflexa* grown on three different hosts was performed using a 6890 Agilent Technologies/MSD 5973 Hewlett Packard, equipped with a MS detector and an HP-5MS capillary column (bonded and cross-linked 5% phenyl-methylpolysiloxane, 30 m × 0.25 µm, film thickness 0.25 µm). The injector and detector temperatures were set at 270 and 280°C, respectively. The oven temperature was set at 80°C and held for 2 min, and then increased at a rate 10°C/min to 120°C and held for 4 min. The oven temperature was then increased at a rate 10°C/min to 155°C and held for 4 min, and then increased at a rate 5°C/min to 280°C and held for 12.50 min. The total running time was 55 min. Helium was used as carrier gas at a flow rate of 1 mL/min. The sample (1 µL) was injected in the splitless mode. GC-MS detection in an electron ionization system with an ionization energy measurement of 70 eV was used. Injector and MS transfer line temperatures were set at 270 and 290°C, respectively. The components were identified based on a comparison of their relative retention times and the mass spectra with W8N08 and Wiley 7th libraries data of the GC-MS system.

Determination of TPC: TPC was estimated using the protocol of Thitilertdecha *et al.* [25], with slight modification. The procedure involved combining 0.25 mL of sample with 2.5 mL deionized water and 0.5 mL of Folin-Ciocalteu reagent. After 5 min, 0.5 mL of 20%, w/v, Na₂CO₃ was added, and the solution was incubated for 1 h at room temperature. Absorbance was then measured at 760 nm. Gallic acid solution was used as a standard for the determination and the results were expressed as mg GAE/g dry extract.

HPLC analysis: The sample extracts were measured using an HPLC technique (Agilent 1200). An Eclipse-C₁₈ column was used (150 x 4.6 mm, i.d.5 µM). The flow rate was set to 0.8 mL/min. The

mobile phase was 0.1% formic acid in water (solvent A) and 100% acetonitrile (solvent B). Gradient elution was performed as follows: 0-50 min, 30% B and 51-60 min, 50% B. The injection volume was 20 µL. The fingerprint profiles were recorded at an optimized wavelength of 280 nm. The quantification of gallic acid, catechin, vanillic acid, protocatechuic acid, syringic acid, caffeic acid, chlorogenic acid, sinapic acid, ferulic acid, *m*-coumaric acid, *o*-coumaric acid, ellagic acid, rosmarinic acid, rutin, myrecetin, quercetin, kaempferol, apigenin, luteolin and isorhamnetin was determined based on peak area measurements.

Antidiabetic assay: α-Glucosidase (AGH) solution was prepared from rat intestinal acetone powder by partial modification of the procedure reported by Oki *et al.* [26]. One hundred mg of intestinal acetone powder was added to 3 mL of 0.9% NaCl solution, homogenized by sonication and kept in an ice bath. After centrifugation at 6,000 rpm for 30 min at 4°C, the resulting supernatant was kept cold and directly subjected to inhibitory assay. The method of Adisakwattana *et al.* [27] was used to determine AGH inhibitory assay. The assay was defined as the percent inhibition under the assay conditions, which was calculated according to the formula:

$$\text{Percent inhibition} = (A_o - A_s / A_o) \times 100$$

where A_o is the absorbance of the control, and A_s is the absorbance of the mixture containing the test compound.

Determination of antioxidant activities

DPPH[•] scavenging assay: The free radical scavenging ability was determined according to the method of Gülcin *et al.* [28], with slight modification. 2,2'-Diphenyl-1-picrylhydrazyl radical solution in ethanol (0.1 mM, 1.5 mL) was mixed with 0.5 mL of different concentrations of each extract, and methanol was used as the control. The mixtures were well shaken and kept at room temperature for 30 min in the dark. The absorbance was measured at 517 nm. The percent of DPPH[•] discoloration of the samples was calculated according to the formula:

$$\text{Percent inhibition} = (A_o - A_s / A_o) \times 100$$

where A_o is the absorbance of the control (containing all reagents except the test compound), and A_s is the absorbance of the mixture containing the test compound. The test sample concentrations providing 50% inhibition (IC₅₀) were calculated from the plot of inhibition percentage against extract concentration values. The radical scavenging ability was presented as IC₅₀ values.

FRAP assay: The FRAP assay was determined according to the protocol of Li *et al.* [29], with some modification. The FRAP reagent containing 10 mM TPTZ solution in 40 mM hydrochloric acid (20 mL), 20 mM ferric (III) chloride (20 mL) and acetate buffer (5 mL, 300 mM, pH 3.6) was prepared freshly prior to being used. Different concentrations of each extract (0.1 mL) were mixed with the FRAP reagent (1.5 mL) and 1.4 mL of acetate buffer (300 mM, pH 3.6) and then incubated at room temperature for 30 min. The absorbance was measured at 593 nm. Gallic acid was used as a standard and FRAP value was calculated as the gallic acid equivalent (mg GAE/g dry extract).

Statistical analysis: The results of all experiments were expressed as mean ± standard deviation. Analysis of variance was performed by ANOVA procedure and Duncan's multiple comparison test was used to determine any significant differences ($P < 0.05$) identified between treatments.

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Chemical Composition, Antioxidant and Anticholinesterase Activities of *Gentianella azurea* from Russian Federation

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Phytochemical study of *Gentianella azurea* (Bunge) Holub (Gentianaceae) collected in Buryatia Republic (Russian Federation) resulted in the isolation of twenty-one compounds including bellidifolin, bellidin, isobellidifolin, norswertianolin, isobellidifolin-8-O-β-D-glucopyranoside, orientin, cynaroside, cosmoisin, apigenin, 4'-O-caffeoyleswertiamarin, swertiamarin-6'-O-β-D-glucopyranoside and sweroside, firstly detected in this species. The extracts and individual compounds were shown to possess antioxidant and anticholinesterase potential.

Keywords: *Gentianella azurea*, Gentianaceae, Xanthones, Flavones, Iridoids, Antioxidant activity, Anticholinesterase activity, HPLC.

Gentianella azurea (Bunge) Holub (azure little gentian) is a small annual herb, common in the forests of Eastern Siberia, China and Mongolia. The decoction of the total herb (local name *sum cu tig ta*) is used in traditional medicine of Buryatia as a choleretic and antidepressant [1]. Previously, some xanthones, flavonoids, and iridoids were reported [2]. All earlier studies were carried out on plant material collected in Mongolia [2a] and China [2b, c]. Hitherto, there is no data about the biological activities of *G. azurea*. As a part of our ongoing research of the chemical composition and bioactivities of bitter gentian teas [3] this study presents the results of a phytochemical and biological investigation of *G. azurea* growing in Buryatia Republic (Russian Federation).

The 60% MeOH-extract of the total herb was partitioned with CHCl_3 , EtOAc, and *n*-BuOH to yield three fractions, which were separated to give twenty-one compounds including eight xanthones {bellidifolin (**1**), bellidin (**2**), isobellidifolin (**3**) [4], 1,3,5-trihydroxy-8-methoxyxanthone (**4**) [3a], swertianolin (**5**), norswertianolin (**6**), isobellidifolin-8-O-β-D-glucopyranoside (**7**) [4], 1,3,5-trihydroxy-8-methoxyxanthone-1-O-β-D-glucopyranoside (**8**) [3a]}, six flavones {orientin (**9**), isoorientin (**10**) [5], cynaroside (**11**), cosmoisin (**12**), luteolin (**13**), apigenin (**14**) [6]}, six iridoids {gentiopicroside (**15**), swertiamarin (**16**), 4'-O-caffeoyleswertiamarin (**17**) [7], swertiamarin-6'-O-β-D-glucopyranoside (**18**) [8], sweroside (**19**), loganic acid (**20**) [9]}, and one triterpene oleanolic acid (**21**) [3b], which were identified by comparison of their m.ps, UV, FTIR, ¹H and ¹³C NMR spectral, and MS data with those reported in the literature. Compounds **1-3**, **6**, **7**, **9**, **11**, **12**, **14**, **17-19** were isolated from *G. azurea* herb for the first time.

HPLC data showed that the flowers and leaves of *G. azurea* accumulated xanthones, flavonoids and iridoids (Table 1). The dominant xanthones of the flowers and leaves were **8** and **5**. The prevailing flavonoid (**9**) concentrated in the leaves. Iridoids were the most significant group of extracted compounds, and **16** was found in higher levels in the flowers and leaves. A characteristic feature of the roots was their ability to concentrate **15**, with small

Table 1: Content of xanthones (Xt), flavonoids (Fl) and iridoids (Ir) in *G. azurea* organs^a, total herb (GaH1-GaH3) and MeOH-extract (GaM)^b.

Comps	Fl	L	St	R	GaH1	GaH2	GaH3	GaM
1	0.4	0.2	tr. ^c	0.1	0.6	0.3	0.4	0.9
2	1.5	0.3	0.3	tr.	2.9	4.6	0.7	12.5
3	0.1	0.1	n.d. ^d	n.d.	0.3	0.4	tr.	1.1
4	0.4	0.4	0.4	0.5	0.5	0.2	0.5	0.6
5	6.1	7.3	1.0	0.5	2.4	1.3	6.4	3.5
6	4.7	5.7	2.0	n.d.	5.1	2.9	5.6	8.0
8	7.1	6.7	3.1	0.9	7.1	5.7	8.2	15.3
9	14.6	20.7	4.3	n.d.	17.9	11.8	19.4	31.6
10	2.4	2.1	0.3	n.d.	3.3	2.7	5.0	7.3
11	7.3	2.4	0.6	n.d.	4.0	2.5	2.6	6.7
12	tr.	0.9	tr.	n.d.	0.6	0.9	0.5	2.4
15	tr.	2.2	3.1	16.0	2.8	2.5	5.2	6.7
16	154.8	121.5	38.2	2.4	107.6	73.6	113.5	197.8
19	31.7	3.8	6.1	0.5	27.6	24.1	37.0	62.7
20	4.0	2.4	4.3	0.6	4.6	4.6	5.9	11.9
ΣXt	20.3	20.7	6.8	2.0	18.9	15.4	21.8	41.9
ΣFl	24.3	26.1	5.2	n.d.	25.8	17.9	27.5	48.0
ΣIr	190.5	129.9	51.7	19.5	142.6	104.8	161.6	279.1

^aFl – flowers, L – leaves, St – stems, R – roots; ^bmg/g, dry plant (extract) weight; ^ctraces (<LOQ); ^dn.d. not detected (<LOD).

concentrations of xanthones and flavonoids. The total content of xanthones, flavonoids and iridoids in *G. azurea* herbs collected in three regions of Buryatia Republic were 15.4–21.8, 17.9–27.5 and 104.8–161.6 mg/g respectively (Table 1). Four components (**8**, **9**, **16** and **19**) were prevalent in all herb samples.

Table 2: Radical-scavenging (DPPH) and anticholinesterase activity (ACE) of *G. azurea* MeOH-extract (GaM) and individual compounds^a.

Compds	DPPH ^a	ACE ^b	Compds	DPPH ^a	ACE ^b
GaM	39.8	22.9	12	>100	>100
1	5.5	1.2	13	10.6	4.96
2	3.4	5.3	14	>100	>100
3	6.3	2.9	15	>100	>100
4	7.1	44.2	16	>100	>100
5	14.1	9.2	17	52.4	>100
6	12.6	47.5	18	>100	>100
7	21.7	20.6	19	>100	>100
8	25.2	>100	20	>100	>100
9	13.8	6.8	21	>100	22.9
10	19.7	28.5	Trolox ^b	11.6	-
11	14.5	35.1	Neostigmine ^b	-	0.91

^aIC₅₀, μg/mL; ^breference compound.

The IC₅₀ value for the radical-scavenging activity of the MeOH-extract (GaM) against DPPH[•]-radicals was 32.3 µg/mL. The most active compounds were xanthone aglycones (**1–4**) (Table 2). The acetylcholinesterase-inhibiting activity of GaM was characterized by an IC₅₀ value 22.9 µg/mL. Comparative analysis of individual compounds showed that xanthone aglycones **1** and **3** with a methoxy-group in positions C-3 and C-5, respectively, were more effective inhibitors than **2** and **4** (Table 2). It should be noted that the activities of flavone **13** and C-glucoside **9** were close to that of xanthone aglycones.

In conclusion, the results obtained confirmed the data of the ethnopharmacological use of *G. azurea* as a natural phytotherapeutic agent.

Experimental

General experimental procedures: UV-Vis, SF-2000 UV-Vis-spectrophotometer (OKB Specter), Uniplan microplate UV-Vis-spectrophotometer (Pikon); Elemental composition, MAT 8200 spectrometer (Thermo Finnigan); IR, FT-801 FTIR spectrophotometer (Simex); Melting point, IA9100 micromelting point apparatus (Electrothermal Thermo Scientific); NMR, VXR 500S spectrometer (Varian); ESIMS (positive mode), Thermo Finnigan LCQ mass spectrometer; CC, Silica gel 60 (Merck, 70-230 mesh), Amberlite XAD7HP (Sigma), Woelm polyamide; GPC, Sephadex LH-20 (25-100 µm; Pharmacia); prep. HPLC-UV, Summit system with UV-Vis detector (Dionex), LiChrosorb RP-18 (4.6×250 mm, 5 µm, Merck), T 35°C, flow rate 1 mL/min; solvent, linear gradient of 5-70% of B (MeCN) in A (H₂O) for 90 min; detector 254 nm; analytical HPLC, Milichrom A-02 system (Econova), as described previously [3b]; Biological assays, radical-scavenging activity – microplate assay [10], acetylcholinesterase inhibition activity – microplate assay [11].

Plant material: The samples of *G. azurea* were collected in the flowering period in the Buryatia Republic (Russian Federation), dried and stored at 4°C in the PlaMeta Group Plant Repository: Nur-Tukhum village (No GaH1; Dzhidinskii region, 15.VIII.2012,

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50°73'31" N, 106°25'98" E; voucher specimen No Gn/h-31/01-64/0812); Tsakir village (No GaH2; Zakameneskii region, 20.VIII.2012, 50°49'26" N, 103°57'36" E; voucher specimen No Gn/h-36/06-82/0812); Turka village (No GaH3; and Pribaikal'skii region, 22.VIII.2012, 52°94'57" N, 108°24'43" E; voucher specimen No Gn/h-54/42-104/0812).

Extraction and isolation: Air-dried, ground herb of *G. azurea* (sample GaH3; 1.7 kg) was extracted with 70% EtOH (60°C, ×3), and the combined extracts were partitioned with CHCl₃ (Ga-F₁, 108.3 g), EtOAc (Ga-F₂, 159.8 g) and *n*-BuOH (Ga-F₃, 283.9 g) respectively. The Ga-F₁ fr. (60 g) was chromatographed using flash chromatography on a silica column [3×100 cm, CHCl₃-MeOH (100:0→0:100)], and by prep. HPLC to yield **1** (63 mg), **2** (82 mg), **3** (22 mg), **4** (18 mg), **13** (12 mg), **14** (8 mg) and **21** (3.35 g). The Ga-F₂ fr. (80 g) was loaded onto Amberlite XAD7HP (400 g) and eluted with a water-MeOH mixt. (100:0→0:100) to give 6 sub-fractions Ga-F₂-1–Ga-F₂-6. Subfr. Ga-F₂-1 was chromatographed by prep. HPLC to yield **17** (12 mg) and **18** (9 mg). Subfr. Ga-F₂-2, Ga-F₂-3 was chromatographed on a polyamide column [2×50 cm, H₂O-MeOH (100:0→0:100)] and a Sephadex LH-20 column [3×100 cm, MeOH-H₂O (100:0→50:50)] to yield **9** (512 mg), **10** (84 mg), **11** (16 mg) and **12** (6 mg). Subfr. Ga-F₂-4 – Ga-F₂-6 was chromatographed on a silica column [2×100 cm, CHCl₃-MeOH (100:0→60:40)], a polyamide column [2×50 cm, H₂O-MeOH (100:0→0:100)] and a Sephadex LH-20 column [3×100 cm, MeOH-H₂O (100:0→50:50)] to yield **5** (87 mg), **6** (245 mg), **7** (12 mg) and **8** (87 mg). The Ga-F₃ fr. (100 g) was loaded onto Amberlite XAD7HP (600 g) and eluted with a water-MeOH mixt. (100:0→0:100) to give 6 sub-fractions Ga-F₃-1–Ga-F₃-6. Subfr. Ga-F₃-1 was chromatographed on a polyamide column [2×70 cm, H₂O-MeOH (100:0→0:100)] and by prep. HPLC to yield **15** (15 mg), **16** (2.63 g), **19** (163 mg) and **20** (32 mg).

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Constituents of Fermented Male Flowers of *Alnus sieboldiana* (Betulaceae)

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Fractionation of the ethyl acetate crude extract of the fermented male flowers of *Alnus sieboldiana* resulted in the isolation of two diarylheptanoids and six flavonoids. Yashabushidiol A, yashabushidiol B and naringenin were known constituents of the male flowers of *A. sieboldiana* while kaempferol, quercetin, pinocembrin dimethyl ether, 5,7-dimethoxy-3-hydroxyflavanone and 5,7-dimethoxy-3-hydroxyflavone were isolated for first time from the fermented male flowers of *A. sieboldiana*. The isolated compounds were characterized using NMR and HRMS and compared with available literature.

Keywords: *Alnus sieboldiana*, Betulaceae, Biotransformation, Flavones, Flavanones, 1,7-Diarylheptanoids.

The genus *Alnus* (family Betulaceae) contains 35 deciduous tree and shrub species. The plants are widely distributed in temperate regions of the northern hemisphere and can be found in wetlands, volcanic soils, glacial deposits and sand dunes [1].

A. sieboldiana Matsum. is a deciduous plant found on the south western coast of Japan. The roots contain *Azotobacter* species, which are involved in nitrogen fixation [2]. Therefore, this species can be planted in poorly nourished mountain and coastal areas to protect them from erosion. The male and female flowers emit a strong aromatic and resinous odor. *A. sieboldiana* has been of special interest to Asakawa's research group who studied the plant over three decades (1969-1995). The male flowers were reported to contain β-phenylethyl cinnamate, stilbene monomers, flavanones, flavones, 1,7-phenylheptanoids and some sesquiterpenoids [3]. The flavone galangin, isolated from the methanol extract of the leaves of *A. sieboldiana*, has been found to inhibit the A549 tumor cell line [4]. It is interesting to note that the whole plant, including the flowers and young leaves, even the cones, are easily infected by the white-colored fungus *Penicillium* sp. [5], causing the powerful aromatic odor of both male and female flowers to diminish dramatically and finally turn unpleasant. The powerful aromatic odor was due to the presence of 2-phenylethanol [6].

The present paper concerns a phytochemical study of fermented *A. sieboldiana* male flowers. From the ethyl acetate extract of the 6-month-old fermented male flowers, two diarylheptanoids and six flavonoids were successfully isolated. Of these isolated compounds, the diarylheptanoids yashabushidiol A (1) [7a] and yashabushidiol B (2) [7], and the flavonoid naringenin (3) [4a, 8] are known constituents of the original male flowers, while the other five isolated flavonoids kaempferol (4), quercetin (5), pinocembrin dimethyl ether (7), 5,7-dimethoxy-3-hydroxyflavanone (8) and 5,7-dimethoxy-3-hydroxyflavone (9) are biotransformation products, resulting from the enzymatic actions of the endophytic white-colour fungus.

Naringenin (3) was previously isolated from fresh male flowers, but in this experiment was only isolated in a minute amount. In



Figure 1: Two diarylheptanoids from the male flowers of *A. sieboldiana*.

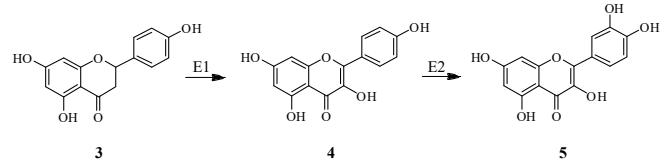


Figure 2: Metabolic pathway of 4 and 5, from naringenin (3).

addition, kaempferol (4) [9] and quercetin (5) [10] were successfully isolated from the fermented male flowers for the first time. Naringenin (3) is biosynthesized in the male flowers of *A. sieboldiana* through the shikimate acid pathway [11a]. Flavonoids 4 and 5 isolated from the fermented male flowers in this work are hydroxylated derivatives of naringenin (3), probably resulting from biotransformation by flavanone 3-hydroxylase (E1) and flavonoid 3'-hydroxylase (E2) enzymes [11b], as proposed in Figure 2.

In addition, three 5,7-dimethoxyflavonoids, 7, 8 and 9, were also isolated from the ethyl acetate crude extract of the fermented male flowers. These compounds have never been reported in the original flowers. We herein report these compounds as biotransformation products in the fermented male flowers of *A. sieboldiana*.

Compound 7 may be produced from pinocembrin (6), which is a known constituent of the male flowers of *A. sieboldiana* [4c], through methylation by plant-O-methyltransferases (OMTs) [11c]. Compound 7 is converted to compound 8 by flavanone 3-hydroxylase (E3) [11d]. Further oxidation of compound 8 led to compound 9 as the final product. In this work, pinocembrin was not detected; presumably it was fully converted into the derivatives. The proposed metabolic pathway of compounds 7, 8 and 9 are illustrated in Figure 3.

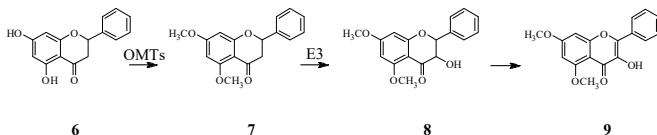


Figure 3: Metabolic pathway of 7, 8 and 9, from pinocembrin (6).

In conclusion, fermentation of *A. sieboldiana* male flowers by the white fungus *Penicillium* sp. caused hydroxylation at positions 3 and 3' of naringenin (**3**) to produce **4** and **5**, while methylation at positions 5 and 7 of pinocembrin (**6**) gave **7**. Further hydroxylation and oxidation of **7** produced **8** and **9**.

Experimental

Plant material: *Alnus sieboldiana* (male flowers) were collected in February 2015 at Naruto (Shikoku Island), Japan at the beginning of spring. The flowers were identified by YA and deposited at the herbarium of Tokushima Bunri University.

The 6-month-old fermented male flowers (0.97 kg) were mechanically crushed and extracted with ethyl acetate. The extract was evaporated to give a viscous extract (49.64 g), which was subjected to VLC (8 cm x 12 cm) eluting with mixtures of *n*-hexane / ethyl acetate (10:0; 9:1; 8:2; 7:3; 6:4; 5:5; 7.5:2.5; 0:10) to obtain 12 fractions. Yashabushidiol B (**2**) was obtained as white crystals upon evaporation of fraction 7 (1.37 g). Fraction 8 was further separated using Sephadex LH-20 eluting with a mixture of dichloromethane:methanol (1:1) giving yashabushidiol A (**1**) as white-yellowish crystals (944.3 mg), and naringenin (**3**) (1.8 mg). Further treatment of fraction 9 using Sephadex, followed by normal phase silica gel column chromatography eluting with a combination of *n*-hexane and ethyl acetate gave kaempferol (**4**) (9.0 mg),

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pinocembrin dimethyl ether (**7**) (1.5 mg), 5,7-dimethoxy-3-hydroxyflavanone (**8**) (2.0 mg) and 5,7-dimethoxy-3-hydroxyflavone (**9**) (2.7 mg). Lastly, quercetin (**5**) (82.88 mg) was obtained as yellow powder isolated from the tenth fraction.

5,7-Dimethoxy-3-hydroxyflavanone (**8**)

Yellow amorphous.

¹H NMR (CDCl₃, 500 MHz): 4.45 (1H, dd, *J* = 0.8, 12.2 Hz, H-2), 5.04 (1H, d, *J* = 12.2 Hz, H-3), 6.13 (2H, dd, *J* = 1.5, 6.5 Hz, H-6, H-8), 7.57 (2H, dd, *J* = 1.5, 7.4 Hz, H-2', H-6'), 7.44 (3H, m, H-3', H-4', H-5'), 3.82 (3H, s, OCH₃-7), 3.92 (3H, s, OCH₃-5).

¹³C NMR (125 Hz): 72.7 (C-2), 83.3 (C-3), 190.8 (C-4), 162.1 (C-5), 93.3 (C-6), 167.0 (C-7), 93.8 (C-8), 164.9 (C-9), 102.9 (C-10), 136.5 (C-1'), 127.5 (C-2', C-6'), 128.7 (C-3', C-5'), 129.2 (C-4'), 56.2 (OCH₃-5), 55.7 (OCH₃-7).

MS *m/z*: 300.0997 C₁₇H₁₆O₅.

5,7-Dimethoxy-3-hydroxyflavone (**9**)

Light yellow powder.

¹H NMR (CDCl₃, 500 MHz): 6.36 (1H, *d*, *J* = 2.15 Hz, H-6), 6.58 (1H, *d*, *J* = 2.2 Hz, H-8), 8.20 (2H, *dd*, *J* = 1.35, 7.9 Hz, H-2', H-6'), 7.52 (2H, *dd*, *J* = 7.9, 7.5 Hz, H-3', H-5'), 7.45 (1H, *m*, H-4'), 3.93 (3H, s, OCH₃-5), 3.99 (3H, s, OCH₃-7).

¹³C NMR (125 Hz): 141.8 (C-2), 138.3 (C-3), 172.1 (C-4), 160.6 (C-5), 95.7 (C-6), 164.5 (C-7), 92.4 (C-8), 159.1 (C-9), 106.3 (C-10), 131.1 (C-1'), 127.1 (C-2', C-6'), 128.5 (C-3', C-5'), 129.6 (C-4'), 55.8 (OCH₃-5), 56.4 (OCH₃-7).

MS *m/z*: 298.0850 C₁₇H₁₄O₅.

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Secondary Metabolites from the Leaves of *Digitalis viridiflora*

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A new phenylethanoid glycoside, named digiviridifloroside (**1**), was isolated from the leaves of *Digitalis viridiflora* Lindley along with a known phenylethanoid glycoside, calceolarioside A (**2**), two flavonoid glycosides, scutellarein 7-*O*- β -D-glucopyranoside (**3**) and hispidulin 7-*O*- β -D-glucopyranoside (**4**), two cleroidicins, cleroidicins B (**5**) and F (**6**), a nucleoside, adenosine (**7**), as well as a mixture of β -glucopyranosyl-(1 \rightarrow 6)-4-*O*-caffeyl- α / β -glucopyranose and 3,4-dihydroxyphenylethanol. The structure of the new compound was established as 3,4-dihydroxy- β -phenylethoxy-6-*O*(*E*)-feruloyl- β -glucopyranosyl-(1 \rightarrow 6)-4-*O*(*E*)-caffeyl- β -glucopyranoside (**1**) based on extensive 1D- and 2D-NMR spectroscopy, as well as HR-ESI-MS. Digiviridifloroside represents a rare type of phenylethanoid glycoside which bears two aromatic acyl units in its structure. In addition to phytochemical studies, the isolates were evaluated for their *in vitro* antimicrobial activities against three pathogenic bacteria and three yeast strains using a microdilution method. Among the tested compounds, **5** exhibited moderate antibacterial activity against *Bacillus cereus* NRRLB 3711 with a MIC value of 25 μ g/mL, whereas compounds **5** and **6** showed relatively high anticandidal activity against *Candida* strains with MIC values down to 12.5 μ g/mL, in comparison to the standard antimicrobial compounds.

Keywords: *Digitalis viridiflora*, Plantaginaceae, Phenylethanoid glycoside, Digiviridifloroside, Antimicrobial activity.

The genus *Digitalis* (Plantaginaceae) contains biennial or perennial species. It is represented by nine species in the flora of Turkey including *D. viridiflora* Lindley [1]. Previous studies on the genus showed the presence of a wide range of secondary metabolites including phenylethanoid glycosides, cardiac glycosides, steroid saponins, pregnane glycosides, clerodendrins, flavonoids and anthraquinones [2-6]. In continuation of our systematic survey on the phytochemical composition of *Digitalis* species from Turkey, five phenylethanoid glycosides were recently reported from the initial work on *D. viridiflora* [7]. Further detailed chromatographic studies on the chemical constituents of the leaves of *D. viridiflora* led to the isolation of one new (**1**) and one known phenylethanoid glycoside, two flavonoid glycosides, two clerodendrins and a nucleoside (Figure 1). This paper reports the isolation, structure elucidation and antimicrobial activities of these compounds.

Compound **1** was obtained as a yellowish amorphous powder. Its UV and IR spectra were characteristic for a phenylethanoid glycoside. It possesses a molecular formula of $C_{39}H_{44}O_{19}$, as determined by the analysis of its HRESIMS (m/z 839.2390 [$M + Na^+$], calcd for $C_{39}H_{44}NaO_{19}$, 839.2375) and ^{13}C NMR data (Table 1). The 1H NMR spectrum (Table 1) of **1** showed resonances at δ_H 7.55 and 6.25 (each d, $J = 15.8$ Hz) as an *AX* system as well as signals at δ_H 7.00 (d, $J = 2.0$ Hz), 6.86 (dd, $J = 8.1, 2.0$ Hz) and 6.74 (d, $J = 8.1$ Hz) as an *ABX* system attributable to an (*E*)-caffeyl moiety. Moreover, the spectrum also contained three aromatic signals as an *ABX* system at δ_H 6.70 (d, $J = 1.9$ Hz), 6.67 (d, $J = 8.0$ Hz), and 6.55 (dd, $J = 8.0$ and 1.9 Hz), two geminal benzylic methylene signals at δ_H 2.78 (t, $J = 8.0$ Hz), and two nonequivalent oxymethylene signals δ_H 4.01 (m) and 3.71 arising from a

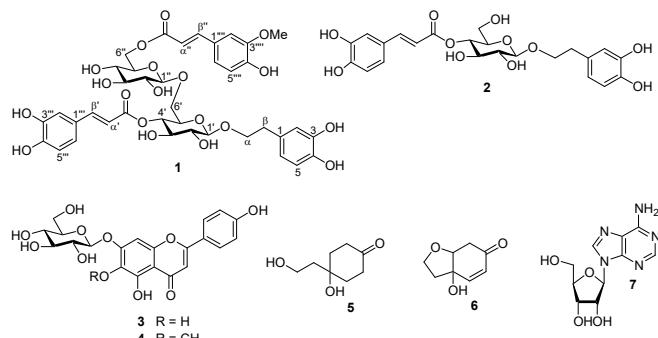


Figure 1: Chemical structures of compounds 1-7.

3,4-dihydroxyphenylethanol moiety. Furthermore, the presence of two anomeric signals at δ_H 4.37 (d, $J = 7.8$ Hz) and 4.35 (d, $J = 7.7$ Hz) revealed the diglycosidic structure of **1**, which was confirmed by the corresponding anomeric carbon resonances at δ_C 105.0 and 104.5 in the ^{13}C NMR spectrum. These findings taken together with 2D NMR experiments (COSY, HSQC and HMBC (Figure 2) revealed the presence of a lugrandoside [2] backbone in **1**. However, the 1H NMR spectrum of **1** contained additional signals at δ_H 7.18 (d, $J = 2.0$ Hz), 7.05 (dd, $J = 8.0, 2.0$) and 6.80 (d, $J = 8.0$ Hz) as an *ABX* type, a pair of *trans*-coupled *AX* type signals at δ_H 7.62 and 6.37 (each d, $J = 15.9$ Hz) and a methoxy signal at δ_H 3.87 (s) suggesting the presence of a (*E*)-feruloyl unit in the structure of **1**.

The deshielded H₂-6'' (δ_H 4.53 and 4.26) and C-6'' (δ_C 64.6) signals of the terminal β -glucopyranose signals indicated that the (E)-feruloyl unit was located at C-6''(OH), which was further

Table 1.¹³C and ¹H NMR data^a for digiviridifloroside (**1**) (CD₃OD, ¹³C 125 MHz; ¹H 500 MHz).

Position	δ_c (ppm)	δ_h (ppm, <i>J</i> in Hz)
Aglycone		
1	131.6	-
2	117.2	6.70 (d, <i>J</i> = 1.9)
3	146.2	-
4	144.7	-
5	116.6	6.67 (d, <i>J</i> = 8.0)
6	121.5	6.55 (dd, <i>J</i> = 8.0, 1.9)
α	72.5	4.01 (m) / 3.71†
β	36.7	2.78 (t, <i>J</i> = 8.0)
Glucose		
1'	104.5	4.37 (d, <i>J</i> = 7.8)
2'	75.4	3.28 †
3'	75.8	3.62 (t, <i>J</i> = 8.0)
4'	72.8	4.83 †
5'	75.1	3.77 (m)
6'	70.3	3.85 †
Glucose		3.69 (dd, <i>J</i> = 11.5, 4.7)
1''	105.0	4.35 (d, <i>J</i> = 7.7)
2''	74.9	3.23 (t, <i>J</i> = 8.2)
3''	77.7	3.35†
4'''	77.4	3.33†
5''	75.7	3.48 (m)
6''	64.6	4.53 (dd, <i>J</i> = 11.7, 1.8) 4.26 (dd, <i>J</i> = 11.7, 5.8)
Caffeoyl		
1'''	127.8	-
2'''	115.4	7.00 (d, <i>J</i> = 2.0)
3'''	146.9	-
4'''	149.8	-
5'''	116.5	6.74 (d, <i>J</i> = 8.1)
6'''	123.2	6.86 (dd, <i>J</i> = 8.1, 2.0)
α'	114.8	6.25 (d, <i>J</i> = 15.8)
β'	147.8	7.55 (d, <i>J</i> = 15.8)
C=O	168.7	-
Feruloyl		
1'''	127.8	-
2'''	111.8	7.18 (d, <i>J</i> = 2.0)
3'''	150.8	-
4'''	149.5	-
5'''	116.8	6.80 (d, <i>J</i> = 8.0)
6'''	124.5	7.05 (dd, <i>J</i> = 8.0, 2.0)
α''	115.4	6.37 (d, <i>J</i> = 15.9)
β''	147.3	7.62 (d, <i>J</i> = 15.9)
C=O	169.2	-
OMe	56.6	3.87 s

^aAssignments are based on COSY, HSQC and HMBC experiments. † Overlapped.

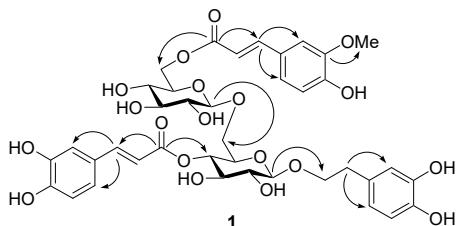


Figure 2: Key HMBC (C-H) correlations for **1**.

confirmed by the long-range correlation of the carbonyl carbon (δ_c 169.2) of the (*E*)-feruloyl unit with H₂-6'' of the terminal glucopyranose unit in the HMBC spectrum (Figure 2). Based on these spectroscopic data, the structure of **1** was elucidated as 2-(3,4-dihydroxyphenyl)ethyl-*O*-6-*O*-(*E*)-feruloyl- β -glucopyranosyl-(1 \rightarrow 6)-4-*O*-(*E*)-caffeoylel- β -glucopyranoside, and named digiviridifloroside.

The known compounds were characterized as calceolarioside A (**2**) [8], scutellarein 7-*O*- β -D-glucopyranoside (**3**) [9], hispidulin 7-*O*- β -D-glucopyranoside (**4**) [10], cleroidicins B (**5**) and F (**6**) [11], and adenosine (**7**) [12] by comparing their spectroscopic data with those published previously. Moreover, a mixture of β -glucopyranosyl-(1 \rightarrow 6)-4-*O*-caffeoylel- α / β -glucopyranose and 3,4-dihydroxyphenylethanol, which could be an artefact formed during the isolation procedure, was characterized.

To the best of our knowledge, digiviridifloroside (**1**) is the third example of a rare phenylethanoid glycoside obtained from the genus *Digitalis*, which contains two aromatic acyl units in its structure; the first two such compounds were reported from *D. lanata* [4]. In a very recent study by Skhirtladze *et al.* [13], another new phenylethanoid glycoside esterified with two aromatic acids was reported. Therefore, the occurrence of such rare phenylethanoid glycosides might possess significant chemotaxonomic importance for the genus *Digitalis* within its new family Plantaginaceae.

The *in vitro* antimicrobial activities of the isolates (except for the mixture) were evaluated against three pathogenic bacteria (*Bacillus cereus* NRRLB 3711, *Pseudomonas aeruginosa* ATCC 10145, *Staphylococcus aureus* ATCC 6538) and three yeast (*Candida albicans* ATCC 90028, *C. parapsilosis* ATCC 22019, and *C. krusei* ATCC 6258) strains using a microdilution method. Compounds **4** and **5** displayed moderate activity against *Bacillus cereus* NRRLB 3711 with MIC values 50 and 25 μ g/mL respectively, while the rest were inactive against the tested bacteria (Table 2). Compounds **4** - **6** displayed moderate activity against all *Candida* strains tested with MIC values ranging from 12.5 to 100 μ g/mL, being **6** the most potent one against *C. parapsilosis* ATCC 22019. To the best of our knowledge the antimicrobial activities of the cleroidicins (**5** and **6**) are being reported for the first time in this study.

Table 2: Antimicrobial activities (MIC, μ g/mL) of compounds **1**-**7**.

Comp.	<i>B. cereus</i> NRRLB 3711	<i>P. aeruginosa</i> ATCC 10145	<i>S. aureus</i> ATCC 6538	<i>C. albicans</i> ATCC 90028	<i>C. parapsilosis</i> ATCC 22019	<i>C. krusei</i> ATCC 6258
1	>100	>100	>100	>100	>100	100
2	>100	100	>100	>100	>100	100
3	>100	>100	>100	>100	>100	>100
4	50	>100	>100	50	100	50
5	25	>100	>100	25	50	25
6	>100	>100	>100	>100	12.5	25
7	>100	>100	>100	>100	>100	>100
S1	-	-	-	0.031	0.062	0.125
S2	0.002	0.062	0.001	-	-	-

S1: Amphotericin B, S2: Chloramphenicol.

Experimental

General experimental procedures: Optical rotation ($[\alpha]_D^{26}$) was measured on a Perkin-Elmer 341 polarimeter. UV spectra and IR spectra were recorded on a HP Agilent 8453 spectrophotometer and a Perkin-Elmer 2000 FT-IR spectrometer, respectively. NMR experiments were performed on a Bruker Avance DRX 500 instrument. COSY, HSQC and HMBC experiments were run under standard conditions at 300 K, dissolving each sample in 550 μ L of 99.8% D CD₃OD (VWR) (¹H, δ = 3.34 ppm; ¹³C, δ = 49.0 ppm). A Q Exactive orbitrap from Thermo Scientific with a HESI ion source was used for HRMS analysis. TLC analyses were carried out on silica gel 60 F₂₅₄ precoated plates (Merck, Darmstadt), and the compounds were stained with 1% vanillin/H₂SO₄ and heating at 105°C. for 1-2 min. For medium-pressure liquid chromatographic (MPLC) separations, Sepacore® Flash Systems X10 / X50 (Büchi) was used with Redi sep columns packed with LiChroprep C₁₈ (13, 43 and 130 g, Teledyne Isco) and SiO₂ (40 g, Teledyne Isco). Open column chromatography (CC) was performed using Silica gel 60 (0.063-0.200 mm; Merck, Darmstadt), polyamide and Sephadex LH-20 (Sigma-Aldrich, St. Louis, MO, USA).

Plant material: The leaves of *Digitalis viridiflora* Lindley were collected from Demirköy, Kırklareli, Turkey, in July 2012 and authenticated by Dr. H. Kırmızıbekmez. A voucher specimen (YEF 12012) has been deposited at the Herbarium of the Department of Pharmacognosy, Faculty of Pharmacy, Yeditepe University, İstanbul, Turkey.

Extraction and isolation: The air-dried and powdered leaves of *D. viridiflora* (185 g) were extracted with MeOH (2 L x 2) at 45°C for 4 h. The solvent was evaporated *in vacuo* to afford the crude MeOH extract (45.1 g, yield 24.3%), which was suspended in H₂O (100 mL) and partitioned with CHCl₃ (100 mL x 3). The H₂O sub-extract (35 g) was subjected to a polyamide column (120 g) eluting with a gradient solvent system of H₂O/MeOH (100:0 to 0:100) to give 7 main fractions, A-G [7]. Fr. B (12.1 g, eluted with 20% MeOH) was applied to silica gel (150 g) CC eluting with a CH₂Cl₂/MeOH/H₂O gradient (90:10:1 to 50:40:10) to obtain 4 sub-fractions, B₁₋₄. Purification of sub-fraction B₁ (340 mg) by medium pressure liquid chromatography (SiO₂, 40 g) eluting with a stepwise CH₂Cl₂/MeOH/H₂O gradient (100:0 to 70:30) gave compounds **6** (50 mg) and **5** (7 mg). Repeated chromatography of sub-fraction B₄ (610 mg) by C₁₈-medium pressure liquid chromatography (C₁₈-MPLC, 43 g, using a H₂O/MeOH gradient, 90:10 to 30:70) and Sephadex LH-20 CC (10 g, MeOH), respectively, gave **7** (2 mg). Fraction D (656 mg, eluted with 40% MeOH) was submitted to C₁₈-MPLC (130 g) eluting with a H₂O/MeOH gradient (85:15 to 35:65) to yield a mixture of β-glucopyranosyl-(1→6)-4-O-caffeoyle-α/β-glucopyranose and 3,4-dihydroxyphenylethanol (11 mg). Fraction G (380 mg, eluted with 100% MeOH) was applied to Sephadex LH-20 CC (60 g) eluting with MeOH to give sub-fraction G₁ as well as **3** (24 mg). Purification of sub-fraction G₁ (65 mg) by C₁₈-MPLC (13 g) eluting with H₂O/MeOH mixtures (85:15 to 0:100) yielded **2** (4 mg), **4** (4 mg) and **1** (15 mg).

Digiviridifloroside (1)

[α]_D²⁶: -59 (*c* 0.1, MeOH).

IR (KBr): 3383, 1698, 1630, 1604, 1515, 1462 cm⁻¹.

UV/Vis λ_{max} (MeOH) nm: 219, 288 (sh), 328.

HR-MS-ESI: *m/z* [M + Na⁺] calcd. for C₃₉H₄₄NaO₁₉: 839.2375; found: 839.2390.

¹H NMR (500 MHz, CD₃OD) and ¹³C NMR (125 MHz, CD₃OD): Table 1.

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Antimicrobial activity assay: A micro-dilution broth susceptibility assay was used, as previously described [14,15]. All microorganisms were stored at -85°C in 15% glycerol prior to the experiments. The bacteria were refreshed on Mueller Hinton agar (MHA, Mast Diagnostics, U.K.), whereas the *Candida* strains were refreshed on Potato Dextrose Agar (PDA, Merck) plates at 37°C. Thereafter, the bacterial suspensions were grown overnight in Mueller-Hinton broth (MHB, Merck, Germany) and were standardized to 1 x 10⁸ CFU/mL versus McFarland No: 0.5 in Mueller-Hinton broth (MHB, Merck, Germany), turbidimetrically. Also, *Candida* strains were inoculated, and standardized in the same way, however in sterile saline (% 0.85) to 5 x 10³ CFU/ per well in RPMI medium (Sigma-Aldrich). Stock solutions of the test samples were prepared in dimethylsulfoxide (DMSO). Dilution series were prepared from 0.6-100 µg/mL accordingly in Mueller Hinton Broth (MHB, Merck) for bacteria and RPMI medium for *Candida* strains in 96-well microtiter plates. Each bacterial (10 µL) and fungal suspension (100 µL) was then added to each well. The last row containing medium with microorganism was used as negative control and medium served as a positive growth control. After incubation at 37°C for 24 h, for staining of viable microorganisms, 0.01% resazurin (20 µL) was added to all of the plates. The first blue well was determined as the minimal inhibitory concentration (MIC, µg/mL). Amphotericin B and chloramphenicol (Sigma, Germany) were used as standard antimicrobial agents at a concentration range of 0.125-64 µg/mL. All experiments were repeated in triplicate and average MICs are given in Table 2.

Supplementary data: HR-MS, ¹H and ¹³C NMR, COSY, HSQC, HMBC spectra of the new compound 1.

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(±)-Perforison A, A Pair of New Chromone Enantiomers from *Harrisonia perforata*

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Dedicated to Prof. Lisheng Ding on the occasion of his 60th birthday.

(+)-Perforison A and (-)-perforison A, a new pair of chromone enantiomers, along with four known compounds, were isolated from the leaves and stems of *Harrisonia perforata*. Their structures and absolute configurations were determined on the basis of extensive analysis of spectroscopic data and electronic circular dichroism (ECD) calculations. The cytotoxic activities *in vitro* of these compounds were evaluated, but none showed significant activity.

Keywords: *Harrisonia perforata*, Simaroubaceae, Chromone, Electronic circular dichroism, Perforison A.

The genus *Harrisonia* (Simaroubaceae) comprises four species and is mainly distributed in south-east Asia, Africa and Oceania [1]. *H. perforata* (Blanko) Merr. is the only species growing in China [1], and has been used as a folk medicine against malaria and boils [2]. Several chromones [3-6], limonoids [4, 7-11], quassinoids [12, 13], and polyketides [14] have been isolated from *H. perforata*. As part of our effort to search for biologically active constituents of this species [7, 12, 15], a new pair of chromone enantiomers, perforisone A, along with four known compounds, peucenin-7-methyl ether (**2**) [16], umtatin (**3**) [17], greveichromenol (**4**) [17], and saikochromone A (**5**) [18], were isolated from the leaves and stems of the title plant. Perforisone A was resolved to (+)-perforisone A (**1a**) and (-)-perforisone A (**1b**) by HPLC using a chiral column, and their absolute configurations were determined by computational methods via calculation of their electronic circular dichroism (ECD) spectra. Herein, we describe the structural elucidation and biological evaluation of the isolated compounds.

Perforisone A (**1**) was obtained as a colorless solid and had the molecular formula $C_{16}H_{18}O_6$ by the positive HRESIMS (m/z 329.0996 [$M + Na$]⁺, calcd: 329.0996), requiring 8 degrees of unsaturation. Its IR spectrum showed absorption bands for hydroxyl (3435 cm^{-1}), carbonyl (1653 cm^{-1}), and aromatic ($1582, 1559, 1490$ and 1457 cm^{-1}) groups. In the ^1H NMR spectrum, two aromatic H-atoms ($\delta_{\text{H}} 6.32, \text{s}, 6.58, \text{s}$), one spin system ($\delta_{\text{H}} 2.95, \text{dd}, J = 13.1, 7.6\text{ Hz}$, $\delta_{\text{H}} 2.86, \text{dd}, J = 13.1, 6.8\text{ Hz}$, $\delta_{\text{H}} 4.34, \text{t}, J = 7.1\text{ Hz}$), one singlet methyl ($\delta_{\text{H}} 1.80, \text{s}$), together with one methoxyl ($\delta_{\text{H}} 3.89, \text{s}$) were observed (Table 1). The ^{13}C NMR (DEPT) spectrum exhibited 16 carbon resonances, assigned to one aryl ketocarbonyl ($\delta_{\text{C}} 184.2$), seven sp^2 quaternary carbons ($\delta_{\text{C}} 171.6, 165.5, 160.0, 158.2, 148.7, 111.1, 106.1$), one sp^2 methylene ($\delta_{\text{C}} 111.3$), two sp^2 methines ($\delta_{\text{C}} 106.9, 91.0$), two sp^3 methylenes ($\delta_{\text{C}} 61.4, 29.4$), one methoxy ($\delta_{\text{C}} 56.6$), and one methyl ($\delta_{\text{C}} 17.3$). These groups accounted for six out of eight indices of unsaturation.

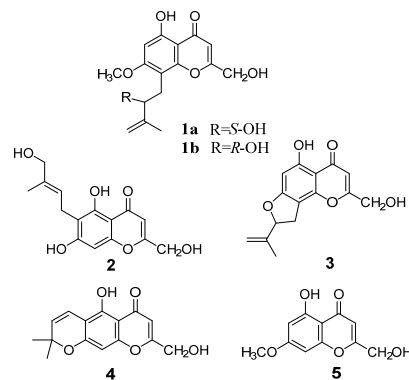


Figure 1: Structures of compounds **1-5**.

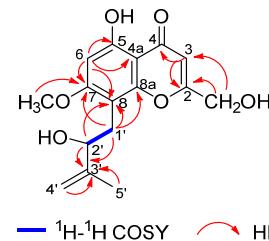


Figure 2: Selected ^1H - ^1H COSY (bold) and key HMBC correlations (Arrow, $\text{H} \rightarrow \text{C}$) of **1**.

From the ^1H - ^1H COSY correlation of $\text{H}_{2-1'}$ and $\text{H}-2'$, combined with the HMBC correlations of $\text{Me}-5'/\text{C}-3'$, $\text{H}-2'/\text{C}-3'$, $\text{H}_2-1'/\text{C}-3'$, and $\text{H}-2'/\text{C}-3'$, the side chain was assigned as a 2-hydroxy-3-methyl-3-butenoate group (Figure 2). The attachment of this unit at C-8 was confirmed by HMBC correlations from $\text{H}_{2-1'}$ to C-7, C-8, and C-8a. The NMR data of **1** were similar to those of perforamone C [6], except for the presence of a hydroxymethyl group instead of a methyl group in the latter. The position of the hydroxymethyl group

Table 1: ^1H and ^{13}C NMR data (600 and 125 MHz, resp.) of compound **1** in MeOD.

No	^{13}C	^1H
2	171.6	
3	106.9	6.32 (1H, s)
4	184.2	
4a	106.1	
5	158.2	
6	91.0	6.58 (1H, s)
7	165.5	
8	111.1	
8a	160.0	
1'	29.4	2.95 (1H, dd, $J = 13.1, 7.6$) 2.86 (1H, dd, $J = 13.1, 6.8$) 4.34 (1H, t, $J = 7.1$)
2'	76.0	
3'	148.7	
4'	111.3	4.61 (1H, br s) 4.60 (1H, br s)
5'	17.3	1.80 (3H, s)
2-CH ₂ OH	61.4	4.48 (2H, s)
7-OCH ₃	56.6	3.89 (3H, s)

at C-2 was supported by HMBC correlations of 2-CH₂OH (δ_{H} 4.48) to C-2 (δ_{C} 171.6) and C-3 (δ_{C} 106.9). Thus, the structure of **1** was established as 5-hydroxy-7-methoxy-2-hydroxymethyl-8-(2-hydroxy-3-methyl-3-but enyl)chromone. It is worth noting that the CD spectrum of **1** is almost a straight line, implying a racemic nature of **1**, which was in accordance with its optical activity ($[\alpha]^{25}_{\text{D}} = 0$). Chiral resolution of **1** was achieved on a CHIRALPAK IC column; the enantiomers **1a** and **1b** were well separated.

Compounds **1a** and **1b** had opposite optical rotations (**1a**: $[\alpha]^{25}_{\text{D}} = +21.5$; **1b**: $[\alpha]^{25}_{\text{D}} = -24.4$) and Cotton effects in their CD spectra. The final assignment of absolute configurations of **1a** and **1b** was achieved by comparison of the calculated electronic circular dichroisms (ECDs) with experimental data [19]. In detail, their ECDs calculated at the b3lyp/6-311++g(d,p)//b3lyp/6-311++g(d,p) level showed good agreement with the experimental spectra (Figure 3).

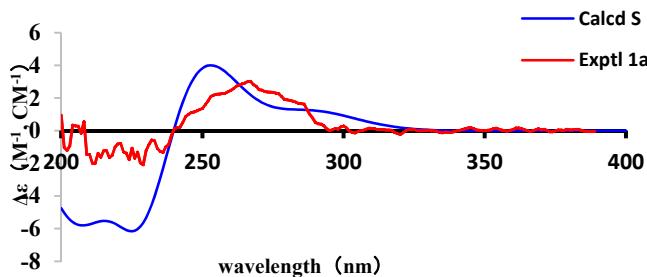


Figure 3: Calculated and experimental ECD spectra of **1a** (blue, at the B3LYP-SCRF (PCM)/6-311+G(2d,p)//B3LYP/6-311+G(2d,p) level in MeOH; red, experimentally observed in MeOH).

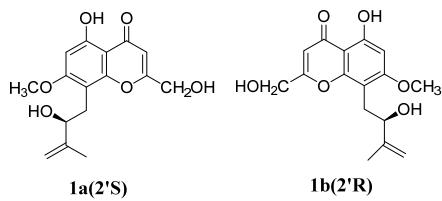


Figure 4: Absolute configuration for **1a** (2'S) and **1b** (2'R).

From the above evidence, the absolute stereochemistry for **1a** (2'S) and **1b** (2'R) were unambiguously determined as shown (Figure 4).

All compounds were evaluated for their *in vitro* growth inhibitory effects against five human cancer cell lines, namely, HL-60 (human

promyelocytic leukemia cell line), SMMC-7721 (human hepatocellular carcinoma cell line), A-549 (human lung cancer cell line), MCF-7 (human breast cancer cell line), and SW480 (colorectal cancer cell line) using MTS method [20]. However, none of them exhibited significant bioactivity ($IC_{50} > 40 \mu\text{M}$).

Experimental

General experimental procedures: Optical rotations were measured with a Jasco P-1020 polarimeter. UV spectra were obtained using a Shimadzu UV-2401A spectrophotometer. A Tenor 27 spectrophotometer was used for IR spectra as KBr pellets. 1D and 2D NMR spectra were recorded on a Bruker AV-500 spectrometer with TMS as internal standard. HRESIMS were obtained on a triple quadrupole mass spectrometer. Semi-preparative HPLC was performed on an Agilent 1100 liquid chromatograph with either a Waters X-Bridge Prep Shield RP18 (10 × 150 mm) column or a CHIRALPAK IC (10 × 250 mm) column. Column chromatography (CC) was performed using silica gel (100-200 mesh and 300-400 mesh, Qingdao Marine Chemical, Inc., Qingdao, P. R. China) and Sephadex LH-20 (40-70 μm , Amersham Pharmacia Biotech AB, Uppsala, Sweden). MCI gel CHP 20P (75-150 μm , Mitsubishi Chemical Industries, Tokyo, Japan)

Plant material: The dried leaves and stems of *H. perforata*, collected in Hainan Province, P. R. China, in October 2014, were identified by Prof. Yanhui Fu (Hainan Normal University). A voucher specimen (HXJ20141208) was deposited at the State Key Laboratory of Phytochemistry and Plant Resource in West China, Kunming Institute of Botany, Chinese Academy of Science (CAS).

Isolation and purification: The air-dried powdered leaves and stems (19.0 kg) were extracted 3 times (4, 3, and 3 h) with 95% ethanol (20 L × 3) at room temperature and concentrated *in vacuo* at 60°C to give a residue (2.5 kg). This was dispersed in water (4.0 L) and extracted with light petroleum and ethyl acetate in turn. The light petroleum and ethyl acetate portions (1 kg) were combined and then passed over a silica gel column, eluting with a gradient of light petroleum-acetone (from 1:0 to 0:1), to yield 5 major fractions (1-5). Subsequently, Fr.3 (200 g) was separated over an MCI column (MeOH-H₂O from 3:7 to 10:0) to obtain 4 sub-fractions (3A-3D). Fr.3B (20 g) was then chromatographed on a silica gel column eluted with chloroform-acetone (100:1 to 5:1), to afford 5 sub-fractions (3B1-3B5). Fr.3B2 (2.2 g) was purified by Sephadex LH-20 (methanol) to obtain **4** (55 mg) and a major fraction (Fr.3B21). Fr.3B21 (200 mg) was separated by semi-preparative HPLC (CH₃CN-H₂O, 4:6) to give **2** (3 mg) and **3** (5 mg). Fr.3B2 (2.2 g) was purified by Sephadex LH-20 (methanol) to obtain Fr.3B22 and Fr.3B24. Fr.3B22 (500 mg) was then chromatographed on a silica gel column eluted with chloroform-acetone (25:1) to obtain **1** (6 mg). Compound **1** was finally subjected to chiral HPLC to afford **1a** ((+)-perforisone A, 1.1 mg) and **1b** ((-)-perforisone A, 2.3 mg) (*n*-hexane/isopropanol, 85:15). Fr.3B24 (200 mg) was separated by semi-preparative HPLC (CH₃CN-H₂O, 5:5) to give **5** (10 mg). The purities of compounds **1-5** were 95%, as determined by TLC and HPLC.

Calculation methodology: Calculations were performed using the Gaussian 03 program package. Geometries were fully optimized with the density functional theory methods of B3LYP at the 6-31+G* level. Harmonic vibrational frequency of the conformation was then calculated at the same level to confirm their stability. Lastly, the “self-consistent reaction field” method (SCRF) was employed to perform the ECD calculation of compounds **1a** and **1b** in methanol at the B3LYP-SCRF/6-311++G (d, p) level.

(±)-Perforison A

Colourless solid.

{ $[\alpha]^{25}_D$: + 21.5 (*c* 0.18, MeOH); **1a**}; { $[\alpha]^{25}_D$: -24.4 (*c* 0.16, MeOH); **1b**}UV (MeOH) λ_{max} (log ε): 312 (3.78), 292 (3.91), 258 (4.16), 232 (4.24) nmIR (KBr) ν_{max} : 3435, 1669, 1653, 1624, 1582, 1559, 1540, 1490, 1457, 1411 cm^{-1} . ^1H and ^{13}C NMR: Table 1.ESI-MS (*m/z*): 329 [$\text{M} + \text{Na}$]⁺, 635 [2 $\text{M} + \text{Na}$]⁺;HR-ESI-MS: *m/z*: 329.0996, [$\text{M} + \text{Na}$]⁺ (calcd for $\text{C}_{16}\text{H}_{18}\text{O}_6\text{Na}$, 329.0996).**References**

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Antiinflammatory and Analgesic Effects in Rodent Models of Ethanol Extract of *Clausena anisata* Roots and their Chemical Constituents

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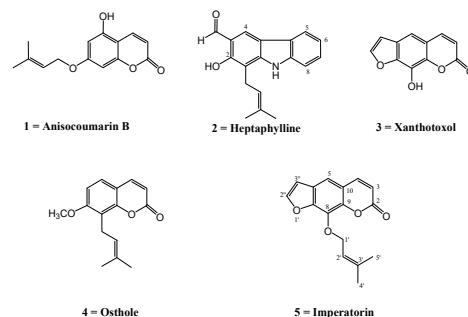
The *in vivo* antiinflammatory and analgesic activities of the crude ethanol extract and chemical constituents of *Clausena anisata* roots were investigated. The crude extract, which was devoid of any visible acute toxicity, displayed significant antiinflammatory effect at the dose of 1000 mg/kg (p.o.) when assessed using the carrageenan-induced oedema model. In the acetic acid-induced writhing and hot plate tests, it produced a very significant ($p < 0.001$), dose-dependent analgesic effect, with maximum analgesic activity of 72.1% at 1000 mg/kg (p.o.). Phytochemical analysis of the crude extract resulted in the isolation of four coumarins (anisocoumarin B, osthole, imperatorin and xanthotoxol) and a carbazole alkaloid, heptaphylline. Among the isolated compounds, osthole and anisocoumarin B produced the highest antiinflammatory activity at 9 mg/kg (p.o.): slightly better than the positive control, indomethacin. Except for xanthotoxol, all the isolated compounds administered at 6 mg/kg (p.o.) produced significant analgesic activity and higher than diclofenac; with heptaphylline being the most potent (48.7%). The analgesic activity of anisocoumarin B (50.4%) was the highest among the isolates tested and the standard, tramadol, in the hot plate test. The nonselective opioid receptor antagonist, naloxone, abolished the analgesic effect of the crude extract and the tested isolates (anisocoumarin B and xanthotoxol) in the hot plate test suggesting an effect *via* the central opioidergic system. These findings provide the scientific basis for the use of *C. anisata* roots in traditional medicine as antiinflammatory and analgesic agents.

Keywords: *Clausena anisata* roots, Antiinflammatory, Analgesic, Coumarins, Carbazole alkaloid, Antinociception, Opioid antagonist.

While self-limiting acute inflammatory response is critical for maintaining homeostasis and defense against a range of pathogens, unregulated inflammation is a feature of many disease conditions from inflammatory pain and allergies to cancer, cardiovascular diseases, and autoimmune disorders. To date, a number of drugs belonging both to the small molecular weight and protein-based modulators are available to combat inflammation and the associated diseases. Due to various limitations (e.g. poor efficacy and unwanted side effects), especially for chronic diseases such as arthritis and inflammatory bowel diseases, the world-wide effort on the search for safe and effective drugs continues in many academic and pharmaceutical industry laboratories [1a,b]. In addition to suppression of inflammatory responses through action like the steroid or non-steroidal mechanisms, drug candidates targeting inflammatory pain through the opiate mechanism of action, like that of tramadol, are highly valuable.

Clausena anisata (Wild.) Hook f. ex. Benth (Rutaceae) is an evergreen shrub or tree that grows up to 10 m high. The plant is widely distributed in higher-rainfall regions of tropical West Africa, as well as Asia and Australia [2a]. The plant is characterised by its distinctive pinnate compound leaves, and by flowers which are highly scented [2a]. In West Africa, the leaves, stems and roots are used to treat countless diseases such as oral candidiasis, fungal infections, epilepsy, convulsion, toothache, gut disturbances, malaria and high blood pressure [2b-f]. Furthermore, the roots serve

as major ingredients in a decoction known as ‘Asena’ in Ghana, which is commonly used to treat arthritis, general bodily pains and other inflammatory conditions at the WHO collaborative Centre for Plant Medicine Research (CPMR) at Mampong-Akwapim. Antiinflammatory, antipyretic, anti-plasmodial and analgesic activities of its leaves extracts and hypoglycemic effect of the roots have been previously reported [3a,3b]. Even though various classes of secondary metabolites (e.g. carbazole alkaloids, coumarins and limonoid) have been isolated from the plant [3c-3g], the antiinflammatory and analgesic activities of the roots so far have not been investigated. This work was therefore designed to investigate these pharmacological properties and isolate and characterize the active principles of *C. anisata* roots (1-5).



The acute toxicity study of the crude ethanoic extract of *C. anisata* roots over a period of 12 days was conducted at exceedingly high doses of either 2500 or 5000 mg/kg p.o, which are 2.5 and 5-times higher than the maximum therapeutic concentrations used in the study respectively. All extract-treated animals neither showed any sign of toxicity nor died during the observation period.

The carrageenan-induced oedema in rat's paw that serves as the gold standard *in vivo* model of antiinflammatory evaluations was employed. The dose regime and protocols were based on our preliminary experiments and previous reports where we have employed the method for the identification of several novel anti-inflammatory agents from medicinal plants [4a-c]. As shown in Figure 1, the maximum carrageenan-induced oedema was attained within 3 h of injection and was sustained during the 5 h observation period. As expected, and in line with our previous report [4c], the positive control indomethacin displayed a time- and dose-dependent antiinflammatory effect within the dose range of 9 and 30 mg/kg (p.o.) (Figure 1).

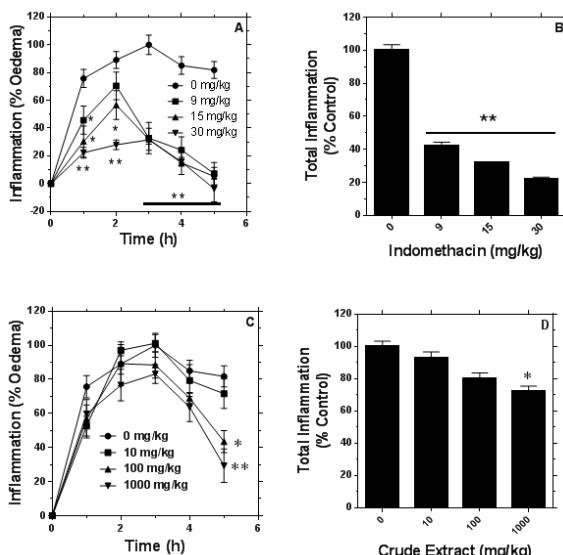


Figure 1: The effect of indomethacin and crude ethanolic extract of *C. anisata* roots on the time-course and overall carrageenan-induced oedema formation in rats paw. * and ** represent significant differences at the level of $p < 0.01$ and $p < 0.001$, respectively. All data represent means and SEM ($n = 5$) values.

The time-course curve and total mean oedema of the crude extract on the carrageenan-induced inflammation are shown in Figure 1c and 1d, respectively. The dose dependent effect of the crude extract was evident though a significant ($p < 0.01$) inhibition of inflammation was observed at the highest dose (1000 mg/kg p.o) tested. At this dose, the overall antiinflammatory activity of 27.5% ($p < 0.05$) was recorded. As compared with the maximum anti-inflammatory activity of indomethacin (78.2% at 30 mg/kg p.o. ($p < 0.001$), the activity of the crude extract appears to be somehow weaker. As the crude extract is a mixture of several hundreds of potentially therapeutically useful and non-useful pharmacological agents, the isolation and identification of the active principles is well justified. On the basis of these observations, our study seems to provide the scientific evidence for the traditional medicinal uses of *C. anisata* roots for inflammatory disorders. It is also worth noting that the carrageenan-induced oedema in rat paws [5a] is an acute inflammatory model that is attributed to three distinct phases marked by sequential release of several mediators [5b]. The 0-1.5 h period after carrageenan injection is known to involve the release of histamine and serotonin and constitutes Phase 1. This is followed by

Phase 2 mediated by the liberation of bradykinin from 1.5 to 2.5 h [5c]; the final phase is mediated by the release of prostaglandins and slow reaction agents from 2.5 to 6 hours [5d]. The fact that the crude extract of *C. anisata* roots significantly inhibited the oedema at 5 h, which corresponds to the 3rd phase, suggests a mechanism of action through inhibition of prostaglandin synthesis and/or actions.

Even though the analgesic activities of *C. anisata* leaves were demonstrated previously [3a], the roots have not yet been studied for such activities. The acetic acid-induced writhing response in animals is one of the commonest methods of analgesic activity evaluation for experimental agents. As expected, the standard positive control, diclofenac, caused a dose-dependent analgesic effect, as evidenced by the reduction in the righting response (Figure 2). Similarly, the crude extract displayed dose-dependent analgesic effects, with all the tested concentrations (10, 100 and 1000 mg/kg, p.o.) inducing significant activities. Hence, the analgesic activity of the crude extract appears to be more pronounced than the antiinflammatory activity as only the 1000 mg/kg dose produced a suppressive effect against the carrageenan-induced oedema (Figure 1).

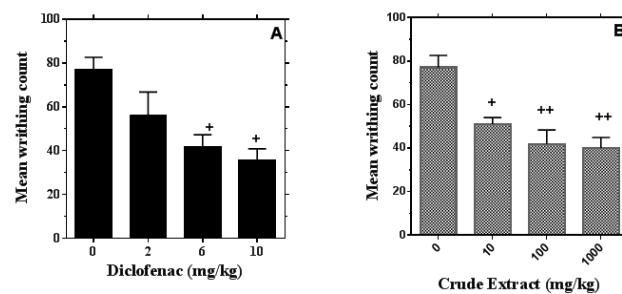


Figure 2: Analgesic effect of the crude ethanolic extract of *C. anisata* and diclofenac on acetic acid-induced pain in mice. + and ++ represent significant differences at the level of $p < 0.05$ and $p < 0.01$, respectively when compared with the vehicle treated control group.

The acetic acid-induced writhing test is a visceral pain model [6a] used in evaluating both central and peripheral analgesic activities of substances [6b]. Intraperitoneal injection of acetic acid into the abdominal cavity of rats or mice results in elevated levels of prostaglandins in the peritoneal exudates after about 30 min [6c]. The stretching and abdominal contractions obtained in the acetic acid-induced writhing test therefore correlates with sensitization of nociceptors to the production of prostaglandins [6b]. Our data on the analgesic effect of the crude extract of *C. anisata* roots are therefore consistent with those obtained in the carrageenan-induced oedema assay, where possible inhibition of prostaglandins production and/or action could serve as the main mechanism of action.

The analgesic activity of the crude extract was further assayed using the hot plate method where the standard tramadol was employed as a positive control. In untreated animals, there was a tendency for the percent mean PTI to reduce over time as the animals appear to be more sensitive to the hot plate stimuli. This sensation was however suppressed in a time-dependent manner when animals were treated by the positive control, tramadol (Figure 3a). The overall analgesic effect, as evidenced from the percent OPTI of over 20% (Figure 3b), was recorded for the two doses of tramadol. As shown in Figure 3c, the crude extract also showed a dose-dependent analgesic effect that appears to be gradually reversed over a period of 5 h. Furthermore, the percent OPTI response showed a remarkable dose-dependent analgesic effect that ranged from about 25 to just over 75% (far superior to that of the positive control) for the

concentration range tested (Figure 3d). It is worth noting that the hot plate test, as a thermally-induced nociception model, is routinely employed to evaluate the central analgesic effect of pharmacological agents [6b]. The dose-dependent antinociception of the crude extract in the hot plate test, similar to that of a known analgesic with an opioidergic mechanism of action, tramadol ($p < 0.001$), may suggest a central mechanism of action.

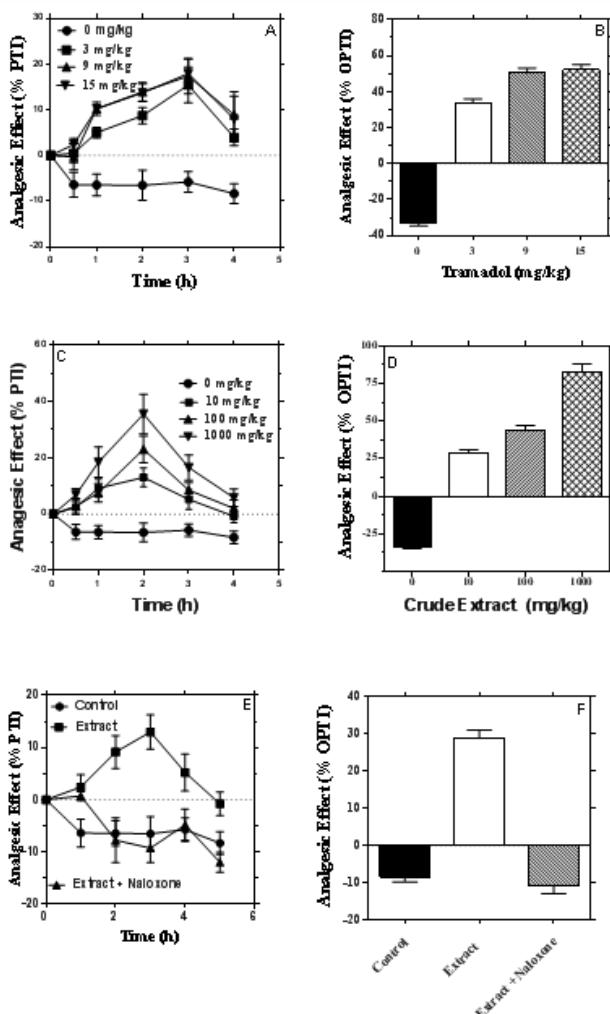


Figure 3: Analgesic effect of crude extract of Tramadol and *C. anisata* roots on the time course curves (left panel) and % overall analgesic response, % OPTI, (right panel) on thermally-induced pain in mice. Effect of naloxone (2 mg/kg, i.p.) on the crude extract (10 mg/kg, p.o.), on thermal mediated antinociception on time course curve (E) and % OPTI (F) are also shown. All data are expressed as means \pm SEM and the % OPTI data for the drug treated groups, except for the naloxone group were significantly different (at least $p < 0.05$) when compared with the vehicle-treated control group.

In order to establish the possible central mechanism of the crude extract in the observed analgesic/antinociceptive effect, the hot plate assay was further reassessed with naloxone co-administration. As shown in Figure 3 (e,f), the activity of the crude extract was completely abolished by the non-selective opioid antagonist, naloxone. This indicates that the analgesic effect of the crude extract was likely to be mediated *via* the central opioidergic system.

Having the antiinflammatory and analgesic activity of the crude extract of *C. anisata* roots established, it was worthwhile to investigate the possible active principles of the roots. Preliminary phytochemical screening of the crude extract using the standard procedure [7a] revealed the presence of coumarins, alkaloids, terpenoids and free reducing sugars. When the crude extract was

suspended in aqueous medium and re-extracted with light petroleum, the presence of coumarins, alkaloids and terpenoids was still evident. Hence, the light petroleum fraction was taken for further purification through repeated column chromatography over silica gel and recrystallization to yield five compounds (**1-5**). The detection and identification of all of the isolated compounds in the various parts of *C. anisata* have been described previously [3c-g, 7b-c]. In the present study, comprehensive spectroscopic analysis including 2D NMR studies (COSY, HMQC, HMBC and NOESY) and accurate mass analyses were employed for unambiguous chemical shift assignments and structural analysis.

As with the crude extract, the isolated compounds were tested for their *in vivo* antiinflammatory activities using the same carrageenan-induced oedema model. The dose-dependent effect of these compounds in an overall total inflammatory effect analysis during the entire observation period is shown in Figure 4. All compounds showed significant antiinflammatory activity at 9 mg/kg (p.o.) suggesting their potent antiinflammatory effects *in vivo*. Furthermore, all compounds, except for compound **2** (heptaphylline) and **5** (imperatorin) showed comparable activity with the positive control indomethacin when tested at the dose of 9 mg/kg ($p > 0.05$).

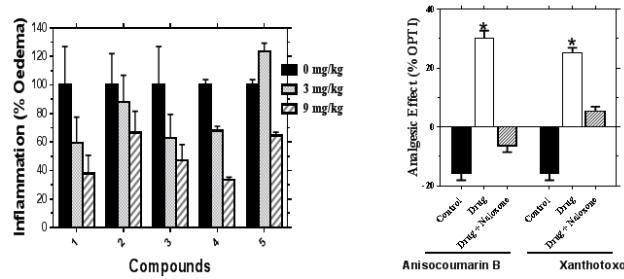


Figure 4: Antiinflammatory and analgesic activity of the isolated compounds. The overall carrageenan-induced oedema formation in rats paw (left) and thermal antinociception effect (right panel) are shown. *, indicate significant difference ($p < 0.001$) vs either the vehicle treated control group or the drug plus naloxone group. Significant antiinflammatory activity (at least $p < 0.05$) was observed at the dose of 9 mg/kg. All data are means \pm SEM ($n = 5$).

Osthole (**4**) is one of the best characterised anti-inflammatory natural products with known effect as an inhibitor of the carrageenan-induced hind paw edema in rats [8a]; carrageenan-induced lung inflammation in rats [8b]; hepatic injury in a rodent model of trauma-hemorrhage [8c]; inhibition of inflammatory reaction following permanent middle cerebral artery occlusion in rats [8d]; and iNOS protein expression at 10 μ M [8e], among others. Imperatorin (**5**) has been shown to suppress the degranulation and eicosanoid generation in activated bone marrow-derived mast cells [8f]; inhibit the release of pro-inflammatory cytokines production, including interleukin (IL)-6 and tumor necrosis factor (TNF)- α in LPS-stimulated RAW 264.7 cells [8g-8h]; block the protein expression of iNOS and cyclooxygenase-2 (COX-2) in LPS-stimulated RAW264.7 macrophages [8i]; inhibit acute lung injury induced by lipopolysaccharide in mice [8j] and anti-inflammatory activity on TPA (12-O-tetradecanoylphorbol-13-acetate) mice ear model [8k]. While xanthotoxol (**3**) is known to exert neuroprotective effects *via* suppression of the inflammatory response in a rat model of focal cerebral ischemia [8l] its antiinflammatory effect is largely unknown. On the other hand, heptaphylline (**2**) has been known for some time to have an *in vitro* antiinflammatory effect such as suppression of inflammation mediators (iNOS, TNF-alpha, and COX-2) expression in the mouse macrophage RAW 264.7 cell line [8f], while anisocoumarin B (**1**) has not previously been shown to display anti-inflammatory

activity. The identification of compounds **1-5** from the roots of *C. anisata* is, therefore, in good agreement with the observed *in vivo* antiinflammatory effect of the crude extract and the proposed mechanism of action, including the prostaglandins system.

Since the yield of the isolated compounds did not permit the undertaking of a comprehensive dose range analysis, they, along with the positive control, were screened for analgesic activity at one fixed dose of 6 mg/kg (p.o.). As shown in Table 1, only compound **3** (xanthotoxol) failed to show analgesic activity at the tested dose. The observed analgesic activity of the active compounds, which was comparable in many cases with the standard diclofenac, was in good agreement with the activity profile of our data for the crude extract in the same assay. It is, therefore, reasonable to conclude that the isolated active compounds as active principles could contribute to the analgesic activity of the acclaimed *C. anisata* roots in the traditional medicine.

Table 1: Analgesic activity of the isolated compounds and diclofenac on acetic acid - induced writhing pain in mice.*

Compounds	% Righting Response Vs Control	% Analgesic Activity
1 (Anisocoumarin B)	61.4 ± 16.0	38.6
2 (Heptaphylline)	49.1 ± 12.6	50.9
3 (Xanthotoxol)	112.1 ± 3.3	No effect
4 (Osthole)	53.6 ± 6.7	46.4
5 (Imperatorin)	61.9 ± 12.7	38.1
Diclofenac	53.9 ± 7.5	46.1

*Tested at 6 mg/kg (p.o.)

The analgesic activity of osthole (**4**) was reported before [81], including in sciatica induced by the lumber disc herniation model [9a-b], while the rest of the isolated compounds appear to be reported here for the first time. Due to the small yields of the compounds, it was not possible to repeat all of the experiments in the hot plate assay and hence data were provided in Figure 4 only for anisocoumarin B (**1**) and xanthotoxol (**3**). While both compounds appear to show significant activity, the effect of anisocoumarin B (**1**) was higher than that of the positive control, tramadol. As with the analgesic effect of the crude extract, the antinociceptive effects of these compounds were suppressed by the non-selective opioid antagonist, naloxone, suggesting a similar central effect (Figure 4).

In conclusion, the present study established that the roots of *C. anisata* possess significant antiinflammatory and analgesic effects similar to NSAIDs and/or opioids and could be safely used to manage some inflammatory and painful disorders. This study also validates the use of *C. anisata* roots in traditional medicine as an anti-inflammatory and analgesic agent. Phytochemical analysis of the crude extract resulted in the isolation of prenyl derivative compounds: four coumarins (anisocoumarin B, osthole, imperatorin and xanthotoxol) and a carbazole alkaloid, heptaphylline. In a further validation of the traditional uses of the plant, the isolated compounds also possess potent antiinflammatory and analgesic activities.

Experimental

Materials: Carrageenan, naloxone hydrochloride dehydrate, and acetic acid were products of Sigma Chemical Co. (St. Louis USA). Indomethacin, diclofenac sodium, tramadol hydrochloride and Tween 80 were purchased from Cayman Chemical Company, Bliss GVS (India), Bristol Laboratories Ltd and VWR International, PROLABO (CE) respectively. Sodium chloride was from Timstar Laboratory Suppliers Ltd., Cheshire, England. All organic solvents and other reagents used for extraction, isolation and purification of the compounds were either analytical or HPLC grade.

Plant material: The roots of *C. anisata* were harvested from the arboretum of CPMR at Ayikumah in the Eastern Region of Ghana in March, 2012. After authentication by Mr H. R. Blaggoge, a voucher specimen (CPM0312) was deposited at the herbarium of CPMR.

Extraction and isolation: The crude extract (140 g) was suspended in 80% ethanol-water (2.4 L) and extracted 4-times with light petroleum (2.4 L). The combined fractions were dried in a rotary evaporator at 40°C to obtain 16.8 g of a bark brown syrup. A portion of this fraction (13.0 g) was subjected to column chromatography (CC) over normal phase silica gel (400 g) and elution made with light petroleum, light petroleum containing increasing amount of ethyl acetate, and finally a gradient of EtOAc containing ethanol. Aliquots of fractions from the column were collected and grouped by similarity in their TLC profiles to give 7 main fractions (P1-P7). Fraction P1 eluted with light petroleum /ethyl acetate (95:5) gave colorless crystals, which were filtered, washed (light petroleum /chloroform 6:1) and recrystallized from light petroleum /ethyl acetate (70:30) to give compound **1** (40.5 mg). Fraction P2 (light petroleum /ethyl acetate 80:20) gave a yellow ppt, which appeared to be a mixture when assessed by TLC (solvent, light petroleum/chloroform (2:3); anisaldehyde spray reagent). Further CC over 40 g of silica and elution with light petroleum /chloroform gave 3 sub-fractions. The sub-fraction eluted with light petroleum /chloroform (90:10) gave compound **2** (10 mg) as bright yellow crystals. Fraction P4 (solvent, eluent, light petroleum /ethyl acetate 60:40) was subjected to further fractionation with a stepwise gradient elution with light petroleum /chloroform/ ethyl acetate. A total of 87 fractions of 30 mL aliquots was obtained and grouped into 4 sub-fractions. Compound **3** (35.0 mg) was obtained from the fraction eluted with light petroleum /chloroform/ethyl acetate (40:40:20) as brown solid particles. Fraction P5, eluted with light petroleum /ethyl acetate (60:40) gave a colorless ppt, which was subjected to repeated recrystallization from light petroleum /chloroform (4:1) to obtain compound **5** (20.5 mg). Fraction P6, eluted with light petroleum /ethyl acetate (30:70) yielded white crystals, which were recrystallized from light petroleum /ethyl acetate (5:1) to obtain compound **6** (30.0 mg).

Animals: All animals used in the study were bred at the Animal House Unit of CPMR, Ghana. The animals were fed on palette feed purchased from Agricare Limited located in Kumasi, Ghana; and were allowed free access to sterile water and feed *ad libitum*. Animals were housed in aluminum cages under standard temperature and pressure. Animals were cared for and handled according to the guidelines and procedures of the Foundation for Biomedical Research on the use of animals in research (F.B.R., 1987).

Acute toxicity studies: The acute toxicity of the crude extract was assessed in groups of 6 male Sprague-Dawley rats, Swiss albino mice and C57/BL6 mice. For each animal group, 2 doses of the crude extract (2500 and 5000 mg/kg p.o.) and a vehicle control (2% Tween 80 aqueous solutions) were administered at 10 mL/kg body weight per animal. They were observed for signs of toxicity such as pilo-erection, motor impairment, sedation, salivation, hyperexcitability and death within 24 h and for an extra 16 days.

Antiinflammatory activity assay: Our carrageenan-induced paw oedema model of antiinflammatory study [5a] has been described previously [4a-c]. Briefly, Sprague-Dawley rats of either sex were divided into groups and received orally either the vehicle (2% Tween 80 in water) or various doses of test agents (extracts, isolated

compounds or the reference drug, indomethacin). After 1 h of drug treatment, inflammation was induced by injection of 0.1 mL of 1%, w/v, carrageenan in 0.9% normal saline into the sub plantar area of the right hind paw of rats. Paw volumes were measured by volume displacement using a Plethysmometer (Ugo Basile 7140) before and at hourly intervals from 1-5 h after carrageenan injection. The mean hourly oedema for each group and the total mean oedema for the 5 h observation period were calculated as described previously [4b].

Analgesic activity assay: Acetic acid-induced writhing assay: The analgesic effect of the test agents was assessed using the acetic acid-induced writhing model as described previously [10a]. Swiss albino mice were divided into 7 groups (n = 5) and treated (p.o.) with various doses of the crude extract, isolated compounds and the standard, diclofenac sodium. Animals receiving 2% Tween 80 in water served as the untreated vehicle control group. Following drug or vehicle treatment (45 min), each mouse was injected with aqueous acetic acid (1%, v/v) at 1 mL/100 g (i.p) and isolated into separate plastic cages. The number of writhing movements and stomach contractions was counted for 20 min. The inhibition of writhing movements in the treated group(s) compared with the control group was taken as the percentage analgesic effect (% AE), which was calculated using the formula:

$$\% \text{ Analgesic effect} = (\text{MRc} - \text{MRt})/\text{MRC} \times 100$$

Where: MRc = mean writhing count of the control; MRt = mean writhing count of extract or other drug treated group.

Analgesic activity assay: Hot plate assay: The mouse hot plate analgesic activity evaluations were performed as described previously [10b]. C57BL6 mice of either sex were divided into 7 groups, each containing 6 animals. The mice were separately placed on an electric hot plate (Ugo Basile hot/cold plate 35 100) maintained at $55 \pm 0.5^\circ\text{C}$ and the time taken to lick, lift, shake or stamp any of the hind limbs or jump constitute latency time and was recorded. Baseline latencies (T_0) were obtained as the mean of 2 determinations prior to any treatment. Only mice with baseline latency of 3.5–9 s were used. Test samples of either the crude extract, purified compounds or standard reference, tramadol (3, 9

and 15 mg/kg, Group 4-6) or vehicle control (2% Tween 80 in water control) were then administered orally. Latency of each mouse was measured at 1 h (T_t) intervals for 5 h after each treatment. The analgesic activity was expressed as percentage of pain threshold inhibition (%PTI) [6b] calculated as:

$$\% \text{PTI} = (T_t - T_0)/T_0 \times 100$$

Where T_t and T_0 refer to treatment time and time zero respectively.

Overall analgesic response (% OPTI) for each group was obtained as the sum of mean % PTI of all animals per group over the experimental period.

Mechanism of antinociceptive action: The effect of the nonselective opioid antagonist, naloxone, on the extract or purified compounds- induced analgesia was evaluated in the hot plate assay using 3 groups (n=6) of C57BL/6 mice. Drug administration was made as follows: CRE (at 10 mg/kg p.o.) was administered and 15 min later naloxone 2 mg/kg (i.p) (in distilled water) at 10 mL/100 g was given (Group 1). The procedure was repeated for CRE (10 mg/kg p.o.) alone (Group 2) and 2% Tween 80 aqueous solution as vehicle control (Group 3). The analgesic activity was then assessed as described above in the hot plate test. The same procedure was repeated for compounds 1 and 3 at 9 mg/kg p.o. each.

Statistical analysis: All statistical analyses were performed using Graph Pad Prism Version 6 with the level of significance set at 95 % confidence interval of difference. The overall mean oedema and overall analgesic response (%OPTI) were calculated as sum of the oedema volume or latency in each group.

Spectroscopic analysis: ^1H NMR, ^{13}C NMR and 2D-NMR (COSY, NOESY, HMQC and HMBC) spectra were obtained on a JEOL 500 MHz instrument, as described previously [10c].

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Preventive Effects of Resveratrol-enriched Extract of Peanut Sprout on Bacteria- and Estradiol-induced Prostatitis in Mice

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The present study investigated the effect of peanut sprout extract (PSE) as a natural resveratrol supplement on chronic bacterial prostatitis (CBP) and estradiol-induced benign prostatic hyperplasia (BPH). PSE contained a high level of resveratrol ($148.51 \pm 3.05 \mu\text{g/g}$), and was tested on the mouse models of CBP (induced by *Escherichia coli* 292 infection) and BPH (induced by treatment with β -estradiol and dihydrotestosterone). PSE toxicity was assessed on the basis of changes in body weight, alanine aminotransferase activity (an indicator of hepatotoxicity), and expression of the kidney injury marker KIM-1. The effects of PSE on the histopathology of prostate tissue, the proportion of neutrophils, and immune cell profiles in the blood and spleen were examined. PSE administration did not result in any toxicity but reduced the bacterial burden and histopathological changes in the prostate. In addition, lymphocytes (CD4⁺, CD8⁺, and CD19⁺) in the spleen were significantly increased after PSE administration in CBP mice, suggesting immune enhancement. PSE treatment of bone marrow-derived macrophages increased the expression of CD40, which is related to the pro-inflammatory function and host defense against pathogens. It is concluded that PSE would be a good supplement for the mitigation of prostate hyperplasia and prostatitis.

Keywords: Peanut sprout extract, Resveratrol, Chronic bacterial prostatitis, Benign prostate hyperplasia, Anti-inflammation, Immune enhancement.

Prostatitis and prostate hyperplasia are diseases that commonly occur during men's lifespan all over the world, and 35–50% of men have prostatitis at some time in their lives [1, 2]. Prostate diseases include chronic bacterial prostatitis (CBP), benign prostatic hyperplasia (BPH), and prostate cancer [2-5]. Prostatitis is frequently present in aging men, and many patients are suffering from chronic prostatitis syndrome [1, 5, 6]. Chronic prostatitis of bacterial and nonbacterial origin is a very common disease in men; nevertheless, there are no established management strategies in everyday life. After the onset of symptoms, anti-inflammatory drugs, α -adrenergic blocking agents, and 5 α -reductase inhibitors are the best remedies [2, 6]. However, in recent years, phytomedicines are becoming popular all over the world for management of prostate diseases [4, 5, 7-9]. The major infection routes in chronic bacterial prostatitis are through the urinary tract, rectum, and blood [10-12]. CBP treatment with antibiotics is difficult because of biofilm formation by bacteria, especially *E. coli* spp. [10]. If the symptoms persist, chronic prostatitis can progress to bacterial cystitis, pelvic pain, and acute pyelonephritis [1, 4]. BPH commonly occurs in aged men, accordingly general preventive and therapeutic measures have to be planned for a long time [5]. Complementary alternative medicine (CAM) is becoming popular all over the world for treatment of various diseases. Among prostate diseases, the usefulness of CAM has been mainly reported for prostate cancer, CBP, and BPH [4, 5, 13]. The small polyphenol compound resveratrol is a natural ingredient of herbal remedies and has received widespread attention as a potential therapy or preventive agent for numerous diseases [14]. The scientific interest to resveratrol is related to its use for prevention of carcinogenesis and cardiovascular disease, and to its anti-inflammatory and anti-oxidant properties [14]. Resveratrol also

exerts anti-inflammatory effects and enhances multiple immune functions of human T-cells in vitro, suggesting its ability to regulate the immune response; it also inhibits proliferation of human prostate cancer cell lines [15, 16]. However, the in vivo and in vitro effects of resveratrol on the prostatitis are not well understood. Resveratrol (3,5,4'-trans-trihydroxystilbene) is mainly found in grapes, a variety of other berries, and peanuts [14]. The most important dietary source of resveratrol is red wine; however, peanut (*Arachis hypogaea* L.), which has a high content of resveratrol, has been recently highlighted as a next-generation alternative dietary source of resveratrol [14, 17]. In particular, the content of resveratrol in peanut sprouts (germinated peanut seeds) is higher than in wines or peanut plants [14, 17]. To take advantage of the high resveratrol content in this dietary source, peanut sprouts are processed by drying and extraction of the sprout powder; these extracts (PSE) have been tested for the biological effects of resveratrol in various disease models [17-24]. In this respect, the present study examined the concentration of resveratrol in PSE, assessed the effect of PSE on prostate inflammation in the mouse models of CBP and BPH, evaluated whether administration of PSE as a dietary source of resveratrol causes toxicity to the liver or kidneys, and finally examined the possible role of the immune-stimulating effect of PSE in its ability to alleviate prostatitis.

The composition of PSE used in the present study is shown in Table 1. The major components were proteins, carbohydrates, lipids, and water (25.6, 20.4, 0.3, and 49.1 g per 100 g of PSE, respectively). Among other nutrients shown in Table 1, trans-resveratrol was present at 14.851 mg per 100g of PSE (Table 1). Trans fat and cholesterol were not detected (data not shown).

Table 1: Composition of the peanut sprout extract (PSE).

Ingredient	Content (per 100 g of PSE)	Nutrient	Content (per 100 g of PSE)
Lipids (g)	0.3	Dietary fiber (g)	5.9
Proteins (g)	25.6	Vt. B1/B2/B3/B6 (mg)	3.1/1.8/30.9/0.5
Carbohydrates (g)	20.4	Ca/Na/P/Zn (mg)	223.5/52.3/833.2/1.8
Ash (g)	4.6	Fe/Mg/K/Mn (mg)	1.9/366.1/1169.7/2.0
Water (g)	49.1	Trans-resveratrol (mg)	14.851 ± 0.305

Before the experiments in prostatitis mouse models, the toxicity of PSE was examined in control BALB/c mice (Figure 1). ALT activity and the expression level of the Kim-1 gene were used as indicators of toxicity to the liver and kidney, respectively. No toxicity was found even after administration of a high dose of PSE (200 µL of PSE prepared in 100 mg/mL) (Figure 1). Therefore, this dose was used in further experiments. The concentration of resveratrol contained in this dose is about 2.97 µg (per 200 µL of PSE prepared in 100 mg/mL), and the injection dose is 0.119 mg/kg.

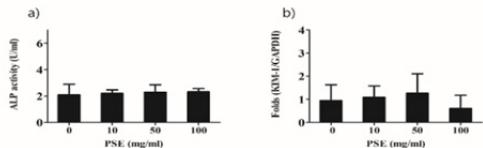


Figure 1: Evaluation of toxicity markers in the liver and kidney after PSE administration ($n=3$). PSE toxicity was evaluated by serum ALP activity for the liver (a) and Kim-1 gene expression for the kidney (b). ALP activity was evaluated using colorimetric assay kit. Kim-1 gene expression was analyzed by RT-PCR and normalized to that of GAPDH.

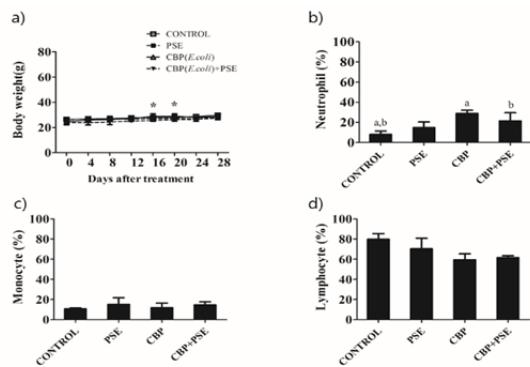


Figure 2: Changes in body weight and in blood cell profiles after PSE administration in CBP mice ($n=5$). Body weight was measured every 4 days (a). Changes (%) in neutrophils (b), monocytes (c), and lymphocytes (d) were examined by microscopic observation. *A statistical significant difference between CBP and CBP+PSE ($p < 0.05$). ^a, ^bA statistical significant results ($p < 0.05$) based on Scheffe post hoc one-way ANOVA analysis (each letter should be compared separately with the letters of other bars).

The mice were divided into four groups ($n=5$): control, CBP (*E. coli*), PSE, and CBP+PSE. The amount of resveratrol in 200 µL of PSE (100 mg/mL) was calculated as 2.97 µg. To investigate the effect of PSE, body weight and blood neutrophils were examined in the four groups. A significant difference in body weight between *E. coli* inoculation (CBP) with or without PSE administration was shown at day 16 and 20 after treatment (Figure 2a, $p < 0.05$). Unless, there was no significant change in body weight of all experimental groups (Figure 2a). The population of inflammatory cells predominantly increased upon the induction of prostatitis-was neutrophils. *E. coli* inoculation (CBP induction) significantly increased the proportion of blood neutrophils (%) compared to that in control mice (28.8 ± 3.4 vs. 8.0 ± 3.4 , respectively, $p < 0.05$) (Figure 2b). PSE administration in CBP mice resulted in a reduction of neutrophils compared to that in CBP mice although it is not statistically different (25.4 ± 11.2 (%) vs. 28.8 ± 3.4 (%)). In contrast, the proportion of blood monocytes was slightly increased in the CBP+PSE group compared to that in the CBP group although the difference is not statistically (14.0 ± 2.9 (%) vs. 11.8 ± 4.6 (%)) (Figure 2c). Increase in neutrophils and monocytes conversely caused the decrease in lymphocytes proportion (Figure 2d).

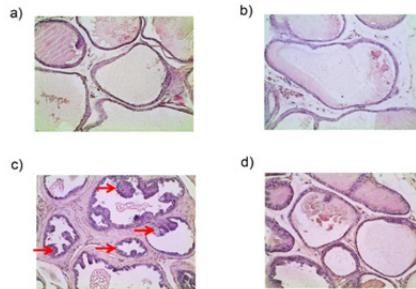


Figure 3: Effect of PSE on histopathology of the prostate in CBP mice. The pathologic changes were examined by H&E staining. Experimental groups were as follows: Control (a); PSE (b); CBP (c); CBP + PSE (d). Arrows in (c) show fibrotic features with enlarged and edematous stroma and tufting of the mucin-secreting epithelium in the granules. PSE administration resulted in a recovery of tissue pathology (d). Tissue sections were examined under $\times 200$ magnification.

Histologically, normal prostate is composed of glandular and stromal tissue (Figure 3a). The prostate of rodents contains 3 separate lobes: dorsal, ventral, and lateral (frequently combined with dorsal as the dorsolateral lobe). Glands (tubules) are lined by two cell layers: the outer layer and the inner layer of columnar mucin-secreting epithelium. The stroma is the fibromuscular layer. The fibromuscular stroma between glandular tissues had an intact muscle layer, and the columnar mucin-secreting epithelium of glands was intact in both control and PSE groups (Figure 3a, b). In contrast, edema was detected in the prostate stroma in CBP mice (2.4×10^5 cfu per gram of prostate tissue, data not shown); in these mice, the prostate was enlarged and had fibrotic stroma suggesting prostatic hypertrophy (Figure 3c). In addition, mucosal epithelial hyperplasia and tufting of the mucin-secreting epithelium in the inner layer of granules were found in CBP mice (Figure 3c). However, these abnormalities disappeared after PSE administration (Figure 3d). In this experiment, the presence of *E. coli* in the prostate was reduced to 0.6×10^5 cfu per gram of prostate tissue (data not shown). In general, PSE administration in CBP mice reduced *E. coli* proliferation to 25% of that without PSE.

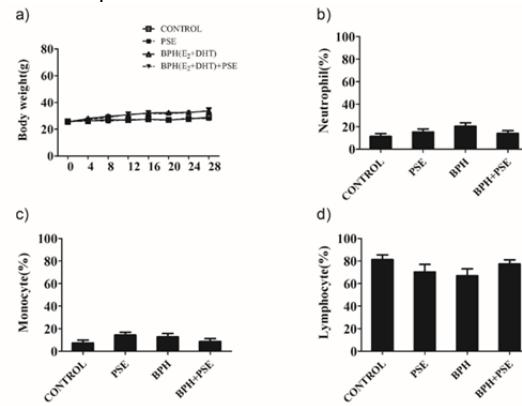


Figure 4: Changes in body weight and blood cell profiles after PSE administration in BPH ($E_2 + DHT$) mice ($n=5$). Body weight was measured every 4 days (a). Changes (%) in neutrophils (b), monocytes (c), and lymphocytes (d) were examined by microscopic observation.

To investigate the effect of PSE on non-bacterial prostate hyperplasia, we used $E_2 + DHT$ to induce estradiol-induced prostatitis with BPH. The mice were divided into four groups ($n=5$): control, BPH, PSE, and BPH+PSE. There was no significant increase in body weights although BPH mice was increased from day 12 to day 28 (from 30.8 ± 0.7 g to 33.6 ± 1.4 g) in comparison with control mice (from 26.8 ± 0.6 g to 29.0 ± 1.1 g) (Figure 4a). During this period, PSE did not affect the body weight both in the control and $E_2 + DHT$ groups. To examine the induction of prostatitis in BPH mice, blood neutrophils were differentially counted as for CBP mice (Figure 4b). The proportion of neutrophils

(%) increased from 11.3 ± 2.1 in the control group to 18.3 ± 5.9 in the BPH group, suggesting the induction of prostatitis. In contrast, in the BPH+PSE group the proportion of neutrophils was reduced to 14.0 ± 2.6 , i.e. by 4.3 percentage points in comparison with the BPH group although it is not statistically different (Figure 4b). The proportion of blood monocytes (%) was also reduced in the BPH+PSE group in comparison with the BPH-group (8.7 ± 2.5 and 12.0 ± 2.8 , respectively) although it is not statistically different (Figure 4c).

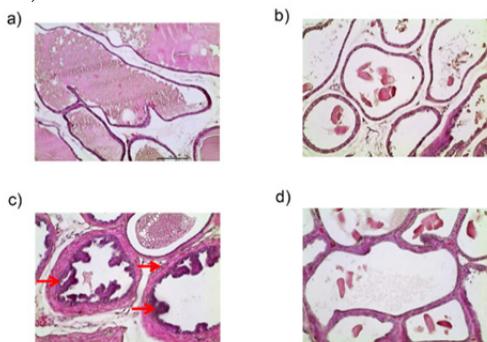


Figure 5: Effect of PSE on histopathology of the prostate in BPH mice. The pathologic changes were examined by H&E staining. Experimental groups were as follows: Control (a); PSE (b); BPH (c); BPH + PSE (d). Arrows in (c) show fibrotic features with enlarged and edematous stroma and tufting of the mucin-secreting epithelium in the granules. PSE administration resulted in a recovery of tissue pathology (d). Tissue sections were examined under $\times 200$ magnification.

In the dorsal lobes of prostate glands, intact muscle layers in the fibromuscular stroma were found both in the control and PSE groups, and glands also had an intact epithelium in the columnar mucin-secreting epithelium of granules (Figure 5a, b). In contrast, epithelial cell hyperplasia and swollen epithelial layers were observed in the prostate glands of BPH mice (Figure 5c). In addition, the stroma in BPH mice was thicker than in control mice, suggesting prostatic hypertrophy (Figure 5c). PSE administration remarkably reduced the pathological signs (edema with enlarged tissue and fibrotic features in the stroma as well as mucosal epithelial hyperplasia in the granules) in the prostates of BPH mice (Figure 5d). Macrophages (BMMs) were isolated from the bone marrow. To investigate the possibility of immune activation after PSE treatment, we incubated BMMs with PSE, *E. coli* LPS, or both, and analyzed the expression of the macrophage activation marker CD11b⁺ and the antigen presenting cell activation marker CD40⁺ (Figure 6).

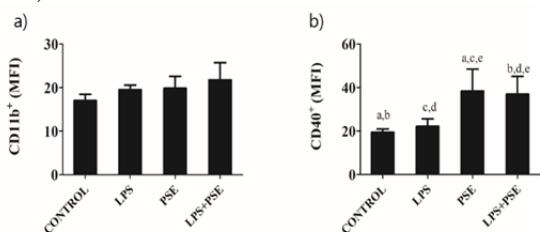


Figure 6: Effect of PSE on the CD11b and CD40 expression. BMMs were incubated with PSE (5 mg/mL), and examined for CD11b (a) or CD40 (b) by FACS analysis. The expression of CD40 was increased in PSE-treated BMMs both at 24 h post-treatment. ^{a,b,c,d,e} A statistical significant results ($p < 0.05$) based on Scheffe *post hoc* one-way ANOVA analysis (each letter should be compared separately with the letters of other bars).

The expression of CD11b⁺ was not significantly increased under any experimental conditions (Figure 6a). However, the expression of the CD40⁺ costimulatory molecule for antigen presentation was significantly increased in PSE-treated cells, suggested that the effect of PSE may be related to the M1 macrophage phenotype and T cell immunity for anti-pathogenic immunity (Figure 6b). Accordingly, our results strongly suggest that the reduction of *E. coli* burden in the prostate of CBP mice may be related to the PSE treatment for 4 weeks.

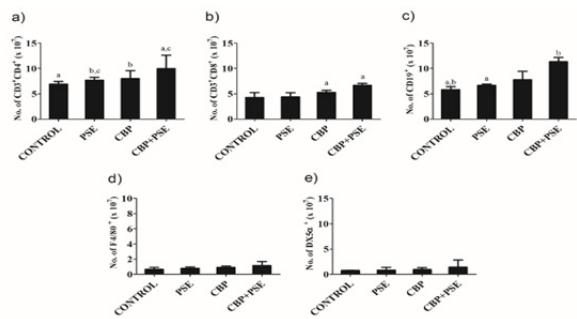


Figure 7: Changes in immune cell phenotypes in the spleens of CBP mice. PSE administration induced the increase in lymphocytes: CD3⁺CD4⁺-T cells (a), CD3⁺CD8⁺-T cells (b), and CD19⁺-B cells (c). Macrophages (F4/80) (d), NK cells (DX5a) (e). ^{a,b,c} A statistical significant results ($p < 0.05$) based on Scheffe *post hoc* one-way ANOVA analysis (each letter should be compared separately with the letters of other bars).

To investigate whether the reduction of bacterial burden in the CBP+PSE-group was related to immune enhancement, the phenotypes of splenocytes were examined by FACS analysis (Figure 7). PSE administration in non-infected mice resulted in a slight increase in CD4⁺ T cells, CD8⁺ T cells, and CD19⁺ B cells in comparison with control mice (CD3⁺CD4⁺: $7.7 \pm 0.57 \times 10^7$ vs. $6.9 \pm 0.53 \times 10^7$; CD3⁺CD8⁺: $4.4 \pm 0.85 \times 10^7$ vs. $4.3 \pm 0.95 \times 10^7$; CD19⁺: $6.7 \pm 0.002 \times 10^7$ vs. $5.8 \pm 0.58 \times 10^7$) (Figure 7a,b,c). The increase in lymphocytes was stronger in the CBP+PSE group in comparison with the CBP group (CD3⁺CD4⁺: $9.97 \pm 0.27 \times 10^7$ vs. $8.01 \pm 0.15 \times 10^7$; CD3⁺CD8⁺: $6.68 \pm 0.37 \times 10^7$ vs. $5.28 \pm 0.4 \times 10^7$; CD19⁺: $11.35 \pm 0.08 \times 10^7$ vs. $7.78 \pm 0.17 \times 10^7$) (Figure 7a,b,c). The number of F4/80-macrophages was also increased in CBP+PSE mice in comparison with CBP mice without a statistical significance ($1.16 \pm 0.52 \times 10^7$ vs. $0.92 \pm 0.16 \times 10^7$) (Figure 7d). In addition, the numbers of DX5a-NK cells were increased in CBP+PSE mice in comparison with CBP mice although there was no statistical significance ($1.41 \pm 1.41 \times 10^7$ vs. $0.92 \pm 0.16 \times 10^7$) (Figure 7e). The increased numbers of lymphocytes in PSE-administered CBP mice were likely related to the anti-bacterial immune response and helped to relieve inflammation by decreasing the *E. coli* burden in the prostate.

Prostate diseases include bacterial (CBP) and non-bacterial prostatitis (BPH) [1]. The therapeutic measures for CBP are mainly antibiotic prescription [6], whereas BPH is treated with α -adrenergic blocking agents and 5- α -reductase inhibitors [2], or often by phytotherapy using phytochemicals such as a polyphenolic flavonoid [4]. BPH therapy by using herbal remedies was attempted with the expectation of their diverse favorable effects such as antibacterial, anti-inflammatory and antioxidant activity [13]. In particular, because the BPH and prostate cancer occurrence in aged men cannot be predicted, general phytomedicines for the prostate continue to attract attention and are becoming more popular all over the world [5]. Resveratrol was introduced as a candidate nutritional substance for prostate cancer prevention [9]; recently, it was suggested to have many beneficial health effects including antioxidant, anti-inflammatory, anti-proliferative, pro-apoptotic, and anti-angiogenic [8]. Those health effects of resveratrol were first recognized when it was realized that red wine consumption results in a lower incidence of cardiovascular diseases in the French population [8, 15]. *Trans*-resveratrol is the biologically active form of resveratrol, and is present in the skin of grapes and in other fruits (such as raspberries, blueberries, and mulberries), as well as in Scots pine, Eastern white pine, and knotweed [8]. Particularly high concentrations of *trans*-resveratrol were found in peanuts and grapes [9]; resveratrol content is 3 $\mu\text{g/g}$, 0.6 $\mu\text{g/g}$, and 110.3 $\mu\text{g/g}$ in grapes, wine, and peanut sprouts, respectively [17]. According to a recent study that used PSE (38.17 mg/mL, 0.05% resveratrol), in which rats were fed a high-fat diet with the addition of PSE (30

mg/kg), PSE had an anti-obesity effect because it controlled the levels PPAR γ and adiponectin proteins in adipose tissue [17]. In another study, *trans*-resveratrol content in PSE obtained by dissolving peanut sprout powder with water was 54.2 $\mu\text{g/g}$ [20]. PSE processed by hot water extraction and concentration in the present study was rich in *trans*-resveratrol ($148.51 \pm 3.05 \mu\text{g/g}$). In the present study, we were interested in the beneficial health effect of PSE on bacterial and nonbacterial prostatitis because PSE is a natural source of resveratrol [8, 9]. PSE administration reduced bacterial proliferation to one-quarter in of that in CBP mice. Although PSE is not an antibiotic, highly concentrated resveratrol in PSE may have a role in restoration of prostate tissue through its anti-inflammatory and immune-enhancing effect [8, 15]. Biomedical effects of resveratrol observed in multiple biological systems include those related to immunity, for example the anti-inflammatory effect and effects on multiple immune functions of T cells [15]. The present study has also shown an increase in the immune cells, CD4 $^+$, CD8 $^+$, and CD19 $^+$ -lymphocytes, in PSE-treated mice. This suggests that the administration of PSE induces immune enhancement due to the presence of resveratrol. Under the same conditions, PSE had no toxicity as assessed by measuring body weight as well as liver and kidney toxicity.

BPH is also an important prostate disease and represents hormone-induced chronic inflammation that results from the infiltration of inflammatory cells; it may show no distinct clinical signs and generally lead to urination problems resulting from urinary tract obstruction by an enlarged prostate [13]. With respect to BPH, the usefulness of phytotherapy and complementary alternative medicine has been suggested with the expectation of the reduction in the risk of death and long-term morbidity [4, 5]. Experimentally, BPH can be induced by E2 and DHT treatment [2, 13, 25]. Fortunately, BPH can be relieved by herbal remedies due to their ability to inhibit cyclooxygenases 1 and 2 (COX-1 and COX-2) [13]. The resveratrol content in PSE is very high compared with a natural phytomedicine [17]. Similar to CBP mice, blood neutrophils were increased in BPH mice; PSE administration reduced the blood neutrophil population by approximately 4.3 percentage points compared to that in untreated BPH mice. At the same time, PSE-administered mice showed a reduction in epithelial cell hyperplasia and swelling in epithelium of the glandular tissues as well as the thickening and fibrous features in the stroma related to prostate hypertrophy. These results are in line with those of a recent study that investigated the effect of resveratrol on BPH induced by testosterone propionate injection [25]. Resveratrol (1 mg/kg) introduced into BPH rats showed favorable effects such as a decrease of prostate weight and DHT production [26]. In the present study, mice were treated with a much lower resveratrol concentration, 0.119 mg/kg, but it still showed the preventive and therapeutic effect in CBP and BPH-induced prostatitis.

The anti-prostatitis effect of PSE is likely to result in part from immune enhancement, because our ex vivo experiment showed activation of 95% pure BMMs by PSE. Activation of these antigen-presenting cells suggests the induction of further immune responses. CD40 is an M1 macrophage costimulatory marker, and M1 macrophages produce considerable amounts of nitric oxide in response to *E. coli* and have important roles in chronic inflammatory diseases [26]. In contrast, CD11b is a monocyte marker and a marker of CD8 $^+$ cytotoxic T cell activation during virus infection [27]. We found that PSE treatment increased CD40 expression on BMMs. This result agrees with the cell phenotype increased in the spleen. The numbers of T cells (CD4 $^+$ and CD8 $^+$), B cells (CD19 $^+$), and macrophages were increased more in the CBP+PSE group than in the CBP group; a decrease in *E. coli* burden in the prostate in the CBP+PSE group suggests the induction

of anti-bacterial immune response. Therefore, it seems that the increase in lymphocytes in PSE-administered mice with prostatitis enhanced the anti-bacterial immune response and helped to relieve inflammation.

Since PSE used in the present study was a mixture of natural compounds, it is impossible to definitely point out the ingredient that had the favorable effect in the prostatitis mouse model. However, peanut contains a high level of resveratrol, and resveratrol biosynthesis in peanut is enhanced by germination [24]; moreover, the extraction of peanut sprouts using steaming and ethanol is a processing method that yields highly concentrated resveratrol [19]. Recently, resveratrol has been reported to have an effect on BPH via regulation of inflammatory and apoptotic proteins [25]. On the basis of our results and the previous studies, we speculate that the effect of PSE on both CBP and BPH prostatitis may be due primarily to the presence of resveratrol.

Experimental

General: Peanut kernel, used for producing peanut sprout in this experiment, was purchased from Laixi city shunxiang peanut product's Co., LTD (shandong, china) and its scientific name is "*Arachis hypogaea L.*" (HS code: 12024200). *Arachis hypogaea L.* is registered in "Bio Resource Information Service center (Sejong-si, Republic of Korea) and its code is "MANUN1201292056". PSE was prepared by the established method with some modifications [17]. Briefly, peanut seeds were germinated by spraying and watering them dropwise for 7 to 8 days in the dark at 24–25°C and 85% humidity, and then processed at a local plant (Jangsuchae Co., Seoul, Korea) for further the preparation of liquid PSE. Liquid PSE was extracted by vacuum distillation and condensation with hot water extraction (100°C for 7 h), concentrated under reduced pressure (95°C for 48 h), diluted 10 times with 10 mM PBS (pH 7.2), filtered through a 0.22- μm PVDF syringe filter (Millipore, Billerica, MA), and then used for in vitro and animal experiment. The final PSE samples were orally introduced by 200 μL of PSE prepared in 100 mg/mL using gavage needle every day into CBP or BPH mice for 4 weeks. The composition of PSE was analyzed by the Korea Advanced Food Research Institute (KAFRI, Seoul, Korea).

Lipid, protein, ash, and moisture contents of PSE were determined using the standard methods of the Association of Official Analytical Chemists (AOAC, 2000; methods 960.39, 981.10, 920.153, and 950.46B, respectively). The total dietary fiber content was measured using a total dietary fiber assay kit (Sigma, St. Louis, MO, USA) according to the method of Prosky-AOAC (1985). Water-soluble vitamins (B1, B2, B3, and B6) were determined using HPLC according to the method of Ji et al. [28]. The contents of Ca, Na, P, Zn, Fe, Mg, K, and Mn were quantified by inductively coupled plasma atomic emission spectrometry (ICP-AES; Spectro Analytical Instruments, Kleve, Germany) according to the method of the Korean Food Standards Codex (Korea Food and Drug Administration, 2009). Resveratrol content of PSE were analyzed using 2695 HPLC system (Waters, Milford, MA) equipped with photodiode array detector. PSE was diluted in 80% MeOH and separated on a YMC-triart C18 column (4.6 x 250mm, 5 μm particle size; YMC, Kyoto, Japan) at 25°C. The flow rate was 1.0 mL/min and the injection volume was 10 μL . The gradient mobile phases were used with solvent (A) water:acetic acid (97.5:2.5 v/v) and solvent (B) acetonitrile:water:acetic acid (77.5:20:2.5 v/v). The gradient condition of the mobile phase was as follows: 0–5 min; solvent (B) 18%, 15 min; solvent (B) 30%, 25 min; solvent (B) 45%, 30 min; solvent (B) 50%, 35 min; solvent (B) 100%, 37 min; solvent (B) 18%. The absorbance was measured at 308 nm and trans-resveratrol (Sigma) was used as standard. Identification of resveratrol in PSE chromatogram was based on comparison with

spectra and retention time of standards. Trans-resveratrol concentration in PSE used in present study was $148.51 \pm 3.05 \mu\text{g/g}$.

Eleven-week-old BALB/c male mice purchased from Orient Bio (Seongnam, Korea) were housed in a specific pathogen-free barrier zone of Seoul National University College of Medicine animal facilities, and treated in accordance with the guidelines of the Institutional Animal Care and Use Committee (IACUC) at Seoul National University (IACUC No: SNU-140127-4-1) under approval for the use of animals for the present study. Mice were housed at 25 °C, 50% relative humidity and 12-h light/dark cycles, and sacrificed by CO₂ inhalation in a chamber after the experiment was finished. For the experiments on CBP and BPH, four experimental groups (5 mice per group) were used: control (no treatment), CBP (or BPH), PSE, and CBP+PSE (or BPH+PSE). PSE solution (200 μL, 100 mg/mL) was administered daily for 4 weeks from following day after *E. coli* was injected (or from that day when β-estradiol dissolved in sesame oil was treated). Body weight was examined every 4 days during the period of PSE administration.

E. coli 292 isolated from humans was kindly donated by Korea Prostate Bank (KPB, Seoul, Korea), and grown overnight in LB broth (Difco, Franklin Lakes, NJ). The genotype of the bacteria was confirmed by sequencing bacteria-specific 16S rRNA using universal 16S rRNA primers (forward primer: 5'-AGA GTT TGA TCM TGG CTC AG-3', reverse primer: 5'-ATT ACC GCG GCT GCT GG-3'). Sequencing was performed by SolGent Co., Ltd (Daejeon, Korea), and the results were analyzed using the Geneious Pro 5.5 software (Biomatters, Auckland, New Zealand).

CBP was induced as follows. 2×10^6 cfu of *E. coli* 292 per 20 μL was slowly injected into the urinary tract over 1 min using a catheter (20 mm in length, 0.61 mm in diameter) with 10-μL Hamilton syringe (Hamilton #80300, Reno, NV) under anesthesia with 30 mg/kg of Zoletil (Virbac Korea Co. Ltd., Seoul, Korea) plus 10 mg/kg of Xylazine (Rumpun; Bayer Korea Co. Ltd., Seoul, Korea) [6, 10]. Then the penis was tied with a sterilized string for 30 min to prevent the backflow of injected bacteria. The success of bacterial infection was confirmed by *E. coli* colony (cfu) counting in the prostate tissue of sacrificed mice at the end of the experiment. Briefly, tissues were removed from the ventral and dorsal lobes of the prostate, weighted, and ground in PBS using a blender in 50-mL conical tubes. The samples were spread on LB agar plates and cultivated at 37°C for 12 h. The bacterial colonies were counted by cfu $\times 10^5$ per 1 g of tissue. The colony number was approximately 2.4×10^5 cfu per gram of prostate tissue, indicating successful infection.

Prostate hyperplasia was induced by subcutaneous injection of β-estradiol (E2; Sigma-Aldrich) and 5α-androstan-17β-ol-3-one (dihydrotestosterone, DHT; Sigma-Aldrich) for 4 weeks [16]. For the first 2 weeks, β-estradiol dissolved in sesame oil (0.1 mL; 0.25 mg/kg/day) was injected. For the next 2 weeks, a mixture of β-estradiol and DHT dissolved in sesame oil (0.1 mL; 0.25 mg/kg/day each hormone) was injected.

To examine the safety of PSE as a dietary supplement, we examined the body weight, hepatotoxicity and renal toxicity. PSE solution (200 μL, 100 mg/mL) was administered for 2 weeks for three mice. Serum alkaline phosphatase (ALP) was measured using an ALP colorimetric assay kit (Abcam, Cambridge, UK) according to manufacturer's instructions. Kidneys were isolated from PSE-administered mice, and Kim-1 gene expression was examined by PCR amplification using specific KIM-1 gene primers (forward primer: 5'-GAG ATA CCT GGA GTA ATC ACA CTG AA-3', reverse primer: 5'-TGA TAG CCA CGG TGC TCA-3'). For PCR

reaction, total RNA was extracted using RNase kit (Qiagen, Hilder, Germany), and synthesized cDNA using oligo d(T)15 primer (EBT-1515, ELPIS biotech, Daejeon, Korea). 200 ng of cDNA synthesized was applied to PCR reaction. The PCR amplification was performed as follows: 2 min at 95 °C, (20 sec at 95 °C for denaturing, 40 sec at 64 °C for annealing, 30 sec at 72 °C for elongation) x 30 cycles, and 5 min at 72 °C. For PCR reaction on house-keeping gene (GAPDH), specific primers were used (forward: 5'-AGTCGGTGTGAAACGGATTG-3', reverse: 5'-TGTAGACC ATGTAGTTGAGGTCA'). The result was captured as gel image (DNR bio imaging, Jerusalem, Israel) after 2 % agarose gel electrophoresis and, and expressed as a fold-value comparing with GAPDH expression using Image J software (NIH, Bethesda, MD).

The prostate was isolated after the end of the experiment, fixed in 10% neutral formaldehyde for 3 days, and washed with tap water for 1 h. Tissues were embedded in paraffin (Leica TP1020; Leica Biosystems, Buffalo Grove, IL). Paraffin blocks were sliced into 7-μm sections on a microtome (Leica RM2145) and stained with H&E staining. Slides were observed and photos were taken using an optical microscope (Olympus, Tokyo, Japan).

Mice were treated orally with PSE (200 μL, 100 mg/mL) for 28 days. At the end of the experiment, mice were euthanized, and whole blood was collected before sacrifice into K2-EDTA tubes (BD Vacutainer, Franklin Lakes, NJ). Blood was thinly smeared on glass slides and stained using Diff-Quik (DQ) staining solution. A total of 100 white blood cells were counted per mouse by microscopic observation, and each cell population (neutrophils, monocytes, and lymphocytes) was counted separately.

Lipopolysaccharide (LPS) was extracted from *E. coli* 292 using an LPS extraction kit (Intron Biotechnology, Seoul, Korea). LPS from *E. coli* 292 was dissolved in HBSS to be concentration of 200 μg/mL, and stored until use. For in vitro cell cultures, LPS was used as a positive control at a final concentration of 1 μg/mL.

Bone marrow-derived macrophages (BMMs) were collected from the femur and tibia. Briefly, cells were collected by flushing the bone marrow with cold complete RPMI medium (Welgene Inc., Daegu, Korea) containing 10% fetal bovine serum (Invitrogen, Carlsbad, CA) and 1% antibiotic-antimycotic (Invitrogen), and then cultured in 30% L929-conditioned medium (M-CSF-containing medium) in a 100-mm dish (SPL, Pocheon, Korea) for 2 weeks with subculturing every 3 days. The purity of harvested BMMs was measured by using a FACSCalibur flow cytometer (BD Biosciences, San Jose, CA) after immunostaining with the pan-macrophage marker F4/80; the purity was confirmed to exceed 95%. For further in vitro culture, 2×10^5 BMMs were cultured with either PSE (5 mg/mL) or *E. coli* LPS (1 μg/mL) in a 6-well plate (SPL) for 24 h. CD11b- and CD40-expression level were determined by FACS analysis using anti-CD11b-FITC and anti-CD40-PE antibodies (both 1:100, eBioscience, San Diego, CA), respectively.

For investigating the inflammatory cell population, spleens were collected from CBP mice (n=5) at the end of the experiment, dissociated into single-cell suspensions, and washed with RPMI. Red blood cells (RBCs) in the spleen were destroyed by hypotonic shock using RBC lysis buffer containing 0.15 M NH₄Cl, 1 M KHCO₃, and 0.1 mM Na₂EDTA (pH 7.4). Splenocytes were counted by Trypan blue exclusion test, and were pooled in equal amount for further FACS analysis; 2×10^6 splenocytes for each cell population were analyzed in duplicate. The cells were incubated with rat anti-mouse CD16/32 (Fc III/II receptor) antibody (1:100

dilution, eBioscience) to block nonspecific binding before staining with specific antibodies, washed and stained with the following FACS antibodies (all from eBioscience; 1:100, 20 µL of each antibody for 10 min at room temperature in the dark): anti-CD19, phycoerythrin (PE)-conjugated; anti-F4/80, PE-conjugated; anti-DX5a, fluorescein isothiocyanate (FITC)-conjugated; anti-CD3, allophycocyanin (APC)-conjugated; anti-CD4, PE-conjugated; and anti-CD8, PE-cy5-conjugated. After washing, labelled cells were fixed in 20 µL of 4% paraformaldehyde solution. Fluorescence was quantified using a FACSCalibur flow cytometer (BD Biosciences). Data, which were originally expressed as percentage of splenocytes, were recalculated as cell number in each stained cell phenotype relative to the total spleen cell number.

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The results are presented as means ± SD. Statistical analysis was performed by a one-way ANOVA followed by Scheffe *post hoc* comparison. A statistical significance is represented by same lowercase letter. P < 0.05 was set as the level of statistical significance. The graphics program used in the present study is GraphPad Prism 5.0a.

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Phenanthrenes from *Eulophia macrobulbon* as Novel Phosphodiesterase-5 Inhibitors

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Phosphodiesterase 5 (PDE5) inhibitors can be used for the treatment of erectile dysfunction and pulmonary hypertension. In order to search for new leads of PDE5 inhibitors, we investigated the chemical constituents of the tubers of *Eulophia macrobulbon* (E.C. Parish & Rchb. f.) Hook. f. A new phenanthrene, 9,10-dihydro-4-(4'-hydroxybenzyl)-2,5-dimethoxyphenanthrene-1,7-diol (**1**) and three known phenanthrenes i.e., 1-(4'-hydroxybenzyl)-4,8-dimethoxyphenanthrene-2,7-diol (**2**), (9,10-dihydro-2,5-dimethoxyphenanthrene-1,7-diol (**3**) and 1,5,7-trimethoxyphenanthrene-2,6-diol (**4**) were isolated. Among these, **2** was the most potent PDE5 inhibitor ($IC_{50} = 1.67 \pm 0.54 \mu M$) evaluated by the [³H]cGMP radioassay method, whereas **1** showed mild activity ($IC_{50} = 62.3 \pm 3.3 \mu M$). Their inhibitory selectivities against PDE5 over PDE6 were also studied. This study suggests phenanthrenes as a new class of PDE5 inhibitors.

Keywords: *Eulophia macrobulbon*, Phenanthrenes, Phosphodiesterase-5 inhibitors, Erectile dysfunction.

Erectile dysfunction (ED) is the inability to achieve or sustain penile erection [1-4] and causes low self-esteem. Its prevalence is associated with aging and in the US 18 million men are affected [5] and the number is predicted to grow to 300 million globally by 2025 [6]. In Brazil, 45% of the male population over 18 years have some degree of ED [7]. PDE5 inhibitors, sildenafil (Viagra[®]), vardenafil (Levitra[®]), tadalafil (Cialis[®]) and Avanafil (Stendra[®]) have become first-line treatments [8-11]. The most important side-effects of PDE5 inhibitors is retinal defects via PDE6 inhibition [12-14]. Consequently, the search continues for both synthetic and herbal PDE5 inhibitors having high selectivity. We have screened Thai medicinal plants for PDE5 inhibitory activity, and *Eulophia macrobulbon* (E.C. Parish & Rchb. f.) Hook. f., a plant belonging to the Orchidaceae family, showed promising PDE5 inhibition. The present study aimed to isolate the biologically active phytochemicals of this plant, which hitherto have not been reported.

The separation of *E. macrobulbon* tuber extracts yielded four compounds (Figure 1). The HREIMS data of compound **1** exhibited a molecular ion peak at *m/z* 378.1478, which represented $[MH]^+$ for $C_{23}H_{22}O_5$. Its IR spectrum showed a typical characteristic band of a hydroxyl group at 3338 (O-H stretching) and aromatic bands at 1614, and 1516 cm^{-1} (C=C stretching). Its ¹³C NMR spectrum in MeOH-*d*₄ showed 21 resonances, which represented 23 carbon atoms. These were determined as two methyls, three methylenes, seven aromatic methines and eleven aromatic quaternary carbons by DEPT and HSQC experiments. The ¹H NMR spectrum indicated resonances for four protons at δ 2.04, 3.20, 2.38, and 2.61 (ddd), which are typical for methylene protons of a 9,10-dihydrophenanthrene derivative. Two methoxyl resonances were observed at δ 3.62 and 3.69. One singlet aromatic proton signal at δ 6.47 was found and a pair of doublets at δ 6.36 and 6.38 with a coupling

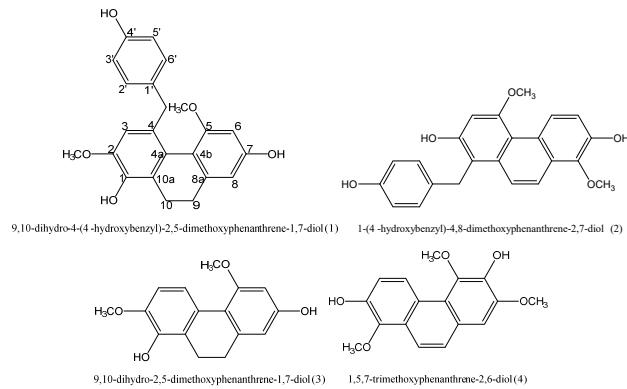


Figure 1: Structures of phenanthrenes isolated from *E. macrobulbon* extract.

constant of 2.0 Hz (*meta* coupling of H-6 and H-8, respectively). The presence of two doublets for a benzylic-CH₂ at δ 3.72 and 3.81 ($J = 15.0$ Hz) indicated a benzyl substitution. Two doublets (2H) at δ 6.81 and 6.61 with *ortho* coupling constants of 8.4 Hz were assigned to H-2'/H-6' and H-3'/H-5', respectively. These data suggested a *para*-disubstituted benzene. The ¹H and ¹³C NMR assignments were based on HMBC and NOESY experiments (Figure 2). The presence of a cross peak between H-3 and the benzylic-CH₂ in the HBMC experiment supported the connectivity between the 4' hydroxybenzyl group and the phenanthrene structure at C-4. The methoxy group positions in the structure were assigned using NOESY. The methoxy at δ 3.62 was located at C-5 since it displayed cross peaks with H-6, while the other methoxy (δ 3.69) showed a cross peak with H-3 in the NOESY experiment, suggesting that it was located at C-2. Finally, compound **1** was unambiguously identified as 9,10-dihydro-4-(4'-hydroxybenzyl)-

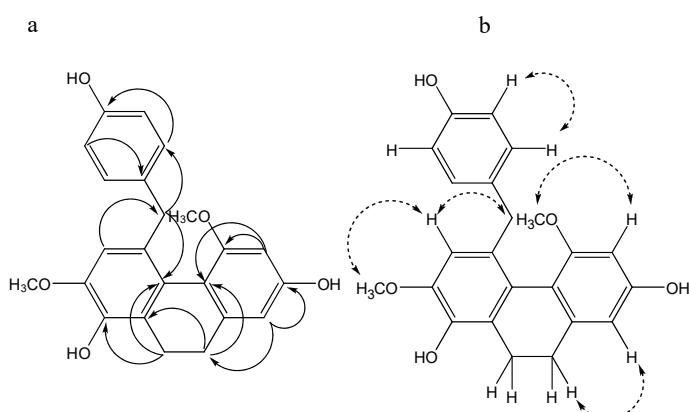


Figure 2: Diagnostic HMBC (a) and NOESY (b) correlations of 9,10-dihydro-4-(4'-hydroxybenzyl)-2,5-dimethoxyphenanthrene-1,7-diol (**1**).

2,5-dimethoxyphenanthrene-1,7-diol, a new 9,10-dihydrophenanthrene derivative. The structures of **2**, **3** and **4** were identified by spectroscopy and by comparing with published spectroscopic data [15-18].

Phenanthrene compounds **1**, **2** and **4** showed >70% inhibition against both PDE5 and PDE6 at 50 µg/mL (Table 1). Compound **2**, the main constituent, was the most potent PDE5 and PDE6 inhibitor, but less than sildenafil (Table 2). Even though PDE5/PDE6 selectivities were poor, they showed high selectivity over PDE1, which is abundant in the heart [19]. This appears to be the first study reporting phenanthrenes inhibiting PDEs and pharmacologically-active constituents of *E. macrobulbon*. Phenanthrenes may offer leads to synthesize more potent and selective PDE5 inhibitors. The limited number of compounds restricted interpretation of structure activity relationships. Nevertheless, the aromaticity of these compounds might be important for PDE5 and PDE6 inhibitory activities. Other *Eulophia* spp. studied have been *E. nuda* [15-16, 20-22], *E. petersii* [23] and *E. ochreata* [24-25], which also contain phenanthrenes. Their pharmacological properties include tumor cytotoxicity, and platelet anti-aggregation, anti-microbial, anti-inflammatory, and spasmolytic actions [22, 26]. PDE inhibition might underlie some actions, particularly those on smooth muscle.

Table 1: % Inhibitory activity of isolated phenanthrenes from *E. macrobulbon* on PDE5, PDE6 and PDE1

Comps	%Inhibitory activity at 50 µg/mL ^a		
	PDE-5	PDE-6	PDE-1
1	78.5 ± 3.6	70.1 ± 4.3	25.5 ± 1.3
2	99.2 ± 0.5	104.2 ± 0.2	0.6 ± 2.8
3	44.6 ± 4.1	66.2 ± 8.7	6.2 ± 3.8
4	72.3 ± 6.8	87.2 ± 4.8	5.8 ± 3.2

^aData are reported as the mean ± 1 S.D and are derived from three repeats.

Table 2: IC₅₀ values of isolated phenanthrenes from *E. macrobulbon* on PDE5 and PDE6

Comps	IC ₅₀ (µM) ^a	
	PDE-5	PDE-6
1	62.3 ± 3.3	94.1 ± 3.9
2	1.7 ± 0.5	1.8 ± 0.07
4	98.1 ± 13.3	51.0 ± 3.8
Sildenafil	0.03 ± 0.01	0.05 ± 0.02

^aData are reported as the mean ± 1 S.D and are derived from three repeats.

Experimental

General: ¹H and ¹³C Nuclear Magnetic Resonance (NMR) spectra were recorded in acetone-*d*₆, MeOH-*d*₄ and dimethylsulfoxide-*d*₆ on a Bruker AV400 (USA) spectrometer at 400 and 100 MHz, respectively. Mass spectra were obtained using an electron

ionization mass spectrometer (EI-MS) (MAT 95 XL, ThermoFinnigan, Germany). Infrared (IR) absorption spectra were recorded using the KBr technique on a Fourier Transform Infrared Spectrometer (Spectrum GX Series, PerkinElmer, USA). Optical rotations were measured in MeOH solution at the sodium D-line (589 nm) (Parameter 341, PerkinElmer). PDE inhibition was measured using a beta counter (TopCount NXT, (Perkin-Elmer, USA)

Plant material: *E. macrobulbon* was collected from Prachinburi Province, Thailand and identified by Asst. Prof. Dr Anupan Kongbangkerd, Faculty of Sciences, Naresuan University. The herbarium specimen (No. 002716) is kept in the Biology Department, Faculty of Science, Naresuan University, Thailand.

Chemicals: cGMP, crude snake venom (*Crotalus atrox*), histone from calf thymus, bovine serum albumin (BSA), ethylene glycol tetra-acetic acid (EGTA), imidazole, Tris, MgCl₂, DEAE-Sephadex, and bovine heart PDE1 were purchased from Sigma Chemical (St. Louis, MO, USA). [³H]cGMP was obtained from Perkin Elmer (Boston, MA, USA). Sildenafil was a gift from Bioequivalence Unit, Department of Pharmacology, Faculty of Medicine, Khon Kaen University, Thailand. The organic solvents (analytical grade) were purchased from Burdick & Jackson (B&J) (UK). All other reagents were bought from LabScan (Bangkok, Thailand). TLC aluminum sheets and silica gel 60 F254 were purchased from Merck (Darmstadt, Germany).

Extraction and isolation: The fresh tubers of *E. macrobulbon* were chopped and dried at 55°C. The dried material (2 kg) was ground into a fine powder, macerated in 95% EtOH (14 L) twice (3 days each time), and the solvent removed under reduced pressure to afford a brown syrupy EtOH extract (225 g). This (200 g) was dissolved in MeOH (200 mL) then partitioned with *n*-hexane (6 x 200 mL), CH₂Cl₂ (8 x 400 mL), EtOAc (8 x 400 mL) and *n*-BuOH (5 x 400 mL). All fractions were then dried under vacuum. PDE5 inhibitory activity was detected in the CH₂Cl₂ fraction, part of which (50 g) was loaded onto a column of SiO₂ and eluted with increasing proportions of MeOH in CH₂Cl₂, followed by elution with 1% 1N HCl in MeOH. Fractions were monitored by TLC (CH₂Cl₂/MeOH) (98:2) and light petroleum/EtOAc (1:1) and the fractions with similar chemical profiles were pooled. Thirteen pooled fractions (ED-Q1 - ED-Q13) were obtained. Fraction ED-Q4 (1.2 g) was further fractionated by CC over SiO₂, yielding 6 pooled fractions. ED-Q4-2 (37 mg) was subjected to SiO₂ preparative TLC (prep-TLC) with light petroleum/EtOAc (27:13) as a solvent system to afford **3** (24 mg). Fraction ED-Q6 (1 g) was separated by Sephadex LH-20 with MeOH to yield fraction ED-Q6-3 (100 mg) and ED-Q6-5 (200 mg). Fraction ED-Q6-3 was further purified by CC using SiO₂ and gradient elution with light petroleum and EtOAc to give 10 fractions. ED-Q6-3-4 (40 mg) was isolated by Sephadex LH-20 with MeOH to give fraction ED-Q6-3-4-3, which was finally purified by SiO₂ prep-TLC with *n*-hexane-EtOAc (3:7) as mobile phase yielding **1** (8 mg). Fraction ED-Q6-5 (200 mg) was further separated on a SiO₂ column using gradient elution with light petroleum and EtOAc to give **2** (110 mg). Fraction ED-Q2 (50 mg) was chromatographed over SiO₂ using *n*-hexane/EtOAc as mobile phase (7:3, isocratic system) to provide fractions ED-Q2-1 to 8. Fraction ED-Q2-8 was further purified by prep-TLC on silica gel using *n*-hexane/EtOAc (6:4) to afford **4** (2 mg).

9,10-Dihydro-4-(4'-hydroxybenzyl)-2,5-imethoxyphenanthrene-1,7-diol (**1**)

Pale yellow amorphous.

[α]_D²⁰: 0 (c 0.1, MeOH).

IR (KBr): ν_{max} 3412, 1612, 1512, 1463 cm^{-1} .

UV (MeOH) λ_{max} (log ϵ): 273 (5.48).

^1H NMR (400 MHz, in methanol- d_4): 2.04 (H, ddd, $J = 14.5, 14.7, 4.0$ Hz, H-10'), 2.38 (H, ddd, $J = 14.7, 14.0, 4.0$ Hz, H-9'), 2.61 (H, ddd, $J = 14.0, 4.0, 2.4$ Hz, H-9'), 3.20 (H, ddd, $J = 14.5, 4.0, 2.4$ Hz, H-10''), 3.62 (3H, s, C-5-OCH₃), 3.69 (3H, s, C-2-OCH₃), 3.73 (1H, d, $J = 15.0$ Hz, benzylic-CH₂), 3.81 (1H, d, $J = 15.0$ Hz, benzylic-CH₂), 6.36 (H, d, $J_{6,8} = 2.0$ Hz, H-6), 6.38 (H, d, $J_{8,6} = 2.0$ Hz, H-8), 6.47 (H, s, H-3), 6.61 (2H, d, $J_{3',2'} \text{ or } J_{5',6'} = 8.4$ Hz, H-3' and H-5'), 6.81 (2H, d, $J_{2',3'} \text{ or } J_{6',5'} = 8.4$ Hz, H-2' and H-6').

^{13}C NMR (100 MHz, methanol- d_4): 24.4 (CH₂, C-10), 32.2 (CH₂, C-9), 40.4 (benzylic-CH₂), 55.3 (C, C-5-OCH₃), 56.2 (C, C-2-OCH₃), 98.6 (CH, C-6), 107.7 (CH, C8), 112.1 (CH, C-3), 115.8 (CH, C-3' and C-5'), 117.3 (C, C-4b), 127.0 (C, C-10a), 127.4 (C, C-8a), 131.0 (CH, C-2' and C-6'), 132.7 (C, C-4a), 135.5 (C, C-1'), 141.0 (C, C-4), 144.4 (C, C-1), 147.1 (C, C-2), 156.0 (C, C-4'), 158.1 (C, C-7), 158.3 (C, C-5).

EIMS m/z (relative intensity): 378 [M]⁺ (100), 363 [M-15]⁺ (23), 347[M-31]⁺ (22), 335 [M-43]⁺ (11), 272 [M-106]⁺ (11), 107 [M-271]⁺ (19).

HREIMS m/z : 378.1478 (calcd. for C₂₃H₂₂O₅).

Enzyme preparation: PDE5 was obtained from rat lung tissues. Briefly, fresh tissue was cut into small sections and homogenized in 1 mL of Tris buffer (50 mM Tris, pH 7.5, 2 mM EDTA, 1 mM DTT and 1:100 of 100 mM phenylmethylsulfonyl fluoride). The homogenate was centrifuged at 14,000 rpm for 20 min at 4°C and the supernatant was used as a source of PDE5. The PDE5 inhibitor, sildenafil, was used to confirm that the supernatant contained abundant PDE5. All of the tissues were obtained from animals for which we received approval from the Institutional Animal Care and Use Committee (IACUC), Naresuan University (NU-AE570408). PDE6 was a gift from Prof. Dr Joseph A. Beavo, Department of Pharmacology, University of Washington, USA.

PDE inhibition assay: The method of Sonnenburg *et al.* [27] was used to assess PDE5 inhibition, with some modification. In brief,

the reaction mixture comprised 20 μL of Reagent A (100 mM Tris-HCl (pH 7.5), 100 mM imidazole, 15 mM MgCl₂, 1.0 mg/mL BSA and 2.5 mg/mL snake venom), 20 μL of 10 mM EGTA, 20 μL of PDE5 solution, and 20 μL of test sample, or only solvent (5% DMSO) as a control. The reaction was started by adding 20 μL of 5 μM [³H]cGMP (~50,000 cpm) and performed at 30°C for 40 min. Then, 100 μL of 50% DEAE resin was added to the reaction. After shaking for 10 min, the resin was allowed to settle (20 min). The supernatant was transferred to a fresh 100 μL of 50% DEAE resin, shaken for 10 min and the resin allowed to settle again. The supernatant (100 μL) was shaken with 200 μL of Microscint® 20 and tritium counted on a TopCount NXT (PerkinElmer, USA) for 2 h. The amount of PDE5 used hydrolyzed 20–25% of the substrate. Each experiment was duplicated in a 96-well plate. Inhibition of calmodulin-dependent PDE1 and PDE6 assays were modified from Sonnenburg *et al.* [27] and Huang *et al.* [28], respectively, similar to that for PDE5, but with Ca-calmodulin and histone in Reagent A. The PDE1 assay needed CaCl₂ (0.8 mM) and calmodulin (4 $\mu\text{g}/\text{mL}$) while histone (0.5 mg/mL) was needed for PDE6. In preliminary screening, 50 $\mu\text{g}/\text{mL}$ of plant extracts was tested and pure compounds tested at 10 μM final conc. All samples were dissolved in DMSO and diluted with water. DMSO was limited to 1% in the final assay medium. In the case of > 80% PDE inhibition, the IC₅₀ values were measured by the determinations of serial concentrations of the samples. The IC₅₀ values were calculated using Prism software (Graph Pad Inc., San Diego, CA). Sildenafil was the positive control.

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Bioactive Glucitol-Core Containing Gallotannins and other Phytochemicals from Silver Maple (*Acer saccharinum*) Leaves

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In the course of our group's investigation of members of the maple (*Acer*) genus, a series of glucitol-core containing gallotannins (GCGs) were isolated and identified (by NMR and HREISMS). Among higher plants, only certain maple species are known to produce GCGs, compounds with potential nutraceutical and cosmetic applications due to their reported antioxidant, antidiabetic, anti- α -glucosidase, anti-glycation, anticancer, and skin health promoting effects. Herein, we sought to investigate whether the previously un-investigated silver maple (*Acer saccharinum*) species was also a source of GCGs. Nine phenolic compounds, including six GCGs, were identified (by HPLC-DAD analyses using previously isolated standards) as ginnalins A-C (1-3), maplexins B, D, and F (4-6), methyl syringate (7), methyl gallate (8), and 3-methoxy-4-hydroxyphenol-1- β -D-(6-galloyl)-glucopyranoside (9). In addition, one sesquiterpenoid, namely, pubinernoid A (10), was isolated and identified (by NMR).

Keywords: Silver maple, *Acer saccharinum*, Phenolic, Glucitol-core containing gallotannins (GCGs).

The maple (*Acer*) genus contains thirteen species which are native to North America including the sugar (*A. saccharum*), red (*A. rubrum*), and silver (*A. saccharinum*) maples. Apart from sap which is primarily consumed as maple syrup [1], other maple plant parts (leaves, bark, etc.) have also been used as traditional medicines by the indigenous people of North America [2].

Our group has reported on the isolation and structure elucidation (by NMR and HREISMS) of over 70 phytochemicals, from the sugar, red, and sycamore (*A. pseudoplatanus*) maples [3-7]. Among these species, only the red maple yielded glucitol-core containing gallotannins (GCGs) [3, 4, 6]. These compounds have been reported, by our group [8-15], and others [16-20], to show a wide range of *in vitro* and *in vivo* biological effects including, antioxidant, anti-diabetic, anti- α -glucosidase, anti-glycation, anticancer, and skin-health promoting effects. Given the potential nutraceutical and cosmetic applications of these compounds, as well as our access to the aforementioned maple-derived phytochemical standards, herein, we sought to investigate whether the silver maple was a source of GCGs.

Using authentic standards (previously isolated by our laboratory), nine phenolics, including six GCGs, were identified from silver maple leaves as ginnalins A-C (1-3), maplexins B, D, and F (4-6), methyl syringate (7), methyl gallate (8), and 3-methoxy-4-hydroxyphenol-1- β -D-(6-galloyl)-glucopyranoside (9) (by HPLC-DAD analyses, see Supplementary data, Table S1) [6, 7]. In addition, one sesquiterpenoid, namely, pubinernoid A (10), was isolated and identified (by NMR; see Supplementary data, Figures S1 and S2). Pubinernoid A was obtained as a white amorphous powder and its ¹H and ¹³C NMR data were consistent with the literature [7]. The chemical structures of the compounds identified from silver maple leaves are shown in Figure 1.

In summary, this is the first reported phytochemical investigation of the silver maple which led to the identification of nine phenolic compounds (six GCGs and three phenolics) and a sesquiterpenoid. Our findings add to the growing body of data of phytochemicals

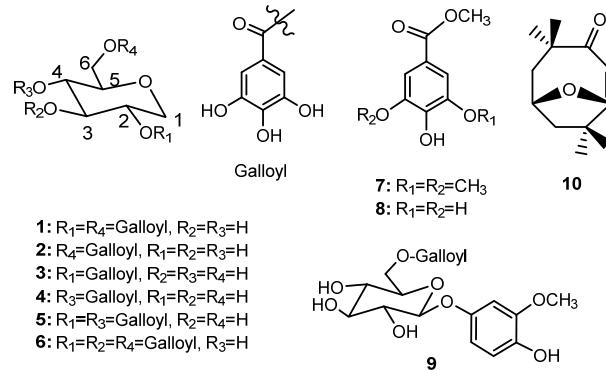


Figure 1: Chemical structures of compounds 1-10.

identified from the maple (*Acer*) genus, and more importantly, about which maple species produce GCGs, a promising class of bioactive plant polyphenols with nutraceutical and cosmetic applications.

Experimental

General: High performance liquid chromatography (HPLC) analyses were performed on a Hitachi Elite LaChrom system consisting of a L-2130 pump, a L-2200 autosampler, and a L-2455 Diode Array Detector. All ¹H and ¹³C NMR spectra were acquired on a Bruker 300 MHz spectrometer using MeOD₄ as solvent.

Plant material: Silver maple (*Acer saccharinum*) leaves were collected on the Kingston campus of the University of Rhode Island (Kingston, RI, USA) in the summer of 2010 and were botanically authenticated by Mr. J. Peter Morgan. Voucher specimens (16JPM1-APS6310) are deposited in the Heber-Youngken Garden and Greenhouse at the College of Pharmacy, University of Rhode Island.

Extraction of plant material: Air-dried silver maple leaves (1007.8 g) were macerated for 9 days with MeOH (5.5 L \times 3). The filtrate

was concentrated under reduced pressure to afford a dried methanol extract (250 g). This was suspended in distilled water and successively partitioned to yield *n*-hexanes (67.5 g), ethyl acetate (EtOAc; 28.7 g), and *n*-butanol (37.5 g) extracts after solvent removal *in vacuo*. The EtOAc extract, which contained the highest total polyphenol content (58.8% GAEs based on the Folin-Ciocalteau assay, see Supplementary data, Table S2), was selected for further phytochemical investigation (described below).

Identification of compounds 1-9 in the EtOAc extract of silver maple leaves by HPLC-DAD analyses: The EtOAc extract (5 mg) was analyzed on a Phenomenex C₁₈ column (250 × 4.6 mm, i.d. 5 µm). The injection volume was 10 µL and the flow rate was 0.75 mL/min. A gradient mobile phase solvent system consisting of solvent A (0.1% aqueous trifluoroacetic acid) and solvent B (MeOH) was used as follows: 0-50 min, 5%-46% B. The HPLC profile was monitored at 220 nm and compounds **1-9** were identified by comparison of their retention times to authentic standards previously isolated by our laboratory from red maple [3, 4, 6].

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Isolation of compound 10 from the EtOAc extract of silver maple leaves:

Having identified the major constituents in the EtOAc extract which were phenolics (**1-9**), we attempted to isolate the (minor) non-phenolic constituent/s (based on UV absorbance; see Supplementary Data, Table S1) present in the extract. Briefly, the EtOAc extract (28.2 g) was purified on a silica gel column eluted with a gradient solvent system of chloroform/MeOH (50:1 - 1:1; v/v) to afford 11 fractions (F1 - F11). Fraction F3 (440 mg) was chromatographed on a Sephadex LH-60 column eluting with MeOH to afford 6 sub-fractions (13A - 13F). Fraction 13B (340 mg) was chromatographed on a silica gel column eluting with a gradient solvent system of hexane/dichloromethane (10:0 - 0:10; v/v) to afford 5 fractions (16A - 17C). Fraction 16C (15 mg) was further purified by C₁₈ reversed-phase semi-preparative HPLC with a mobile phase consisting of MeOH and 0.1% aqueous trifluoroacetic acid (60:40; v/v) to afford compound **10** (3.4 mg, tR = 12.2 min).

Supplementary data: Supplementary data are available for this article and available free of charge.

Pterostilbene and Its Glucoside Induce Type XVII Collagen Expression

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The glycosylation of pterostilbene by cultured plant cells of *Phytolacca americana* gave pterostilbene 4'-O-β-D-glucoside. Both pterostilbene and its 4'-O-β-D-glucoside induced type XVII collagen expression in the EpiDermFT EFT-400 human skin cell model. Pterostilbene 4'-O-β-D-glucoside strongly induced type XVII collagen expression rather than pterostilbene.

Keywords: Glycosylation, *Phytolacca americana*, Pterostilbene, Collagen expression.

Recently, it has been reported that the aging of hair follicle stem cells is triggered by DNA damage induced type XVII collagen (COL17A1) proteolysis [1]. Once aged hair follicle stem cells are activated during the hair cycle, they leave the niche and terminally differentiate into epidermal keratinocytes and are then eliminated from the skin surface.

Glycosylation occurs readily in plant cells, i.e., many kinds of secondary metabolites such as saponins and anthocyanins are produced in the form of glycosides in higher plants. Many of these secondary metabolites exert specific physiological activities and have been widely used in folk medicines [2]. Therefore, the glycosides of natural compounds are of pharmaceutical importance. Although studies on the production of glycosides of stilbenes have been reported [3-5], there are few reports on their physiological activities.

Herein, we report, for the first time, the rapid induction of type XVII collagen (COL17A1) expression in hair follicle stem cells by a pterostilbene glucoside.

The level of type XVII collagen (COL17A1) was examined in the cells of the EpiDermFT EFT-400 human skin model treated with pterostilbene (**1**) and pterostilbene 4'-O-β-D-glucoside (**2**) (Figure 1), which had been prepared by glycosylation of pterostilbene with cultured cells of *P. americana*. One week after treatment of the cells of the EpiDermFT EFT-400 human skin model with pterostilbene (**1**), the COL17A1 level was increased in the cells (Figure 2). The level was enhanced dose dependently; the COL17A1 level of the 50 μM pterostilbene-treated cells (1.35) was higher than that of the 25 μM pterostilbene-treated cells (1.13). The level of the 50 μM pterostilbene-treated cells was elevated in the cells by ca. 1.35-fold, compared with that of control cells, but the 100 μM pterostilbene-treatment significantly reduced the COL17A1 level (0.69), probably due to the cytotoxicity of the pterostilbene.

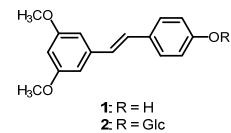


Figure 1: Chemical structures of pterostilbene (**1**) and pterostilbene 4'-O-β-D-glucoside (**2**).

A similar tendency was found in the case of pterostilbene 4'-O-β-D-glucoside (**2**) treatment of the EpiDermFT EFT-400 human skin cell model. At one week after treatment of the cells with pterostilbene 4'-O-β-D-glucoside (**2**), the COL17A1 level was elevated (Figure 3). The enhancement of the COL17A1 level was dose dependent; the COL17A1 level of the 50 μM treated cells (1.61) was higher than that of the 25 μM pterostilbene 4'-O-β-D-glucoside-treated cells (1.36). The COL17A1 level of the 50 μM pterostilbene 4'-O-β-D-glucoside-treated cells was elevated by ca. 1.6-fold, compared

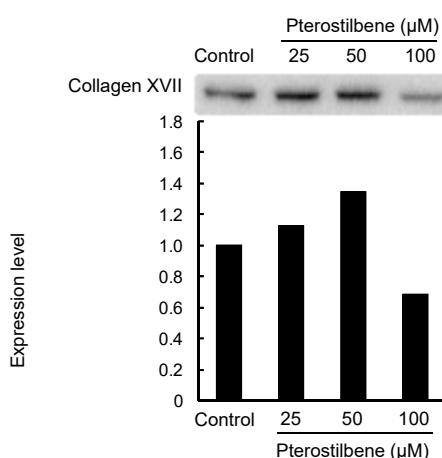


Figure 2: Effect of pterostilbene (**1**) on the expression of COL17A1 in EFT-400 3D human skin cell model.

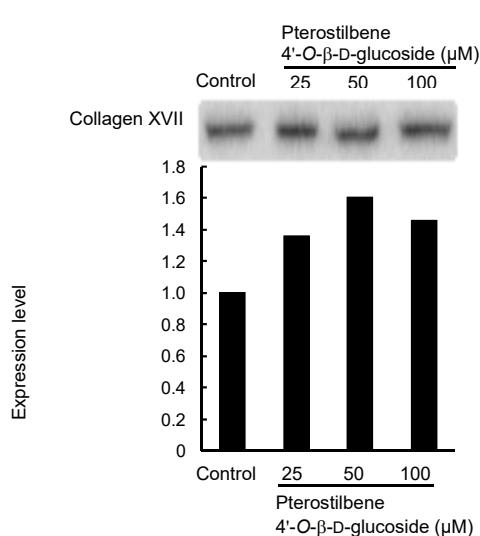


Figure 3: Effect of pterostilbene 4'-O-β-D-glucoside (2) on the expression of COL17A1 in EFT-400 3D human skin cell model.

with that of the control cells. However, the COL17A1 level of the 100 μM treated cells (1.46) was slightly decreased, but was higher than that of the 25 μM pterostilbene 4'-O-β-D-glucoside-treated cells.

Thus, both pterostilbene and pterostilbene 4'-O-β-D-glucoside, which had been prepared by glucosylation of pterostilbene with cultured cells of *P. americana*, quickly induced type XVII collagen (COL17A1) expression in cells of the EpiDermFT EFT-400 human skin model. This is the first report on the induction of COL17A1 expression by plant polyphenols. The COL17A1 level of pterostilbene 4'-O-β-D-glucoside-treated cells was higher than that of pterostilbene-treated cells. Stilbenes and, particularly, their glucosides may be useful for treatment of hair thinning.

Experimental

General: Pterostilbene used as a substrate was purchased from Wako Pure Chemicals Co. and was used without further purification. The cultured plant cells of *Phytolacca americana* were sub-cultured at 4-week intervals on a solid medium containing 2% glucose, 1 ppm 2,4-dichlorophenoxyacetic acid, and 1% agar (adjusted to pH 5.7) in the dark. A suspension culture was started by transferring 20 g of the cultured cells into 300 mL of liquid Murashige and Skoog medium in a 500 mL-conical flask.

Biotransformation procedure: The suspension plant cells (20 g) were incubated in conical flasks for 2 days. The cultured cells in the stationary growth phase were used for the experiments. After the cultivation period, 10 mg of the substrate was added. The biotransformation was performed by incubating the mixture at 25°C on a rotary shaker for 2 days. The culture medium and cells were separated by filtration. The culture medium was extracted with ethyl acetate. The cells were extracted (x3) by homogenization with methanol. The methanol fraction was concentrated and partitioned between H₂O and ethyl acetate. The ethyl acetate fractions were concentrated and analyzed by HPLC. The yield of product was determined on the basis of the peak area from HPLC using a calibration curve prepared with authentic glycosides.

Western blot analyses: EpiDermFT EFT-400 human skin culture cells (MatTek Corp., Ashland, MA, USA) were cultivated at 37°C in a 10% CO₂ atmosphere for 24 h before being mixed with 25, 50, and 100 μM of either pterostilbene or pterostilbene 4'-O-β-D-glucoside. After one week of incubation with the compounds, the cells were lysed, sonicated, and analyzed by Western blot analysis using anti-collagen XVII antibody (Abcam, Cambridge, MA, USA) and anti-GAPDH antibody (standard) (Sigma-Aldrich, St. Louis, MO, USA). The protein solution was fractionated by 10% sodium dodecyl sulfate polyacrylamide gel electrophoresis and transferred onto nitrocellulose membranes. Membranes containing fractionated proteins were blotted with anti-collagen XVII antibody. Membranes were then blotted with horseradish peroxidase conjugated second antibody and the immunoreactive protein bands were visualized by enhanced chemiluminescence.

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Metabolites from *Penicillium* sp. Associated with *Paris polyphylla*

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One new gentisyl alcohol derivative and seven known compounds were isolated from the culture of *Penicillium* sp. Their chemical structures were elucidated by extensive spectroscopic analysis. Compounds **1**, **4**, and **6** inhibited *Bacillus subtilis*, *Staphylococcus aureus*, and *Escherichia coli* with MICs of 32–128 µg/mL.

Keywords: *Penicillium* sp., Metabolite, Endophyte, Antimicrobial activity.

Paris polyphylla Sm., family Trilliaceae, a perennial herb mainly distributed in the Provinces of Yunnan, Sichuan and Guizhou in southwest China, has been used as an important and traditional Chinese medicine (TCM) for the treatment of injuries from falls, fractures, contusions, bleeding and immunity adjustment [1,2]. Endophytic micro-organisms live inside plants without causing them any apparent damage. Recently, endophytic microorganisms have attracted attention because they can produce bioactive compounds of biotechnological interest. The endophytic microorganisms in *P. polyphylla* are under scrutiny because they may be responsible for producing the bioactive metabolites associated with the plant [3]. As part of our program to discover structurally unique and biologically active secondary metabolites from fungi associated with medicinal plants, a *Penicillium* sp. isolated from the rhizosphere soil of *P. polyphylla* from Mile, Yunnan Province, China has drawn our interest. One new metabolite along with seven known compounds were isolated and determined as terrestrol I (**1**), (E)-ascladiol (**2**) [4], (Z)-ascladiol (**3**) [4], gentisyl alcohol (**4**) [5], 3-hydroxybenzyl alcohol (**5**) [6], (22E,24R)-ergosta-5 α ,6 α -epoxide-8,22-diene-3 β ,7 α -diol (**6**) [7], 5 α ,8 α -epidioxy-ergosta-6,22-dien-3 β -ol (**7**) [7], and 7-hydroxy-4,6-dimethylphthalide (**8**) [8].

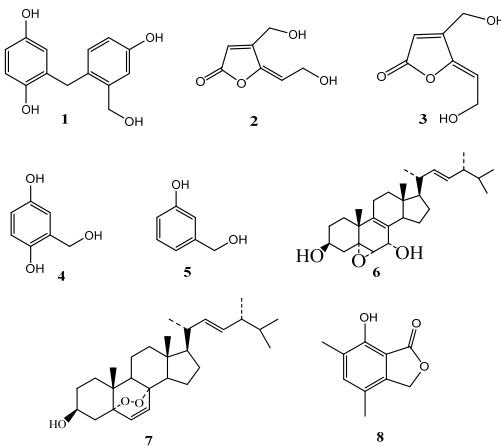


Figure 1: Structures of compounds 1-8.

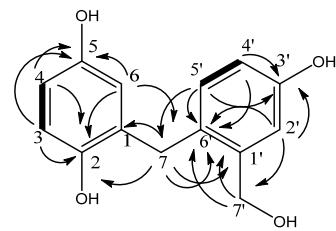


Figure 2: ¹H-¹H COSY correlations and the selected HMBC correlations of compound 1.

Table 1: ¹³C NMR (125 MHz) and ¹H NMR (500 MHz) data for compound 1 in CDCl₃.

Pos.	δ_{C}	δ_{H}	Pos.	δ_{C}	δ_{H}
1	130.7		1'	130.8	
2	149.7		2'	116.3	6.93 (s, H-2')
3	117.4	6.63 (d, $J=8.4$ Hz)	3'	157.6	
4	115.1	6.48 (dd, $J=8.4, 3.2$ Hz)	4'	115.8	6.63 (d, $J=8.8$ Hz)
5	151.9		5'	133.1	6.93 (d, $J=8.8$ Hz)
6	118.3	6.28 (d, $J=3.2$ Hz, H-6)	6'	142.5	
7	33.0	3.83 (s)	7'	63.7	4.53 (s)

The HR-ESI-MS analysis of compound **1** revealed a sodium adduct at *m/z* 269.0791[M+Na]⁺ (calcd for C₁₄H₁₄O₄Na 269.0790). The molecular formula was determined as C₁₄H₁₄O₄, which was further corroborated by the NMR data. The δ 6.0–7.0 region of the ¹H NMR spectrum of **1** (Table 1) showed resonances for three aromatic protons of the 1, 2, 5-trisubstituted phenyl ring A, at δ 6.28 (d, $J=3.2$ Hz, H-6), 6.48 (dd, $J=8.4, 3.2$ Hz, H-4), and 6.63 (d, $J=8.4$ Hz, H-3), as well as three aromatic protons of 1', 3', 6'-trisubstituted phenyl ring B at δ 6.63 (d, $J=8.8$ Hz, H-4'), 6.93 (d, $J=8.8$ Hz, H-5'), and 6.93 (s, H-2'). In addition, two singlets were observed at δ 3.83 and 4.53 for CH₂ at H-7 and H-7'. By comparing NMR data, the structure of compound **1** was shown to be similar to the gentisyl alcohol derivatives isolated previously from *Penicillium* [9]. The connection of ring A to ring B through C-7 was confirmed by the HMBC correlations from H-7 to C-1, 2, 1', and 6', and from H-5' and H-6 to C-7; C-7' was connected to C-1' based on the HMBC correlations from H-7' to C-1' and C-6', and from H-2' to C-7'. Three hydroxyls connected to C-2, 5, and 3' were determined by the HMBC correlations from H-3, 4, and 6 to C-2, and from H-2', H-4' and H-5' to C-3. Other HMBC correlations also determined the structure of the gentisyl alcohol derivative **1** (Figure 2).

Table 2: Antimicrobial activities (MICs, $\mu\text{g/mL}$) of compounds **1**, **4**, and **6** from *Penicillium* sp.

Strain	1	4	6	N	K
<i>B. subtilis</i>	64	32	64	—	8
<i>F. oxysporum</i>	—	—	—	1	—
<i>F. solani</i>	—	—	—	1	—
<i>S. aureus</i>	64	64	—	—	16
<i>M. albicans</i>	—	—	—	1	—
<i>E. coli</i>	—	128	—	—	64

—: MICs > 512 $\mu\text{g/mL}$; N: nystatin; K: kanamycin.

The antimicrobial activities (*Bacillus subtilis*, *Fusarium oxysporum*, *Fusarium solani*, *Staphylococcus aureus*, *Monilia albicans*, *Escherichia coli*) of compounds **1**, **4**, and **6** were evaluated. The MICs for these compounds are listed in Table 2.

Experimental

General experimental procedures: Silica gel (200–300 mesh; Qingdao Marine Chemical Group Co., Qingdao, China), and Sephadex LH-20 (GE Healthcare Co., Buckinghamshire, UK) were used for column chromatography (CC). 1D and 2D NMR spectra were obtained on a Bruker AVANCE 500 MHz NMR instrument (Bruker, Karlsruhe, Germany). MS were recorded with an Agilent G3250AA (Agilent, Santa Clara, USA).

Biological material and cultivation of strain Y-12: The endophytic fungus was isolated by using PDA medium from *Paris polyphylla* collected at Mile, Yunnan, China, and was identified as *Penicillium* sp. by ITS gene sequence. The strain is preserved at the School of Chemical Science and Technology, Yunnan University, Kunming, China. The strain was cultured in 500 mL Erlenmeyer flasks each containing 200 mL potato dextrose broth (PDB) seed culture (1000 mL water, 20 g glucose, 200 g potato). After 3 days of incubation at 28°C on a rotary shaker (125 r.p.m.), 10 mL portions of the culture were inoculated into 1000 mL Erlenmeyer flasks, each containing 250 mL PDB fermentation medium for 7 days at 28°C on a rotary shaker (150 r.p.m.). The fermented whole broth (30 L) was filtered through hospital gauze to separate the culture broth and mycelia.

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Antibacterial assays: For the *in vitro* antimicrobial test, PDB was used as an incubation medium for fungi and LB medium for bacteria. The final volume of each well was 100 μL . Aliquots (5 μL) of the metabolite solutions in dimethyl sulfoxide (DMSO) were added into the 96-well sterilized microplates, and their final concentrations ranged from 512 to 1 $\mu\text{g/mL}$ using a twofold serial dilution method. Wells containing kanamycin and nystatin (Taicheng Pharmaceutical Co., Ltd, Guangdong, China) were introduced as the positive controls. The minimum inhibitory concentrations of the metabolites that completely inhibited pathogenic bacteria growth were defined as the MICs.

Extraction and isolation of compounds: The fermentation broth was extracted 3 times with EtOAc to give a crude residue (25 g). This was separated by silica gel CC with $\text{CHCl}_3/\text{MeOH}$ (1:0–0:1) to give 5 fractions (Fr. 1 to Fr. 5). In the further purification, silica gel CC with light petroleum/ethyl acetate, and $\text{CHCl}_3/\text{MeOH}$ were used. Finally, compound **1** (14.5 mg) was obtained from Fr. 2. The mycelium extract (12.8 g) was separated by silica gel CC with $\text{CHCl}_3/\text{MeOH}$ to give 5 fractions (Fr. 1 to Fr. 5), Compound **8** (3.4 mg) was isolated from Fr. 2–1 by silica gel CC with light petroleum/ethyl acetate, and $\text{CHCl}_3/\text{MeOH}$. Compounds **2** (2 mg), **3** (2 mg), **4** (9.2 mg), and **5** (12.4 mg) were isolated from Fr. 2–3 using silica gel CC with $\text{CHCl}_3/\text{MeOH}$ and Sephadex LH-20 (MeOH). Compounds **6** (6.7 mg), and **7** (10.3 mg) were obtained from Fr. 2–2 using silica gel CC with $\text{CHCl}_3/\text{MeOH}$ and Sephadex LH-20 (MeOH).

Terrestrol I

^1H and ^{13}C NMR: Table 1.

HR-ESIMS m/z : 269.0791 [M+Na]⁺ (calcd for $\text{C}_{14}\text{H}_{14}\text{O}_4\text{Na}$ at 269.0790).

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Ferulic Acid Esters of Oligo-glucose from *Allium macrostemon*

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Two new ferulic acid esters of oligo-glucose, 1-O-(E)-feruloyl- β -D-gentiobioside (**1**) and 1-O-(E)-feruloyl- $\{\beta$ -D-glucopyranosyl (1 \rightarrow 6)-[β -D-glucopyranosyl (1 \rightarrow 2)]}- β -D-glucopyranoside (allimacronoid D, **2**) were isolated together with 1-O-(E)-feruloyl- β -D-glucopyranoside (**3**) and *trans*-ferulic acid (**4**) from the leaves of *Allium macrostemon* Bunge. The chemical structures were elucidated based on the analyses of the spectroscopic and chemical data.

Keywords: Amaryllidaceae, *Allium macrostemon*, Allimacronoid, Ferulic acid, Glucoside.

Allium macrostemon Bunge (Amaryllidaceae), known as wild onion, is widely distributed in East Asian countries. Some steroidal saponins [1] in the bulb showed various activities with the potential for treatment of acute myocardial ischemia, hyperglycemia, hyperlipidemia and visceral obesity [2-4]. An antidepressant-like activity of the aqueous extract of the plant was also reported [5]. Nakane *et al.* reported the isolation and HPLC analysis of the flavonoids in the leaves of *A. macrostemon* [6]. Recently, Usui *et al.* also isolated ferulic acid esters with oligo-glucose (allimacronoids A-C) [7], 1-O-(E)-caffeooyl- β -D-sophoroside [8] and kaempferol glycosides [8] from the leaves of this species. In the present study of the phenolic compounds of the leaves of *A. macrostemon*, two new ferulic acid esters with oligo-glucose (**1** and **2**) were isolated, together with two known phenolics (**3** and **4**). The chemical structures of these compounds were elucidated from their spectroscopic and chemical data. The known phenolics **3** and **4**, were identified as 1-O-(E)-feruloyl- β -D-glucopyranoside (**3**) [9] and *trans*-ferulic acid (**4**) [10], by comparison of their spectroscopic data (Figure 1).

Compound **1** gave a quasi-molecular ion peak at *m/z* 517.1584 [M-H]⁻, which corresponded to the molecular formula C₂₂H₃₀O₁₄. The ¹H NMR spectrum (Table 1) showed ABX-type (δ 6.81, *J*=8.4 Hz, 1H; δ 7.10 *J*=1.8, 8.4 Hz, 1H and δ 7.21, *J*=1.8 Hz, 1H) aromatic proton signals, together with *trans*-olefinic (δ 6.40 and 7.73, *J*=15.6 Hz, each 1H), methoxyl (δ 3.89, 3H), and two sugar anomeric (δ 5.55, *J*=7.6 Hz and δ 4.33, *J*=7.6 Hz) proton signals. In the ¹³C-NMR spectrum of **1** (Table 1), 9 carbon signals of an aglycone (δ 112.0, 114.8, 116.6, 124.5, 127.7, 148.5, 149.6, 151.1 and 167.8) with a methoxyl signal (δ 56.5), which were attributable to ferulic acid [9, 10], were observed, as well as other sugar signals (12 carbons), which indicated the presence of two hexose moieties. Acid hydrolysis of **1** with 0.5 N HCl yielded an aglycone component, which was identified as *trans*-ferulic acid (**4**) by HPLC analysis. The hexose units of **1** were also identified as D-glucose by the method of Tanaka *et al.* [11]. Therefore, **1** was supposed to be a diglucoside of *trans*-ferulic acid. The NMR data of **1** were very similar to those of phenylpropanoid esters of gentiobioside such as

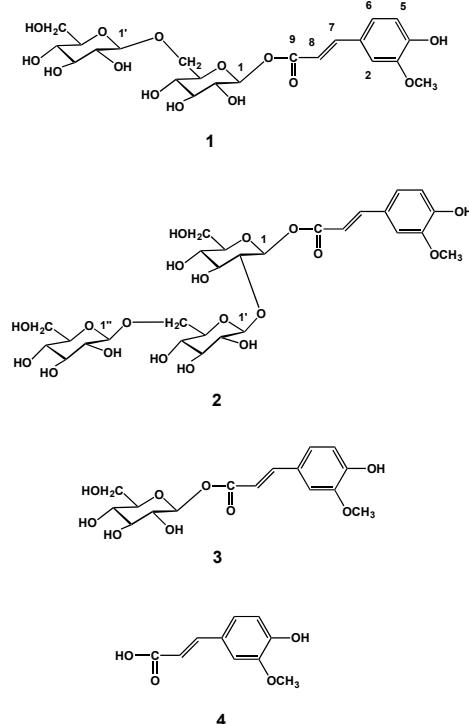


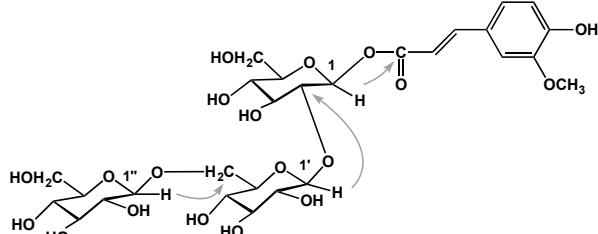
Figure 1: Chemical structures of **1-4**.

1-O-(E)-caffeooyl- β -D-gentiobioside [12], 1-O-(E)-cinnamoyl- β -D-gentiobioside [13] and 1-(3'', 4''-dihydroxy-5''-methoxyl)-O-(E)-cinnamoyl- β -D-gentiobioside [14], with the only difference being the existence of ferulic acid as an aglycone moiety. The large *J* values (7.6 Hz) of the two anomeric proton signals in the ¹H NMR spectrum of **1** (Table 1) also confirmed the β configurations. From the data mentioned, **1** was concluded to be as 1-O-(E)-feruloyl- β -D-gentiobioside (Figure 1).

Table 1: ^{13}C - and ^1H -NMR data for **1** and **2** (δ , CD₃OD).

Compound Position	1 ^{13}C	1 ^1H	2 ^{13}C	2 ^1H
Ferulic acid				
1	127.7		127.5	
2	112.0	7.21 d (1.8)	112.0	7.23 d (1.9)
3	149.6		149.5	
4	151.1		151.0	
5	116.6	6.81 d (8.4)	116.6	6.83 d (8.2)
6	124.5	7.10 dd (1.8, 8.4)	124.5	7.14 dd (1.9, 8.2)
7	148.5	7.73 d (15.6)	148.2	7.71 d (15.9)
8	114.8	6.40 d (15.6)	115.1	6.41 d (15.9)
9	167.8		167.4	
OMe	56.5	3.89 s	56.6	3.91 s
Glucose-1				
1	95.9	5.55 d (7.6)	94.5	5.73 d (8.1)
2	74.0	3.42 m	81.8	3.78 m
3	78.0	3.31 m	77.6	3.66 m
4	71.0	3.44 m	70.8	3.46 t (8.0)
5	78.0	3.58 m	78.8	3.41 m
6	69.6	3.77 dd (11.6, 5.2)	62.3	3.71 m
		4.17 br. d (11.6)		3.85 dd (11.8, 2.0)
Glucose-2				
1'	104.6	4.33 d (7.6)	104.9	4.60 d (7.8)
2'	75.1	3.20 t (7.6)	75.8	3.18 m
3'	77.8	3.30 m	77.7*	3.36 m
4'	71.5	3.46 m	71.5	3.26 m
5'	77.9	3.25 m	77.3	3.40 m
6'	62.7	3.64 dd (12.0, 4.8)	69.4	3.71 m
		3.83 dd (12.0, 1.6)		4.00 dd (11.7, 2.0)
Glucose-3				
1''		104.5	4.42 d (7.8)	
2''		75.0	3.16 m	
3''		77.6*	3.34 m	
4''		71.3	3.30 m	
5''		77.8	3.24 m	
6''		62.7	3.62 m	
			3.78 m	

The numbers in parentheses are J values in Hz. * assignments may be interchangeable.

**Figure 2:** The selected HMBC (^1H - ^{13}C) correlations of **2**.

Compound **2** showed a quasi-molecular ion peak at m/z 679.2130 [$\text{M}-\text{H}$]⁺ (corresponded to the molecular formula $\text{C}_{28}\text{H}_{40}\text{O}_{19}$). The ^1H - and ^{13}C -NMR spectral data of **2** (Table 1), the assignments of which were also certified by COSY, HSQC and HMBC spectra, were similar to those of **1**, with the difference being the signals of an additional hexose moiety, suggesting **2** to be a ferulic acid ester of a triglycoside. Acid hydrolysis of **2** yielded *trans*-ferulic acid (**4**), which was identified by HPLC analysis. The sugar components of **2** were also identified as D-glucose by HPLC analysis [11]. The positions of the glucoses were confirmed by the HMBC correlations between glucose H-1 to ferulic acid C-9, glucose H-1' to glucose C-2 and glucose H-1'' to glucose C-6' (Figure 2). The J values of the three anomeric proton signals in the ^1H NMR spectrum of **2** (Table 1) indicated β configurations of the anomeric carbons. From the data described above, **2** was concluded to be 1-*O*-(*E*)-feruloyl- $\{\beta$ -D-glucopyranosyl (1 \rightarrow 6)-[β -D-glucopyranosyl (1 \rightarrow 2)] $\}-\beta$ -D-glucopyranoside, and named as allimacronoid D (Figure 1).

Although ferulic acid is commonly distributed in *Allium* species [15], the ester derivatives with oligo-glucose are rarely found. Han *et al.* reported tuberonoid A [1-*O*-(*E*)-feruloyl- β -D-sophoroside] and

tuberonoid B [1-*O*-(*Z*)-feruloyl- β -D-sophoroside] from *A. tuberosum* [16]. Including the reported compounds allimacronoids A-C [7], it is very noteworthy that *A. macrostemon* contains various types of ferulic acid esters of oligo-glucose.

Experimental

General: Nuclear magnetic resonance (NMR) spectra were recorded in CD₃OD using JEOL JNM-A500 (500 MHz for ^1H NMR and 125 MHz for ^{13}C NMR) and JEOL JNM-A400 spectrometers (400 MHz for ^1H NMR and 100 MHz for ^{13}C NMR). Optical rotation was measured with a JASCO DIP-1000 Digital Polarimeter, and UV spectra using a JASCO V-530 UV/VIS spectrophotometer. Negative-ion HRESI-TOF-MS were recorded on an Agilent Technologies 6540 Accurate-Mass Q-TOF LC/MS spectrometer. For column chromatography, DIAION HP20SS (Mitsubishi Chemical Corporation), Sephadex LH20 (Pharmacia Corporation), TOYOPEARL HW-40F (Tosoh Corporation) and Cosmosil 140C18-OPN (Nacalai Tesque Inc.) were used. Preparative HPLC was performed on a JASCO preparative HPLC system (Pump: PU-2086 Plus, Detector: UV-2075 Plus, detected at 254 nm) equipped with a COSMOSIL Cholester (10 mm *i. d.* x 250 mm, Nacalai Tesque Inc.) column.

Plant material: The leaves of *Allium macrostemon*, grown in the campus of Saga University, Japan, were collected in October 2015. The plant material was identified by one of the authors (K. Ohshima, professor of the laboratory of Plant Virology, Saga University). A voucher sample is deposited in the laboratory of Analysis of Plant Metabolism at the Faculty of Agriculture, Saga University.

Extraction and isolation: The fresh leaves (1800 g) of *Allium macrostemon* were extracted with 60% aqueous EtOH (7 L) for 2 days. The solvent was concentrated *in vacuo* and subjected to a column of Cosmosil 140C18-OPN, which was eluted with H₂O containing increasing amounts of MeOH to afford 6 fractions {Fr. 1 (20 g), Fr. 2 (25 g), Fr. 3 (10 g), Fr. 4 (8 g), Fr. 5 (8 g) and Fr. 6 (10 g)}. Fr. 1 was purified by DIAION HP20SS (stepwise elution with H₂O and MeOH) CC to afford 3 fractions (Fr. 1-1 and Fr. 1-3). Fr. 1-2 (1 g) was applied to a column of TOYOPEARL HW-40F (stepwise elution with H₂O and MeOH), followed by preparative HPLC separation on COSMOSIL Cholester (elution with 30% aqueous MeOH, 3 mL/min) to give **1** (2 mg, Rt=13.5 min). Fr. 1-3 (2 g) was purified by Sephadex LH-20 (80% aqueous EtOH) and DIAION HP-20SS (stepwise elution with H₂O and MeOH) CC to give **4** (14 mg). Fr. 2, on DIAION HP-20SS (stepwise elution with H₂O and MeOH) CC, afforded 3 fractions (Fr. 2-1 to Fr. 2-3). Fr. 2-1 (0.5 g) was purified by TOYOPEARL HW-40F (stepwise elution with H₂O and MeOH) CC, followed by preparative HPLC separation with COSMOSIL Cholester (elution with 12% aqueous CH₃CN, 3 mL/min) to give **2** (2 mg, Rt=13.5 min). Fr. 2-3 (0.5 g) was purified by TOYOPEARL HW-40F (stepwise elution with H₂O and MeOH) CC, followed by preparative HPLC separation with COSMOSIL Cholester (elution with 15% aqueous CH₃CN, 3 mL/min) to give **3** (2 mg, Rt=15.0 min).

HPLC-QTOF-MS analysis of **1 and **2**:** Negative-ion HRESI-TOF-MS of **1** and **2** were obtained in the HPLC-QTOF-MS analysis using the following conditions. HPLC system (Agilent 1100 series); Column: Agilent Eclipse Plus C18 (2.1 mm *i. d.* x 100 mm, 3.5 μm), Mobile phase: 0.1% formic acid with 2.5 mM AcONH₄-CH₃CN [100 : 0 (0 min) \rightarrow 0 : 100 (20 min)], Flow rate: 0.2 mL/min, Column temperature: 40°C, Detection: 280 nm (UV). Retention times (min): **2** (3.9) and **1** (4.2). TOFMS system (Agilent

G1969A); Ionization: ESI (negative), Drying gas: N₂, 350°C, 10 L/min, Nebulizer gas: N₂, 50 psig, Capillary: 4000 V, Fragmentor: 150 V, Mass range: 50-1200 m/z.

1-O-(E)-feruloyl-β-D-gentiobioside (1)

Off-white amorphous powder.

[α]_D : -28.6 (c 0.09, MeOH).

UV_{λmax} (MeOH) nm (log ε): 330 (4.29).

¹H NMR (400 MHz, CD₃OD): Table 1

¹³C NMR (100 MHz, CD₃OD): Table 1

HRESI-TOF-MS (negative-ion mode) m/z: 517.1584 [M-H]⁻ (Calcd for C₂₂H₂₉O₁₄: 517.1557).

Allimacronoid D (2)

Off-white amorphous powder.

UV_{λmax} (MeOH) nm (log ε): 330 (4.18).

[α]_D : -30.1 (c 0.22, MeOH).

¹H NMR (500 MHz, CD₃OD): Table 1

¹³C NMR (125 MHz, CD₃OD): Table 1

HRESI-TOF-MS (negative-ion mode) m/z: 679.2130 [M-H]⁻ (Calcd for C₂₈H₃₉O₁₉: 679.2086).

Identification of the sugar moieties of 1 and 2: Identification of the sugar moieties of **1** and **2** was made by the following analytical method of Tanaka *et. al.* [11]. Solutions of **1** and **2** (0.5 mg, respectively) in 0.5 M HCl (0.1 mL) were heated at 95°C in screw-capped vials for 2 h. The mixtures were neutralized by addition of

Amberlite IRA400 (OH- form) and filtered. The filtrates were dried *in vacuo*, dissolved in 0.1 mL of pyridine containing L-cysteine methyl ester (5 mg/mL) and reacted at 60°C for 1 h. To the mixtures a solution (0.1 mL) of o-torylisothiocyanate in pyridine (5 mg/mL) was added and they were heated at 60°C for 1 h. The final mixtures were directly analyzed by HPLC [COSMOSIL Cholester (4.6 mm i. d. x 250 mm, Nacalai Tesque Inc.); mobile phase, 0.1% formic acid - CH₃CN (9:1 to 1:4 in 30 min); flow rate, 0.6 mL/min; column temperature, 40°C; detection, 250 nm]. The tRs of the peaks at 28.65 min (from **1**) and at 28.60 min (from **2**) coincided with that of D-glucose (28.58 min, *cf.* tR of the peak for L-glucose was 28.23 min).

Acid hydrolysis of 1 and 2: Solutions of **1** and **2** (0.1 mg, respectively) in 0.5 M HCl (0.1 mL) were heated at 90°C in screw-capped vials for 1 h. The mixtures were directly analyzed by HPLC [COSMOSIL Cholester (4.6 mm i. d. x 250 mm, Nacalai Tesque Inc.); mobile phase, 0.1% formic acid - CH₃CN (9:1 to 1:4 in 30 min); flow rate, 0.6 mL/min; column temperature, 40°C; detection, 280 nm]. The tRs of the peaks at 25.50 min (from **1**) and at 25.49 min (from **2**) coincided with that of *trans*-ferulic acid (**4**, 25.24 min).

Supplementary data: ¹H and ¹³C NMR spectral data for compounds **1-2**, COSY, HSQC and HMBC NMR for compound **2** and TOFMS for compounds **1** and **2** are also available.

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Quantitative Analysis of Phenolic Acids and Antiplatelet Activity of *Melissa officinalis* Leaf Extracts

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The contents of total phenolic acids and rosmarinic acid were determined and the antiplatelet potential assessed of extracts of lemon balm (*Melissa officinalis* L., Lamiaceae) leaf samples collected during 2014 and 2015. The quantity of total phenolic acids ranged from 4.8 to 8.5%, while the rosmarinic acid content ranged from 1.78 to 4.35%. Results of statistical analysis showed that the undiluted extracts have antiplatelet activity in the ADP test (final concentration 31.25 mg/L). While there are reports on antiplatelet activity of lemon balm essential oil, to the best of our knowledge, this is the first report on antiplatelet activity of lemon balm leaf extracts.

Keywords: *Melissa officinalis*, Polyphenols, HPLC, Spectrophotometry, Antiplatelet activity.

Lemon balm (*Melissa officinalis* L., Lamiaceae) is an endogenous plant of western Asian regions and the Eastern Mediterranean, and is cultivated in almost all of Europe and North America [1]. It grows as a shrub, 0.3 to 0.9 m high, usually with several stems. The major chemical constituents are polyphenolic acids namely rosmarinic (most abundant), *p*-coumaric, caffeic and chlorogenic acids. Lemon balm essential oil (0.02–0.37%) is composed of more than 40% monoterpenes and more than 35% sesquiterpenes [2]. Other constituents determined in leaf samples include flavonoids, tannins and triterpenes [3]. Externally, lemon balm extracts are used for symptomatic treatment of herpes labialis while orally applied extracts act as a carminative and a sedative [1].

Antiplatelet activity, i.e. prevention of platelet aggregation, of polyphenols is of interest as it has been shown that they can act in submicromolar concentrations on platelet aggregation. This is a useful characteristic as polyphenols could serve as new sources of antiaggregatory drugs in prevention of cardiovascular diseases (e.g. prevention of thrombosis, heart attack or stroke). On the other hand, polyphenols from everyday diet could interfere with platelet functional tests and subsequently hinder diagnosis of platelet related disorders [4]. It has been shown that lemon balm essential oil has antiplatelet activity [5] that can be attributed to major terpene constituents [6,7]. As major constituents of lemon balm leaves are phenolic acids, primarily rosmarinic acid, the objective of this work was to determine the contents of total phenolic acids and rosmarinic acid, and the antiplatelet activity of ethanolic lemon balm leaf extracts.

Total phenolic acid concentrations ranged from 4.8 to 8.5% (Table 1) fulfilling the requirements of European Pharmacopoeia (more than 4% of phenolic acids). Rosmarinic acid, as a major constituent, was determined by HPLC and its content ranged from 1.78 to 4.35% (Table 1).

Lemon balm ethanolic extracts (non-diluted, 10%, 1%) were tested for antiaggregatory activity in a thrombocyte functional test induced with adenosine diphosphate (ADP). Statistical analysis (one-way

Table 1: Content of total phenolic acids and rosmarinic acid.*

Sample	Total phenolic acids	Rosmarinic acid
1	6.5 ± 0.5	1.78 ± 0.02
2	7.1 ± 0.1	2.14 ± 0.02
3	4.8 ± 0.1	1.99 ± 0.01
4	8.5 ± 0.2	4.35 ± 0.02
5	6.5 ± 0.1	2.30 ± 0.02

*Results are expressed as percentage (± SD) of dry mass of leaf powder.

ANOVA with post hoc Tukey test of the results showed that only the undiluted extracts had antiplatelet activity in the test (Figure 1). Standard rosmarinic acid was analyzed in concentrations of 0.09, 0.9, 9 and 90 µM. Only 90 µM rosmarinic acid solution showed statistically significant reduction of platelet aggregation.

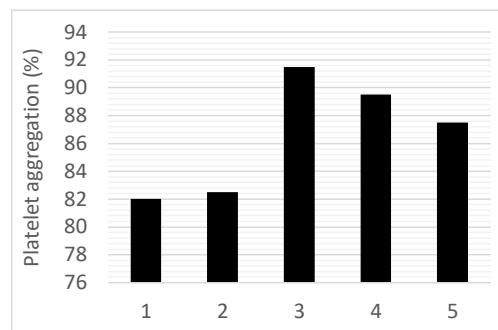


Figure 1: Antiaggregatory effect of *Melissa officinalis* leaf extracts (1-5) (see Table 1 for concentrations). Platelet aggregation is expressed as percentage to the solvent control.

To the best of our knowledge this is the first report of antiplatelet effect of lemon balm leaf extracts. This activity can be attributed mainly to rosmarinic acid as all extracts showed antiplatelet activity at concentrations of 50–120 µM (based on HPLC results, % rosmarinic acid {see Table 1} x concentration of extract (1 mg/mL) / M (rosmarinic acid, 360.3 g/mol}), which is comparable with 90 µM concentration of rosmarinic acid in the antiplatelet test.

Previously, lemon balm essential oil was shown to have an antiplatelet effect at a concentration of 4 µg/mL using collagen and thrombin as inducers of aggregation that was attributed to terpene constituents [5]. The lower results that we have observed can be attributed to the different chemical composition of the leaf extract compared with the essential oil. The advantage of our approach is that the analysis was on whole blood rather than platelet rich plasma used in the previous study [8].

Experimental

Plant material: Some of the leaf samples collected in 2014 and 2015 (samples 1 and 2) were cultivated in the Pharmaceutical Botanical Garden "Fran Kušan", Faculty of Pharmacy and Biochemistry, University of Zagreb, while the others (samples 3-5) were obtained from Suban (Strmec, Croatia). Voucher specimens are deposited in the Herbarium of the Department of Pharmaceutical Botany, University of Zagreb.

Determination of total phenolic acids: The content of phenolic acids in the plant samples was determined by the official Eur. Ph. method [3]. Extracts were prepared by reflux extraction of 0.2 g of the powdered drug with 190 mL of ethanol (50%, v/v) for 30 min. Filtrate and washings were collected and diluted to 200 mL with diluted ethanol. Total phenolic acid content was determined by measuring the absorbance of the complex formed between phenolic acids and sodium nitrite – sodium molybdate at 505 nm [9]. The measurements were carried out using a Varian Cary 50 Bio spectrophotometer (Varian Inc., USA). The content of phenolic acids was expressed as the equivalent of rosmarinic acid based on 3 independent analyses.

HPLC analysis of rosmarinic acid: Rosmarinic acid content was determined by reverse phase high performance liquid chromatography coupled with a diode array detector on an Agilent 1100 system (Agilent, USA) using the method described by Medić-Šarić *et al.* [10]. Extracts used for determination of phenolic acids

were filtered (Whatman 0.22 µm syringe filter, Sigma Aldrich, USA). A Zorbax SB-C18 (250 mm x 4.6 mm, particle size 5 µm) column with precolumn (12.5 mm x 4.6 mm, particle size 5 µm) was used as the stationary phase. Elution was achieved with a gradient of mobile phases A (water, methanol and formic acid in ratio 93:5:2) and B (water, methanol and formic acid in ratio 3:95:2) following the timetable (t/min, %B): (0, 20), (10, 40), (35, 50), (47, 50), (70, 80), (80, 20). Detection was based on retention time and UV spectra compared with standard; quantification was based on calibration curves recorded at 254 nm.

Determination of antiplatelet activity: Platelet aggregation was analyzed in whole blood utilizing a Multiplate® impedance analyzer (Roche, Switzerland). Blood samples were obtained from 9 healthy volunteers with their written consent. This study has been approved by the ethical committee of the Croatian Institute of Transfusion Medicine. The following procedure was used for all analyses: 300 µL of blood was incubated at 37°C for 6 min with 20 µL of lemon balm extract solution and 300 µL of saline with CaCl₂. Aggregation cascade was induced according to protocol by adding 20 µL of agonist ADP reagent (6.3 µM) [3]. Aggregation was measured for 6 min and expressed as area under the curve in arbitrary units (AU). Untreated samples (negative control, 100% of aggregation) were analyzed with 20 µL of solvent. All measurements were performed in triplicate. Final results for extracts were expressed as percentage of residual platelet aggregation compared with the untreated sample. Rosmarinic acid standard was used as positive control. Statistical analysis was performed using the Microsoft Office Excel 2003 (Microsoft Corporation, USA) program; *p* values lower than *a* = 0.05 were regarded as statistically significant.

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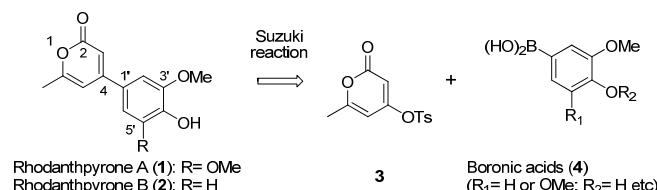
A Concise Synthesis of Rhodanthpyrone A and B, Natural 4-(Hydroxyphenyl)-substituted α -Pyrone

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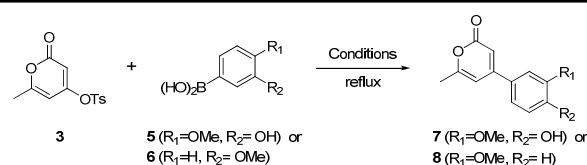
A concise synthesis of rhodanthpyrone A and B was accomplished *via* a Suzuki coupling reaction. To find the conditions appropriate to install hydroxyphenyl moieties to the α -pyrone skeleton, a model study was conducted using commercially available boronic acids. It was revealed that the hydroxy moiety of the phenylboronic acids should be concealed when reacted with labile 4-tosyl α -pyrone. Consequently, rhodanthpyrone A and B could be synthesized in high yields by Suzuki reaction using TIPS-protected arylboronic acids. This procedure provided a concise and versatile route for the synthesis of rhodanthpyrones and their 4-aryl substituted α -pyrone analogs.

Keywords: Synthesis, Rhodanthpyrones, α -Pyrene, Suzuki coupling reaction.

α -Pyrene (=2H-pyran-2-one) is a six-membered cyclic unsaturated ester found in numerous biologically active natural products, which exhibit strong and a broad variety of biological efficacies such as antimalarial, antimicrobial, and anti-cancer [1-3]. The pyrone moiety and its fused derivatives such as coumarin and chromone structures have been recognized as a privileged scaffold [4]. In this regard, these natural α -pyrones and their derivatives have been considered as an attractive target in terms of both synthetic and medicinal chemistry [5]. Recently, rhodanthpyrone A (**1**) and B (**2**), rare 4-aryl substituted α -pyrones, were isolated by Wang *et al.* from *Gentiana rhodantha*, a plant used as a traditional medicine in China for the treatment of inflammation, cholecystitis, and tuberculosis [6-7]. Considering not only the therapeutic properties of *G. rhodantha*, but also the usefulness of the privileged α -pyrone structure in medicinal chemistry, it is believed that these rhodanthpyrones can exhibit therapeutically useful biological activities, and the 4-aryl substituted α -pyrone structure may be a novel scaffold for further medicinal chemistry research.


Figure 1: Structure of rhodanthpyrones and retrosynthetic analysis.

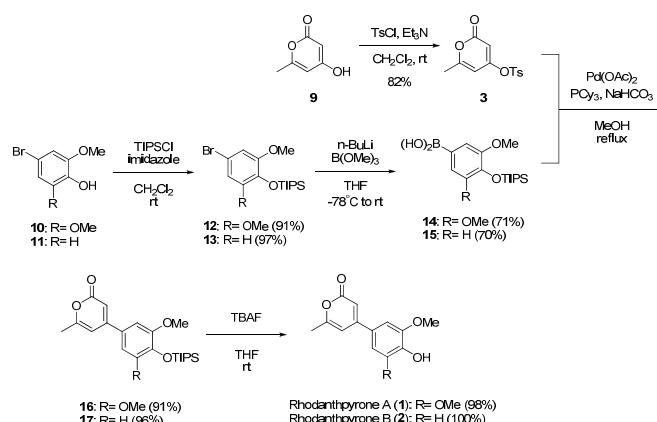
Palladium catalyzed cross-coupling reactions such as the Suzuki reaction have become a general and versatile method in organic synthesis for the introduction of aryl and heteroaryl substituents into diverse molecules. For instance, Rossi and his colleagues reported palladium-catalyzed Suzuki reactions between known 4-tosyloxy-pyrone **3** [8] and several substituted arylboronic acids in high yields using Pd(OAc)₂ with tricyclohexylphosphine (PCy₃) in methanol as an optimized catalyst system [9]. Based on this previous study, it was supposed that a concise synthetic method for rhodanthpyrones could be established by the Suzuki reaction between 4-tosylate (**3**) and suitable 4-hydroxyphenyl boronic acids (**4**), as shown in Figure 1.


Table 1: Model study of Suzuki reaction

Entry	Condition	Result
1	5, Pd(OAc) ₂ (5%), PCy ₃ (10%), NaHCO ₃ , MeOH	
2	5, Pd(OAc) ₂ (5%), PCy ₃ (10%), NaHCO ₃ , toluene/H ₂ O	mixture of
3	5, Pd(OAc) ₂ (5%), PCy ₃ (10%), NaHCO ₃ , dioxane	unidentifiable
4	5, Pd(OAc) ₂ (10%), PCy ₃ (20%), NaHCO ₃ , MeOH	side products
5	5, Pd(OAc) ₂ (5%), PCy ₃ (10%), K ₂ HPO ₄ , MeOH	
6	5, Pd(PPh ₃) ₄ (5%), CuI, NaHCO ₃ , benzene/EtOH	
7	6, Pd(OAc) ₂ (5%), PCy ₃ (10%), NaHCO ₃ , MeOH	8 (95%)

Initially, as shown in Table 1, the Suzuki reactions between 4-tosylate (**3**) and commercially available boronic acids such as regioisomeric 3-hydroxy-4-methoxyphenylboronic acid (**5**) were carried out as model studies. The Suzuki coupling reaction of the phenylboronic acid (**5**) with 4-tosyloxypryanone (**3**) using Pd(OAc)₂ with tricyclohexylphosphine in methanol, reported in entry 1, resulted in only a mixture of unidentifiable side products. Several modifications of the reaction conditions, such as solvent (entry 2 and 3), loading amount of catalyst (entry 4), and base (entry 5), were tried, but 4-aryl α -pyrone (**7**) could not be observed. Moreover, copper-facilitated Suzuki reaction (entry 6), used for the synthesis of 4-arylcoumarin analogues [10], also did not afford the 4-aryl α -pyrone (**7**). On the other hand, the Suzuki coupling reaction using 3-methoxyphenylboronic acid (**6**) (entry 7) afforded 4-aryl α -pyrone (**8**) in a high yield. Based on these observations, it was supposed that the electrophilic phenol of the boronic acid **5**, enough to lead to various undesired reactions [11]. Based on this idea, introduction of a protecting group to the phenol of boronic acid was decided. Among the various protecting groups for phenol, the triisopropylsilyl (TIPS) group was selected as it is known to be a stable and easily removable silyl protecting group in substrates bearing reducible olefin and labile ester.

As shown in Scheme 1, syntheses of rhodanthpyrone A and B commenced with TIPS-protection of bromophenols (**10** [12] and **11**).

**Scheme 1:** Synthesis of rhodanthpyrone A and B

Bromobenzenes (**12** and **13**) were converted to arylboronic acids (**14** and **15**) in moderate yields by treatment with *n*-butyllithium, followed by addition of trimethylborate and hydrolysis using 3N hydrochloric acid. Suzuki reaction using $\text{Pd}(\text{OAc})_2$ with tricyclohexylphosphine in methanol between TIPS protected arylboronic acids (**14** and **15**) and 4-tosylate (**3**), prepared from 4-hydroxypyranone (**9**) [8], readily afforded 4-aryl α -pyrones (**16** and **17**), respectively. Rhodanthpyrone A (**1**) and B (**2**) could be obtained from 4-aryl α -pyrones (**16** and **17**) by deprotection using tetrabutylammonium fluoride (TBAF), without any side product. The spectral data of synthetic rhodanthpyrone A (**1**) and B (**2**) were identical to those of the reported data [6].

In summary, this article describes a concise synthesis of natural rhodanthpyrones via the Suzuki coupling reaction of labile 4-tosyl α -pyrone using silyl protected boronic acids. To find the Suzuki reaction conditions appropriate to install hydroxyphenyl moieties at the 4-position of the α -pyrone, model, a study was conducted using commercially available boronic acids. It was supposed that the hydroxy moiety of phenylboronic acids should be concealed. In this connection, the TIPS group, stable in the Suzuki reaction and easily removable without side-reactions, was introduced to protect the 4-hydroxy moiety of the arylboronic acids. Rhodanthpyrone A and B could be synthesized in high yields by the Suzuki reaction using TIPS-protected arylboronic acids, followed by deprotection. To my best knowledge, this study is the first practical example of the synthesis of 4-aryl substituted α -pyrones bearing a hydroxy moiety. This will be helpful in the synthesis of derivatives of rhodanthpyrone and construction of a chemical library of 4-aryl substituted α -pyrones.

Experimental

General: Unless noted otherwise, all starting materials and reagents were obtained commercially and were used without further purification. All solvents utilized for routine product isolation and chromatography were of reagent grade and glass-distilled, and reaction flasks were dried at 100°C before use. Flash column chromatography was performed using silica gel 60 (230-400 mesh, Merck) with the indicated solvents. Thin-layer chromatography (TLC) was performed using 0.25-mm silica gel plates (Merck). Mass spectra were obtained using a VG Trio-2 GC-MS instrument, and high-resolution mass spectra were obtained using a JEOL JMS-AX 505WA unit. Infrared spectra were recorded on a FT-IR spectrometer. ^1H and ^{13}C spectra were recorded on a Bruker Avance III 700 (700 MHz) spectrometer in either deuteriochloroform (CDCl_3), or deuteriomethanol (CD_3OD). Chemical shifts are expressed in parts per million (ppm, δ) downfield from

tetramethylsilane and are referenced to the deuterated solvent. ^1H NMR data are reported in the order of chemical shift, multiplicity (s, singlet; bs, broad singlet; d, doublet; t, triplet; q, quartet; m, multiplet; and/or multiple resonances), number of protons, and coupling constant in Hertz (Hz).

6-Methyl-2-oxo-2*H*-pyran-4-yl tosylate (3): Tosylated compound **3** was obtained in 82% yield (915 mg) as a white solid from 500 mg (3.97 mmol) of **9** via the reported procedure, and the NMR spectra data were identical with those reported [8].

(4-Bromo-2,6-dimethoxyphenoxy)triisopropylsilane (12): To a solution of 4-bromo-2,6-dimethoxyphenol (**10**) [11] (350 mg, 1.50 mmol) and imidazole (153 mg, 2.25 mmol) in CH_2Cl_2 (20 mL) was added triisopropylsilyl chloride (0.42 mL, 1.95 mmol) at ambient temperature. The reaction mixture was stirred for 48 h at the same temperature and then quenched with H_2O . The reaction mixture was diluted with CH_2Cl_2 . The organic layer was washed with water and brine, dried over MgSO_4 , and concentrated *in vacuo*. Purification of the residue via flash column chromatography on silica gel ($\text{EtOAc}/n\text{-hexane} = 1 : 70$) afforded 532 mg (91%) of **12** as a colorless oil.

IR (thin film, neat): 2943 (m), 2866 (m), 1587 (m), 1504 (s), 1131 (s), 822 (m) cm^{-1} .

^1H NMR (700 MHz, CDCl_3) δ : 6.65 (s, 2H, Ph-H), 3.77 (s, 6H, -OCH₃), 1.26-1.21 (m, 3H, -Si-CH-), 1.08 (d, 18H, $J = 7.7$ Hz, -CH₃).

^{13}C NMR (CDCl_3 , 175 MHz) δ : 152.0 (2C, C), 134.3 (1C, C), 112.3 (1C, C), 108.5 (2C, CH), 55.7 (2C, CH₃), 17.9 (3C, CH), 13.4 (6C, CH₃).

LR-MS: (ESI) m/z : 411 ($\text{M}+\text{Na}^+$).

HR-MS (FAB+): Calcd for $\text{C}_{17}\text{H}_{30}\text{BrO}_3\text{Si}^+$ ($\text{M}+\text{H}^+$): 389.1142. Found: 389.1148

(4-Bromo-2-methoxyphenoxy)triisopropylsilane (13): Silyloxy ether **13** was prepared by the procedure for **12**, using **11** (530 mg, 2.61 mmol) instead of **10**. Purification of the residue via flash column chromatography on silica gel ($\text{EtOAc}/n\text{-hexane} = 1 : 70$) afforded 909 mg (97%) of **13** as a colorless oil.

IR (thin film, neat): 2943 (s), 2867 (s), 1584 (s), 1496 (s), 1227 (s), 835 (s) cm^{-1} .

^1H NMR (700 MHz, CDCl_3) δ : 6.94 (d, 1H, $J = 2.1$ Hz, Ph-H), 6.91 (dd, 1H, $J = 2.1, 8.4$ Hz, Ph-H), 6.73 (d, 1H, $J = 8.4$ Hz, Ph-H), 1.26-1.20 (m, 3H, -Si-CH-), 1.08 (d, 18H, $J = 7.0$ Hz, -CH₃).

^{13}C NMR (CDCl_3 , 175 MHz) δ : 151.9 (1C, C), 145.1 (1C, C), 123.7 (1C, CH), 121.7 (1C, CH), 115.6 (1C, HC), 113.2 (1C, C), 55.8 (1C, CH₃), 18.1 (3C, CH), 13.0 (6C, CH₃).

LR-MS (ESI): m/z 359 ($\text{M}+\text{H}^+$).

HR-MS (FAB+): Calcd for $\text{C}_{16}\text{H}_{28}\text{BrO}_2\text{Si}^+$ ($\text{M}+\text{H}^+$): 359.1036.

3,5-Dimethoxy-4-(triisopropylsilyloxy)phenylboronic acid (14): To a solution of bromobenzene **12** (330 mg, 0.85 mmol) in fresh distilled THF (10 mL) was added slowly 2.5 M *n*-BuLi in *n*-hexane (0.44 mL, 1.10 mmol) at -78°C. After 30 min, to the reaction mixture was added trimethyl borate (0.19 mL, 1.70 mmol) and the mixture was stirred for 1 h before warming to ambient temperature. After 1 h, the reaction mixture was quenched with 3N HCl, and diluted with EtOAc. The organic layer was washed with water and brine, dried over MgSO_4 , and concentrated *in vacuo*. Purification of the residue via flash column chromatography on silica gel ($\text{EtOAc}/n\text{-hexane} = 1 : 2$) afforded 213 mg (71%) of **14** as a white solid.

MP: 185 ~ 187°C.

IR (thin film, neat): 2944 (m), 2867 (m), 1129 (w), 1414 (m), 1343 (s), 1129 (m), 886 (m) cm^{-1} .

¹H NMR (700 MHz, CDCl₃) δ: 7.43 (s, 2H, Ph-H), 3.93 (s, 6H, -OCH₃), 1.34-1.28 (m, 3H, -Si-CH-), 1.12 (d, 18 H, J = 7.0 Hz, -CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 151.3 (2C, C), 140.0 (1C, C), 112.2 (1C, C), 110.1 (1C, CH), 55.6 (2C, CH₃), 17.9 (3C, CH), 13.4 (6 C, CH₃).

LR-MS (ESI): *m/z* 355 (M+H⁺).

HR-MS (FAB+): Calcd for C₁₇H₃₂BO₅Si⁺ (M+H⁺): 355.2107. Found: 355.2113

3-Methoxy-4-(triisopropylsilyloxy)phenylboronic acid (15):

Boronic acid **15** was prepared by the procedure for **14**, using **13** (350 mg, 0.97 mmol) instead of **12**. Purification of the residue via flash column chromatography on silica gel (EtOAc/n-hexane = 1 : 5) afforded 220 mg (70%) of **15** as a white solid.

MP: 150 ~ 151°C.

IR (thin film, neat): 2945 (m), 2868 (m), 1595 (m), 1413 (s), 1371 (m), 1274 (m), 881 (m) cm⁻¹.

¹H NMR (700 MHz, CDCl₃) δ: 7.73 (dd, 1H, J = 1.4, 7.7 Hz, Ph-H), 7.67 (d, 1H, J = 1.4 Hz, Ph-H), 7.00 (d, 1H, J = 7.7 Hz, Ph-H), 3.93 (s, 3H, -OCH₃), 1.32-1.26 (m, 3H, -Si-CH-), 1.11 (d, 18 H, J = 7.7 Hz, CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 150.9 (1C, C), 150.2 (1C, C), 129.7 (1C, CH), 126.8 (1C, C), 118.9 (1C, CH), 117.3 (1C, CH), 55.7 (1C, CH₃), 18.1 (3C, CH), 13.1 (6C, CH₃).

LR-MS (ESI) *m/z*: 347 (M+Na⁺).

HR-MS (FAB+): Calcd for C₁₆H₃₀BO₄Si⁺ (M+H⁺): 325.2001.

General procedure for Suzuki coupling reaction: To a solution of 4-tosyloxypyranone (1 equiv.), NaHCO₃ (3 equiv), and arylboronic acid (1.2 equiv.) in methanol (0.05 M) under an argon atmosphere, were added a solution of tricyclohexylphosphine in toluene (0.60 M, 0.1 equiv.) and Pd(OAc)₂ (0.05 equiv.). The reaction mixture was heated to reflux for 3 h, and then cooled to ambient temperature. The reaction mixture was concentrated *in vacuo*, and diluted with EtOAc and H₂O. The organic layer was washed with water and brine, dried over MgSO₄, and concentrated *in vacuo*. Purification of the residue via flash column chromatography on silica gel (EtOAc/n-hexane = 1 : 5~10) afforded the corresponding 4-aryl substituted α-pyrone.

4-(3-Methoxyphenyl)-6-methyl-2*H*-pyran-2-one (8): 4-Aryl α-pyrone **8** was obtained in 95% yield (37 mg) as a white solid from 50 mg (0.18 mmol) of **3** *via* the general procedure.

MP: 86 ~ 87°C.

IR (thin film, neat): 2349 (w), 1714 (s), 1549 (m), 1290 (w), 1039 (w) cm⁻¹.

¹H NMR (700 MHz, CDCl₃) δ: 7.41 (t, 1H, J = 8.4 Hz, Ph-H), 7.17 (qd, 1H, J = 0.7, 7.7 Hz, Ph-H), 7.09 (t, 1H, J = 2.1 Hz, Ph-H) 7.04 (ddd, 1H, J = 0.7, 1.4, 7.7 Hz, Ph-H), 6.37 (s, 1H, H-5), 6.31 (s, 1H, H-3), 3.88 (s, 3H, -OCH₃), 2.35 (s, 3H, -CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 163.5 (1C, C), 162.1 (1C, C), 160.1 (1C, C), 155.5 (1C, C), 137.3 (1C, CH), 130.2 (1C, C), 119.0 (1C, CH), 116.0 (1C, CH), 112.3 (1C, CH), 108.3 (1C, CH), 103.6 (1C, CH), 55.4 (1C, CH₃), 20.2 (1C, CH₃).

LR-MS (ESI) *m/z*: 217 (M+H⁺).

HR-MS (FAB+): Calcd for C₁₃H₁₃O₃⁺ (M+H⁺): 217.0859. Found: 217.0860

4-(3,5-Dimethoxy-4-(triisopropylsilyloxy)phenyl)-6-methyl-2*H*-pyran-2-one (16): 4-Aryl α-pyrone **16** was obtained in 91% yield (179 mg) as a white solid from 132 mg (0.47 mmol) of **3** and 200 mg of boronic acid **14** (0.56 mmol) *via* the general procedure.

MP: 152 ~ 153°C,

IR (thin film, neat): 2943 (w), 2865 (w), 1700 (s), 1516 (m), 1129 (m) cm⁻¹.

¹H NMR (700 MHz, CDCl₃) δ: 6.75 (s, 2H, Ph-H), 6.32 (s, 1H, H-5), 6.31 (s, 1H, H-3), 3.85 (s, 6H, -OCH₃), 2.32 (s, 3H, CH₃), 1.29-1.24 (m, 3H, -Si-CH-), 1.07 (d, 18 H, J = 7.7 Hz, CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 163.7 (1C, C), 161.7 (1C, C), 155.6 (1C, C), 151.8 (2C, C), 137.8 (1C, C), 127.5 (1C, C), 106.8 (1C, CH), 103.7 (2C, CH), 103.4 (1C, CH), 55.8 (2C, CH₃), 20.2 (1C, CH₃), 17.9 (3C, CH), 13.4 (6C, CH₃).

LR-MS (ESI) *m/z*: 419 (M+H⁺).

HR-MS (FAB+): Calcd for C₂₃H₃₅O₅Si⁺ (M+H⁺): 419.2248.

4-(3-Methoxy-4-(triisopropylsilyloxy)phenyl)-6-methyl-2*H*-pyran-2-one (17):

4-Aryl α-pyrone **17** was obtained in 96% yield (152 mg) as a white solid from 116 mg (0.41 mmol) of **3** and 200 mg of boronic acid **14** (0.62 mmol) *via* the general procedure.

MP: 145 ~ 146°C.

IR (thin film, neat): 2944 (w), 2868 (w), 1767 (s), 1521 (m), 1296 (m), 1034 (m), 774 (m) cm⁻¹.

¹H NMR (700 MHz, CDCl₃) δ: 7.10 (dd, 1H, J = 2.1, 8.4 Hz, Ph-H), 7.07 (d, 1H, J = 2.1 Hz, Ph-H), 6.95 (d, 1H, J = 8.4 Hz, Ph-H), 6.33 (s, 1H, H-5), 6.32 (s, 1H, H-3), 3.89 (s, 3H, -OCH₃), 2.33 (s, 3H, -CH₃), 1.31-1.28 (d, 3H, -Si-CH-), 1.11 (d, 18H, J = 7.7 Hz, -CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 163.8 (1C, C), 161.7 (1C, C), 155.2 (1C, C), 151.3 (1C, C), 148.4 (1C, C), 128.6 (1C, C), 120.7 (1C, CH), 120.0 (1C, CH), 110.2 (1C, CH), 106.6 (1C, CH), 103.3 (1C, CH), 55.6 (1C, CH₃), 20.2 (1C, CH₃), 17.9 (3C, CH), 12.9 (6C, CH₃).

LR-MS (ESI) *m/z*: 411 (M+Na⁺).

HR-MS (FAB+): Calcd for C₂₂H₃₃O₄Si⁺ (M+H⁺): 389.2143.

Rhodanthpyrone A (1): To a solution of silyloxy ether (**16**) (100 mg, 0.24 mmol) in THF (6 mL) was added 1.0 M solution of TBAF in THF (0.36 mL, 0.36 mmol) at ambient temperature. The reaction mixture was stirred for 30 min at the same temperature and then quenched with H₂O. The reaction mixture was diluted with EtOAc. The organic layer was washed with water and brine, dried over MgSO₄, and concentrated *in vacuo*. Purification of the residue via flash column chromatography on silica gel (EtOAC/n-hexane = 1 : 1) afforded 62 mg (98%) of rhodanthpyrone A (**1**) as a yellow solid.

MP: 214 ~ 215°C.

IR (thin film, neat): 3377 (w), 2992 (w), 2947 (w), 2838 (w), 1688 (m), 1633 (m), 1549 (m), 1517 (m), 1454 (m), 1326 (m), 1214 (m), 1119 (m), 847 (m), 718 (m) cm⁻¹.

¹H NMR (700 MHz, CD₃OD) δ: 7.03 (s, 2H, H-6'), 6.69 (d, 1H, J = 1.6 Hz, H-5), 6.41 (d, 2H, J = 1.6 Hz, H-3), 3.94 (s, 6H, -OCH₃), 2.35 (s, 3H, CH₃).

¹³C NMR (CDCl₃, 175 MHz) δ: 166.3 (1C, C), 163.5 (1C, C), 158.1 (1C, C), 149.7 (2C, C), 140.2 (1C, C), 126.9 (1C, C), 106.3 (1C, CH), 105.7 (2C, CH), 104.8 (1C, CH), 57.0 (2C, CH₃), 19.9 (1C, CH₃).

LR-MS (FAB+) *m/z*: 263 (M+H⁺).

HR-MS (FAB+): Calcd for C₁₄H₁₅O₅ (M+H⁺): 263.0914. Found: 263.0914.

Rhodanthpyrone B (2): Rhodanthopyrone B (**2**) was prepared by the procedure for rhodanthopyrone A (**1**) using **17** (30 mg, 0.077 mmol) instead of **16**. Purification of the residue via flash column chromatography on silica gel (EtOAC/n-hexane = 1 : 2) afforded 18 mg (100%) of rhodanthopyrone B (**2**) as a yellowish solid.

MP: 138 ~ 140°C.

IR (thin film, neat): 2926 (w), 1708 (s), 1632 (m), 1594 (m), 1516 (m), 1321 (w), 1286 (m), 1210 (m), 1131 (w), 1033 (w), 819 (w), 769 (w), 627 (w) cm⁻¹.

¹H NMR (700 MHz, CD₃OD) δ: 7.26 (d, 1H, *J* = 2.1 Hz, H-2'), 7.24 (dd, 1H, *J* = 2.1, 8.4 Hz, H-6'), 6.89 (d, 1H, *J* = 8.4 Hz, H-5'), 6.64 (s, 1H, H-5), 6.36 (s, 1H, H-3), 3.93 (s, 3H, -OCH₃), 2.32 (s, 3H, -CH₃).
¹³C NMR (CDCl₃, 175 MHz) δ: 166.4 (1C, C), 163.5 (1C, C), 157.9 (1C, C), 151.2 (1C, C), 149.6 (1C, C), 127.8 (1C, C), 121.9 (1C, CH), 116.8 (1C, CH), 111.2 (1C, CH), 105.9 (1C, CH), 104.6 (1C, CH), 56.6 (1C, CH₃), 19.9 (1C, CH₃).
LR-MS (FAB+) *m/z*: 233 (M+H⁺).
HR-MS (FAB+): Calcd for C₁₃H₁₃O₄ (M+H⁺): 233.0808. Found: 233.0809.

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A New Lignan from the Leaves of *Zanthoxylum armatum*

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A new furofuran lignan, zanthonin (**1**) together with 13 known compounds including seven furofuran lignans (**2-8**), one isobutyl amide (**9**), a furanocoumarin (**10**) and four flavonoids (**11-14**) have been isolated from the leaves of *Zanthoxylum armatum*. The chemical structures of these compounds were elucidated mainly on the basis of NMR (1D and 2D) and MS data. This is the first report on the isolation of methylxanthoxylool (**4**) from *Z. armatum*.

Keywords: *Zanthoxylum armatum*, Rutaceae, Eurofuran lignan, Zanthonin.

Zanthoxylum armatum DC [syn. *Z. alatum* Roxb.] (Rutaceae), is a deciduous aromatic tree distributed in tropical and subtropical regions of India at altitudes of 600-2100 m. It is commonly known as Indian prickly ash, Nepal pepper and toothache tree. The bark, seeds and fruit have been widely used in traditional medicine for the treatment of stomach ache, fever and dyspepsia, and is also used for catching fish (piscicidal). The fruit pericarp is used as a flavoring agent and spice in certain traditional dishes because of its pleasant flavor and delectable pungency. The fruit, bark and leaves exhibit anti-inflammatory, hepatoprotective, anti-fungal and anti-microbial activities [1]. Previous phytochemical studies on *Z. armatum* showed the presence of alkaloids, amides, lignans, flavonoids and terpenoids. Among these, lignans and amides are the main active constituents [1,2]. As a part of our search for bioactive compounds from Western Himalayan medicinal plants [3], herein we report the isolation and characterization of a new furofuran lignan (**1**), along with 13 known compounds from the leaves of *Z. armatum*.

Compound **1** was isolated as colorless oil with optical rotation $[\alpha]^{25}_D = -116$ (*c* 0.12, MeOH) and its molecular formula was deduced to be C₂₄H₂₈O₆ by positive HR-ESI-QTOF-MS, exhibiting a protonated ion at *m/z* 413.1964 [M+H]⁺ (calcd. for C₂₄H₂₉O₆ 413.1951 [M+H]⁺). The IR spectrum indicated the presence of a hydroxyl group (3537 cm⁻¹) and an olefinic double bond (1606 cm⁻¹). The UV spectrum in methanol showed absorption maxima at λ_{max} 211, 231 and 281 nm, suggesting the presence of aromatic and unsaturated moieties. The ¹H NMR data displayed resonances for two oxymethylene groups at δ_H 3.83 (1H, m, H-9'a) and 3.35 (1H, t, *J* = 8.5 Hz, H-9'b) and at δ_H 4.13 (1H, m, H-9a) and 3.88 (1H, m, H-9b), two oxymethine groups at δ_H 4.48 (1H, d, *J* = 6.9 Hz, H-7) and 4.81 (1H, d, *J* = 6.0 Hz, H-7') and two methine groups at δ_H 2.94 (1H, q, *J* = 7.3 Hz, H-8) and 3.29 (1H, m, H-8'), indicating the presence of a furofuran moiety. The ¹H NMR spectrum also showed resonances for six aromatic protons as two ABX systems at δ_H 6.90 (1H, s, H-2), 6.83-6.88 (3H, m, H-5, H-6 and H-5'), and 6.69-6.78 (2H, m, H-2' and H-6'), representing the presence of 1,3,4-trisubstituted benzene rings, one prenyloxy group at δ_H 5.48 (1H, t, *J* = 6.6 Hz, H-2"), 4.55 (2H, d, *J* = 6.6 Hz, H-1"), 1.71 (3H, s, H-4") 1.74 (3H, s, H-5") and one methoxy group at δ_H 3.85 (3H, s). The ¹³C and DEPT NMR spectroscopic data showed twenty-four signals. Except for the resonances for a methoxy group and one prenyloxy group, the remaining eighteen carbon signals were

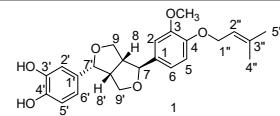


Table 1: ¹³C (150 MHz) and ¹H (600 MHz) NMR data of **1** in CDCl₃, δ in ppm, *J* in Hz.

Position	δ _C	δ _H	Position	δ _C	δ _H
1	133.2	-	4'	143.1	-
2	109.4	6.90 (1H, s)	5'	112.8	6.83-6.88 (1H,m)
3	149.5	-	6'	117.9	6.69-6.78 (1H, m)
4	147.9	-	7'	82.1	4.81 (1H, d, <i>J</i> = 6.0)
5	112.8	6.83-6.88 (1H, m)	8'	49.9	3.29 (1H, m)
6	118.1	6.83-6.88 (1H, m)	9'a	69.6	3.83 (1H, m)
7	87.7	4.48 (1H, d, <i>J</i> = 6.9)	9'b		3.35 (1H, t, <i>J</i> = 8.5)
8	54.1	2.94 (1H, q, <i>J</i> = 7.3)	1"	65.8	4.55 (2H, d, <i>J</i> = 6.6)
9a	71.0	4.13 (1H, m)	2"	119.7	5.48 (1H, t, <i>J</i> = 6.6)
9b		3.88 (1H, m)	3"	137.8	-
1'	130.5	-	4"	18.2	1.71 (3H, s)
2'	115.1	6.69-6.78 (1H, m)	5"	25.8	1.74 (3H, s)
3'	143.7	-	OCH ₃	55.9	3.85 (3H, s)

consistent with a furofuran lignan derivative. The ¹H and ¹³C NMR spectra of **1** were similar to those of planispine-A (**8**) with the major difference being that the methoxy group of **8** was replaced by a hydroxyl group in **1**.

The position of these groups was established from 2D NMR (HMQC, HMBC and NOESY) data. In the HMBC spectra, the correlations of -OCH₃ (δ_H 3.85) to C-3, C-4; H-2 (δ_H 6.90) to C-3, C-4, C-6, and H-1" (δ_H 4.56) to C-4, C-3', C-2" indicated that one methoxy and one prenyloxy group were located at the C-3 and C-4 positions, respectively. The positions of the methoxy and prenyloxy groups were also confirmed by NOESY experiments, which showed the correlation of -OCH₃ (δ_H 3.85) to H-2 (δ_H 6.90) and H-1" (δ_H 4.55) to H-5 (δ_H 6.83). The two hydroxyl groups were connected at C-3' (δ_C 143.7) and C-4' (δ_C 143.1), supported by the HMBC correlations of H-2' to C-1', C-3' and H-6' to C-5', C-4'. The NOESY correlations of H α -7 to H α -9' and H β -8' to H β -7' showed that the proton of H α -7 and H α -9' were on the same face and had α orientations, whereas H β -8' and H β -7' had β orientations. This observation was further supported by the levorotatory optical rotation, which was similar to that of **8** ($[\alpha]^{25}_D = -41$, *c* 0.12, MeOH). Thus, based on the above analysis and literature survey [4] the structure of zanthonin (**1**) was established as (-)-(7 α , 7' β , 8 β)-3-methoxy-4-prenyloxy-7,9': 7',9'-diepoxylinan-3',4'-diol.

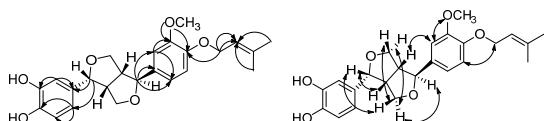


Figure 2: HMBC (→) and NOESY (↔) correlations of compound 1.

Along with the new compound (**1**), 13 known compounds were also isolated and identified as seven furofuran lignans: asarinin (**2**) [5], sesamin (**3**) [6], methylxanthoxylol (**4**) [7], kobusin (**5**) [8], fargesin (**6**) [9], eudesmin (**7**) [10] and planispine-A (**8**) [4]; isobutyl amide: α -sanshoil (**9**) [11], a furanocoumarin: psoralin (**10**) [12] and four flavonoids: catechin (**11**) [13], vitexin (**12**) [14], isovitexin (**13**) [14] and hesperidin (**14**) [15] by comparing their spectroscopic data with those published. Among the known compounds, **4** was isolated for the first time from *Z. armatum*.

Experimental

General: Optical rotation, HORIBA SEPA-300 digital polarimeter; IR, Shimadzu IR Prestige-21 FT-IR spectrophotometer; UV, Shimadzu UV-2600; NMR, Bruker Avance-600; HR-ESI-MS, Waters QTOF Micro mass; TLC, silica gel 60 F254 plates and the compounds were visualized with iodine vapor and 20% H₂SO₄ in ethanol, followed by heating at 110°C. Normal phase (Merck 60-120 and 230-400 mesh) and C18-RP, 23% C (Acros organics, 40-63 micron) silica gel was used for column chromatography.

Plant material: The leaves of *Z. armatum* were collected from Palampur, Himachal Pradesh (alt. 1220 m) in June 2014. A voucher specimen (PLP-16528) was deposited in the Herbarium of CSIR-Institute of Himalayan Bioresource Technology, Palampur, Himachal Pradesh-176061, India.

Extraction and isolation: Air dried and powdered leaves of *Z. armatum* (3.5 kg) were extracted with methanol (3 x 10 L) for 24 h at room temperature by percolation and the percolate was evaporated under reduced pressure at 50°C to give 530 g of methanol extract. This was suspended in water (1000 mL) and subsequently partitioned with *n*-hexane, chloroform, EtOAc and *n*-butanol to obtain *n*-hexane (116.7 g), chloroform (34.2 g), EtOAc (30.7 g) and *n*-butanol (88.5 g) fractions. The water insoluble part

was crystallized in methanol to yield **12** (8.617 g). The CHCl₃ fraction (34.2 g) was chromatographed over silica-gel (60-120 mesh) and eluted using *n*-hexane: EtOAc mixtures (10:0 to 0:10). A total of 46 fractions (250 mL each) were collected. Fractions 13-22 were subjected to silica gel column chromatography (CC) with a gradient of increasing EtOAc in *n*-hexane (8.5:1.5-8:2) to afford **2** (0.984 g) and **3** (0.735 g), which were further purified by crystallization. Fractions 26-38 were subjected to silica gel CC using *n*-hexane: EtOAc (8:2) to furnish **10** (0.047 g), **9** (0.423 g), **6** (0.981 g) and **5** (0.826 g). Fraction 40-42 was subjected to RP C18 CC using MeOH: H₂O (8:2) as eluent to yield **7** (0.220 g). The EtOAc fraction (30.7 g) was separated by CC over silica gel by eluting with a gradient of *n*-hexane: EtOAc (10: 0 to 0: 10) and EtOAc: MeOH (9.5:0.5 to 0:10) yielding 33 fractions (250 mL each). Fractions 18-21 were subjected to CC with EtOAc: *n*-hexane (1:3) resulting in the isolation of **11** (2.141 g). Similarly fractions 24-28 afforded **13** (2.537 g) using EtOAc: MeOH (9.5: 0.5) as eluent. Fraction 12 (7.0: 3.0) was subjected to RP C18 CC using MeOH: H₂O (8:2) to afford **4** (0.030 g). Fractions 14-16 were chromatographed over silica gel CC (230-400 mesh) using *n*-hexane: EtOAc (1:1) yielding 5 fractions (B-1 to B-5). B-3 was subjected to RP C18 CC using MeOH: H₂O (8:2) to yield **8** (0.070 g). Likewise, B-5 yielded **1** (0.030) using MeOH: H₂O (8:2) as eluent. The *n*-butanol fraction (88.5 g) was chromatographed over HP-20 diaion resin and eluted with H₂O: MeOH (10:0 to 0:10) to give 6 fractions. Fraction 3 was subjected to silica gel CC using CHCl₃: MeOH: H₂O (7.8: 2.0: 0.2) to give **14** (0.128 g).

Zanthonin (**1**)

Colorless oil

[α]_D²⁵: -116 (c 0.12, MeOH)

IR (neat): 3537, 1514, 1377, 1259, 1031 cm⁻¹.

UV/Vis λ_{max} (MeOH) nm (log ε): 211 (3.344), 231 (2.820), 281 (1.136).

¹H and ¹³C NMR (CDCl₃): Table 1.

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Production of Podophyllotoxin by Plant Tissue Cultures of *Juniperus virginiana*

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Plant tissue cultures are a potential source of secondary metabolites. However, their production, when compared with intact plants, is usually lower. Phenylalanine, a biogenetic precursor of podophyllotoxin, was used to stimulate podophyllotoxin production in callus and suspension cultures of *Juniperus virginiana* L. The best phenylalanine effect on podophyllotoxin production was manifested in three-years-old callus cultures after a 21-days application of a 10 mmol/L concentration. A podophyllotoxin content of 0.15 mg/g DW was determined, which was about 400% higher in comparison with the control. The maximum content (0.48 mg/g DW) in newly derived suspension cultures (the 4th passage) was induced by 14-days application of a 1 mmol/L concentration; this was about 243% higher than the control. In one-year-old suspension cultures the highest podophyllotoxin content (0.56 mg/g DW) was recorded also after 14-days application of a 1 mmol/L concentration; this was about 211% higher than in the control cultures.

Keywords: *Juniperus virginiana*, Callus cultures, Suspension cultures, Podophyllotoxin production.

A natural lignan, podophyllotoxin, is the preferred precursor for the semi-synthesis of the anticancer drugs etoposide, teniposide and etopophos [1-5]. The traditional plant sources of podophyllotoxin are *Podophyllum* species, but it has also been detected in other plant genera, such as *Linum*, *Juniperus*, *Hypitis*, *Teucrium*, *Nepeta*, *Dysosma*, *Jeffersonia*, *Thymus* and *Thuja*, yet the availability of this lignan remains limited [6,7]. Chemical synthesis of podophyllotoxin is also an expensive process, and so the biotechnological production of this compound using plant cell and tissue cultures can be considered as a promising alternative procedure.

Callus cultures of *Juniperus virginiana* L. (varieties 'Hetzii', 'Glauca', 'Grey Owl') were derived from leaves on Schenk and Hildebrandt medium supplemented with α-naphthaleneacetic acid, kinetin and ascorbic acid. *J. virginiana* suspension cultures (varieties 'Hetzii', 'Glauca', 'Grey Owl') were derived from two-years-old callus cultures on the same medium and under the same conditions as callus cultures. The callus and suspension cultures of the variety 'Glauca' had the best viability and thus provided the most biomass [8].

Plant tissue cultures are a potential source of secondary metabolites, but their production, when compared with intact plants, is usually lower and this is the reason why we are looking for different methods and approaches to increase the production. Phenylalanine, a podophyllotoxin biogenetic precursor [5], was used to stimulate podophyllotoxin production in callus and suspension cultures of *J. virginiana* (variety 'Glauca').

The results proved that podophyllotoxin production in three-years-old callus cultures of *J. virginiana* is low (Figure 1). The best effect of phenylalanine on the production was manifested after a 21-days application of a 10 mmol/L concentration; a maximum podophyllotoxin content of 0.15 mg/g DW was determined, which is about 400% higher than that of the control.

Production of podophyllotoxin in suspension cultures of *J. virginiana* Glauca was larger. The maximum content (0.48 mg/g

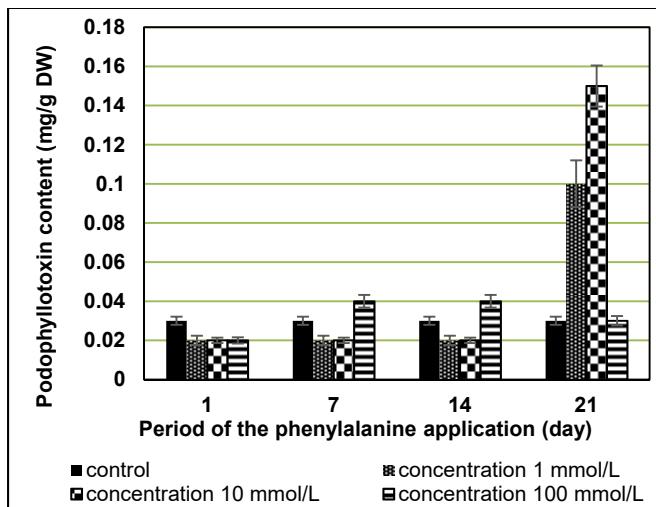


Figure 1: Production of podophyllotoxin in three-years-old callus cultures. Data represent the average values from 3 separate experiments ± SD.

DW) in newly derived suspension cultures (4th passage) was induced by 14-days application of phenylalanine (concentration 1 mmol/L). Production was increased by about 243% in comparison with the control (Figure 2). Increased production (about 213%) was also recorded after 1-day application of the strongest (100 mmol/L) concentration (0.47 mg/g DW).

In the one-year-old suspension cultures of *J. virginiana* (Figure 3), the best effect of phenylalanine was recorded also at the concentration of 1 mmol/L. The highest podophyllotoxin content (0.56 mg/g DW) was also recorded after 14-days application. This production was about 211% higher than that in the control. A positive phenylalanine effect was detected after the shortest, 1-day, application of the strongest concentration (100 mmol/L); podophyllotoxin production (0.55 mg/g DW) was increased by about 150%.

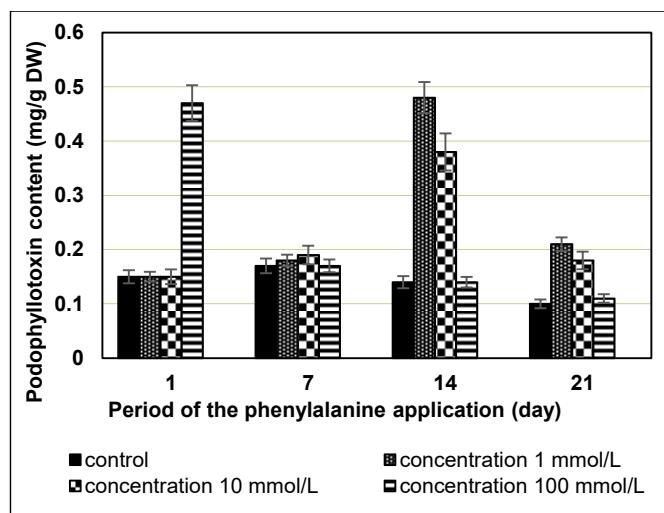


Figure 2: Production of podophyllotoxin in newly derived suspension cultures (the 4th passage). Data represent the average values from 3 separate experiments \pm SD.

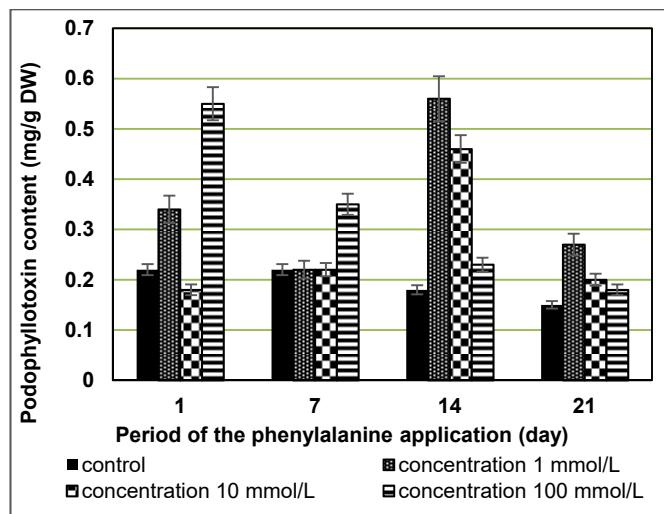


Figure 3: Production of podophyllotoxin in one-year-old suspension cultures. Data represent the average values from 3 separate experiments \pm SD.

Comparison of podophyllotoxin production in the phenylalanine-treated *J. virginiana* suspension cultures (the 4th passage and one-year-old suspension cultures) confirms that the maximum increase in both cases was induced during the 14-days application of the 1 mmol/L concentration. Longer application of this concentration did not increase these values. In contrast, the results show that in three-years-old callus cultures of *J. virginiana* the podophyllotoxin content was the best stimulated after 21-days phenylalanine application (concentration 10 mmol/L). The higher production in the suspension cultures can be explained by the direct contact of cells with nutrient medium, which guarantees easier access of nutrients and the exchange of respiratory gases.

Stimulated podophyllotoxin production is documented also in the callus cultures of *J. chinensis* [9]. Production in the control cultures is low (0.05 mg/g DW), with the highest content being recorded (nearly 0.16 mg/g DW) 15 days after the addition of phenylalanine (concentration 0.5 mg/mL), similarly as in the callus cultures of *J. virginiana*.

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The accumulation of lignans in plant tissue and organ cultures has been discussed based on the work of different authors [4,10-12]. It has been shown that production of podophyllotoxin and related lignans in cell cultures is possible. The newly derived plant tissue cultures as a source of anticancer lignans can play an important role in this respect.

Experimental

Plant material and growth conditions: The three-years-old callus cultures of *J. virginiana* ('varieta 'Glaucia'), the newly derived suspension cultures in the 4th passage and the one-year-old suspension cultures of *J. virginiana* ('varieta 'Glaucia') were cultivated in SH medium [6], supplemented with 3.0 mg/L of NAA, 0.2 mg/L of kinetin and 15 mg/L of ascorbic acid at 25°C and with a 16-h light/8-h dark photoperiod. A sub-cultivation interval of 28 days lasted for callus cultures and 21 days for suspension cultures.

Effect of a biogenetic precursor on production of podophyllotoxin: On the 7th day of cultivation, the callus and suspension cultures were treated with phenylalanine. One mL of an aqueous phenylalanine solution (concentrations of 1 mmol/L, 10 mmol/L and 100 mmol/L) was added to the SH medium (30 mL) and the cultures were cultivated for 1, 7, 14 and 21 days. The control cultures received 1.0 mL of distilled water. The cells were separated from the media by vacuuming, rinsed in distilled water and dried at room temperature.

Determination of podophyllotoxin: Preparation of the samples: 500 mg of control and treated cultures were extracted with 10.0 mL of methanol (100%) using ultrasound for 1 h at 30°C. The supernatant after centrifugation (at 4500 rpm for 5 min) was collected in vials and analyzed by HPLC-DAD. A JASCO 2000-Plus series liquid chromatograph (Jasco, Tokyo, Japan) equipped with diode array was used.

The HPLC conditions were as follows: the main wavelength was set at 280 nm and reference wavelengths for better peak identification were 270 and 290 nm. An RP-18 Lichrospher column (250 x 4 mm, particles size 5 µm) with a precolumn made of the same material was used. The injected volume of the samples and standard solutions was 20 µL and the column was operated at 25 °C.

The mobile phase consisted of (A) methanol/water/phosphoric acid (60:39.7:0.3; v/v/v) and (B) methanol/phosphoric acid (99.7:0.3; v/v). Gradient elution was designed as follows: 0 – 8 min, 0 – 80%; 8 – 9 min, 80 – 100%; 9 – 10 min, 100%; 10 – 11 min, 100% - 0%; 11 – 13 min, 0% of solvent B. The flow rate of the mobile phase was 1 mL/min.

For the podophyllotoxin quantification, a six-point calibration curve was obtained with a pure standard.

All the listed results are averages calculated from the results of 3 independent determination processes.

Acknowledgment - The study was supported by the research programs PRVOUK P40 and SVV 260294.

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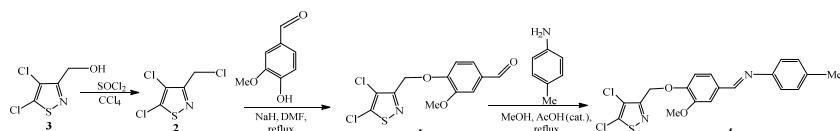
New Data on Vanillin-Based Isothiazolic Insecticide Synergists

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By alkylation of vanillin with 4,5-dichloro-3-chloromethylisothiazole the corresponding ether was synthesized. The latter was then reacted with *p*-toluidine to afford the corresponding azomethine. During the bioassays of synthesized isothiazolic derivatives of vanillin in mixtures with insecticides (imidacloprid and α -cypermethrin) a strong synergistic effect was observed.



Keywords: Vanillin, Isothiazole, Ether, Azomethine, Synergistic effect, Insecticide.

The development of new effective insecticides is one of the key tasks of agrochemistry. The constantly developing resistance of insects to the known insecticides and the aim to reduce the harmful impact on the environment are the main driving forces of this process [1,2]. Among insects with constantly growing resistance to insecticides, the Colorado potato beetle (CPB) is one of the highly notable examples as a serious threat to potato production. Since the middle of the last century, the beetle has developed resistance to 52 different compounds belonging to all major insecticide classes [3]. There are promising results on the creation of transgenic potato lines to control the damage and spread of CPB [4]. However, the development of new substances for chemical protection of potato does not lose its relevance.

Vanillin, a natural aldehydophenol, is widely used in the food industry, medicine and cosmetics. Previously we had reported a series of azomethins based on the vanillin ester of 4,5-dichloroisothiazolic acid, which showed high synergistic effects against CPB in the presence of the neonicotinoid insecticide Kerber® (imidacloprid) [5], whereas the vanillin ester of 4,5-dichloroisothiazolic acid itself showed a potentiating effect with the pyrethroid insecticide Vitan® (α -cypermethrin) [6]. During subsequent studies we found that reduction of the corresponding vanillin azomethins iminic fragment did not lead to significant change in their synergistic activity whereas substitution of chlorine in the 5th position of the isothiazolic ring for bulky fragments, except for the benzylthio fragment, results in a decrease of the isothiazolic vanillin derivatives potentiating activity [7]. Related compounds based on *iso*-vanillin also did not show a noticeable synergistic effect when mixed with the insecticides. For further study of structure activity relationships of isothiazolic vanillin conjugates we decided to synthesize isosters of initially tested compounds by changing the linker between the isothiazole and vanillin fragments from ester to ether. Ether 1 was synthesized by reaction of vanillin with 4,5-dichloro-3-chloromethylisothiazole (2)

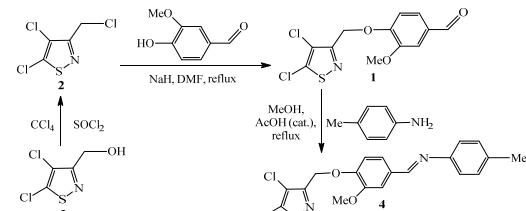


Figure 1: Preparation of vanillin and isothiazole conjugates.

Table 1: Insecticidal activity of insecticides Vitan®, Kerber®, and their mixtures with new vanillin and isothiazole conjugates against potato beetle larvae.

Entry	Active ingredient	a.i. concentration, mg/ha*	Average number of larvae after 24 h, % control	Average number of larvae after 72 h, % control
1	1	200	98.0	98.0
2	4	200	99.0	98.0
3	Vitan	160	83.3	36.7
4	Vitan	80	90.0	76.7
5	Kerber	200	43.3	33.3
6	Kerber	100	86.7	73.3
7	Vitan + 1	80	60.0	60.0
8	Vitan + 4	80	83.3	76.7
9	Kerber + 1	100	46.7	46.7
10	Kerber + 4	100	3.3	0.0

* a.i. means active ingredient, in entry 3–10 – concentration of basic insecticides.

in the presence of sodium hydride in refluxing DMF for 5 h. Chloromethyl derivative 2 was obtained by treatment of previously synthesized (4,5-dichloroisothiazol-3-yl)methanol (3) [8] with thionyl chloride. Condensation of ether 1 with *p*-toluidine in boiling methanol for 4 h in the presence of a catalytic amount of AcOH led to azomethine 4, which was precipitated from the solution by cooling to 0°C. Product 4 was filtered, washed with cold methanol, dried in vacuum over P₂O₅ and further used without purification. The synthesized compounds were identified on the base of IR and ¹H- and ¹³C-NMR spectra data. Azomethine 4 relates to aldimines, for which the *E*-form is preferable, as was established in the results of spectral studies of this group of compounds [9].

The synergistic activity of the synthesized isosters **1** and **4** in mixtures with Vitan® and Kerber® was evaluated by the greenhouse bioassay. Potato beetle larvae were treated with insecticide containing 5%, w/w, of either ether **1** or azomethine **4** with respect to the insecticide. The results of the bioassays are shown in Table 1. Individual compounds **1** and **4** had no insecticidal activity. However, their addition to the insecticides led to promotion of the insecticidal activity of both Vitan® and Kerber®. Vanillin ether **1** showed higher potentiating activity with Vitan® (α -cypermethrin), whereas azomethine **4** showed higher activity in composition with Kerber® (imidacloprid). The revealed potentiating activity of the tested compounds is similar to the activity of the initially tested vanillin ester of 4,5-dichlorothiazolic acid and *p*-toluidine azomethine on its base. The obtained results, as well as our previous ones, indicate the crucial significance of the vanillin-isothiazolic scaffold for demonstration of biological activity in combination with α -cypermethrin and imidacloprid, though the linker nature between isothiazole and vanillin seems to be irrelevant. We assume this information will be useful for further research in this domain.

Experimental

General: Melting points, Boetius heating table; IR, Nicolet Protégé spectrometer; NMR, Brucker Avance-500 spectrometer.

4,5-Dichloro-3-(chloromethyl)isothiazole (2): A solution of 1.84 g (0.01 mol) of (4,5-dichloroisothiazol-3-yl)methanol (**3**) and 2.4 g (0.02 mol) of thionyl chloride in 15 mL dry tetrachloromethane was stirred for 10 h at room temperature, and then refluxed for 8 h. The reaction mixture was cooled, solvent and excess thionyl chloride were removed under vacuum, and 30 mL of *n*-hexane and 0.56 g (2 mmol) of phosphorous pentoxide were added to the crude product. The obtained suspension was stirred at room temperature for 1 h and filtered through a thin layer of silica gel. *n*-Hexane was evaporated under reduced pressure to give the product as a yellow oil. Yield 1.34 g (66%).

IR (KBr): 3022, 2964, 2851, 1507, 1435, 1391, 1375, 1310, 1262, 1213, 1154, 1100, 980, 925, 830, 807, 751, 675, 596, 523 cm⁻¹.

¹H NMR (500 MHz, CDCl₃): 4.63 (2H, s, CH₂).

¹³C NMR (126 MHz, CDCl₃): 39.5 (CH₂), 123.2 (C), 149.1(C), 161.9 (C).

Anal. Calcd for C₄H₂Cl₃NS: C, 23.73; H, 1.00; Cl, 52.52; N 6.92; S 15.83. Found C, 23.82; H 1.18; Cl 52.60; N 6.99; S 15.97.

4-{(4,5-Dichloroisothiazol-3-yl)methoxy}-3-methoxybenzaldehyde (1**):** 0.3 g (1.85 mmol) of vanillin and 0.04 g (1.81 mmol) of sodium hydride (as 60% suspension in mineral oil) was stirred in 10 mL of DMF for 5 min. After that, 0.25 g (1.23 mmol) of 4,5-dichloro-3-(chloromethyl)isothiazole **2** was added, and the reaction mixture was refluxed for 5 h. The reaction mixture was cooled down, and quenched with 50 mL of sodium bicarbonate saturated solution, and stirred for 20 min. The precipitate was filtered off, dried under vacuum and purified by column chromatography on silica gel using benzene as the eluting solvent. Yield 0.28 g (71%). MP 127-128°C.

IR (KBr): 3080, 3007, 2961, 2931, 2852, 2831, 1682, 1588, 1508, 1461, 1422, 1395, 1377, 1350, 1280, 1264, 1232, 1140, 1105, 1033, 1013, 1004, 859, 817, 734, 654 cm⁻¹.

¹H NMR (500 MHz, CDCl₃): 3.91 (3H, s, OCH₃), 5.27 (2H, s, CH₂), 7.09 (1H, d, J=8.7, CH_{arom}), 7.39-7.42 (2H, m, CH_{arom}), 9.84 (1H, s, HC=O).

¹³C NMR (126 MHz, CDCl₃): 56.2 (OCH₃), 66.3 (CH₂), 109.9 (CH_{arom}), 113.0 (CH_{arom}), 123.2 (C), 126.3 (CH_{arom}), 131.2 (C), 149.0 (C), 150.3 (C), 152.9 (C), 161.2 (C), 191.0 (CH=O).

Anal. Calcd for C₁₂H₉Cl₂NO₃S: C, 45.30; H, 2.85; Cl, 22.28; N 4.40; S 10.08. Found C, 45.41; H 2.99; Cl 22.35; N 4.43; S 10.14.

(E)-1-((4,5-Dichloroisothiazol-3-yl)methoxy)-3-methoxyphenyl-N-(*p*-tolyl)methanimine (4**):** To the solution of 0.32 g (1 mmol) of 4-((4,5-dichloroisothiazol-3-yl)methoxy)-3-methoxybenzaldehyde (**1**) and 0.11 g (1 mmol) of *p*-toluidine in 5 mL dry methanol, 1 drop of acetic acid was added and the mixture was refluxed for 4 h. Then the reaction mixture was cooled down to 0°C, the precipitate was filtered, washed with 1 mL of cooled methanol, and dried under vacuum over P₂O₅. Yield 0.30 g (72%).

MP 122-123°C.

IR (KBr): 3080, 3038, 3029, 3008, 1625, 1595, 1581, 1512, 1460, 1417, 1377, 1275, 1232, 1215, 1138, 1107, 1025, 1011, 808 cm⁻¹.

¹H NMR (500 MHz, CDCl₃): 2.41 (3H, s, Me), 4.01 (3H, s, OMe), 5.30 (2H, s, CH₂), 7.08 (1H, d, J=8.2, CH_{arom}), 7.16 (2H, d, J=8.3, 2CH_{arom}), 7.23 (2H, d, J=8.1, 2CH_{arom}), 7.28-7.33 (1H, m, CH_{arom}), 7.66 (1H, d, J=1.8, CH_{arom}), 8.40 (1H, s, CH=O).

¹³C NMR (126 MHz, CDCl₃): 21.1 (CH₃), 56.3 (OCH₃), 66.7 (CH₂), 110.0 (CH_{arom}), 114.0 (CH_{arom}), 120.9 (2CH_{arom}), 123.3 (C), 123.9 (CH_{arom}), 129.9 (2CH_{arom}), 131.2 (C), 135.7 (C), 148.8 (C), 149.7 (C), 150.5 (C), 158.0 (C), 159.0 (C), 161.9 (CH=O).

Anal. Calcd for C₁₉H₁₆Cl₂N₂O₂S: C, 56.03; H, 3.96; Cl, 17.41; N 6.88; S 7.87. Found C, 56.15; H 4.07; Cl 17.39; N 6.78; S 7.92.

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Phileucin - A Cyclic Dipeptide Similar to Phevalin (Aureusimine B) from *Streptomyces coelicolor* M1146

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Overexpression of a putative type III polyketide synthase (PKSIII) from the marine myxobacterium *Enhygromyxa salina* SWB007 in *Streptomyces coelicolor* M1146 led to the accumulation of a novel monoketopiperazine consisting of phenylalanine and isoleucine. This compound was named phileucin and shows high structural similarity to phevalin (aureusimine B). The protease inhibiting activity was tested against human cathepsin L, human leukocyte elastase, bovine trypsin and bovine chymotrypsin. In contrast to phevalin, no protease inhibition was observed.

Keywords: Myxobacteria, Streptomyces, Monoketopiperazine.

Marine myxobacteria, e.g. of the genus *Enhygromyxa*, have the potential to biosynthesize so far unknown natural products [1]. For *E. salina* SWB007 production of enhygrolides and of salimabromide (**1**) was shown (Figure 1) [2, 3]. Compound **1** is a brominated polyketide with an unprecedented carbon skeleton, which exhibited moderate antibiotic activity against Gram-positive bacteria. However, the biosynthesis of salimabromide (**1**) remained elusive up to now. The structure of **1** gives hints towards the involvement of a polyketide synthase (PKS), a terpene synthase, and a halogenase. Thus, a putative type III PKS identified within the genome of *E. salina* SWB007, which is clustered with a putative halogenase, should be tested towards its involvement in the biosynthesis. For this purpose, the corresponding gene named *ks007* was heterologously expressed in *Streptomyces coelicolor* M1146 [4]. Using the vector pGM1202 (kindly provided by Dr Günther Muth, University of Tübingen, Germany), the expression of *ks007* was inducible under the control of the *tipA* promoter. The identity of the construct pGM1202-*ks007* was verified by sequencing, and it was transferred to *S. coelicolor* M1146 by bi-parental conjugation, yielding strain *S. coelicolor* M1146 pGM1202-*ks007*. Cells of this strain were grown in ISP2 medium and the expression of the gene *ks007* was induced by adding thiomstreptone to the culture. After seven days of incubation, the cultures were extracted using ethyl acetate. Analysis of the resulting crude extract by chromatographic methods revealed one additional peak in comparison to the host strain. To elucidate the identity of the metabolite causing the additional peak, the compound was purified by vacuum liquid chromatography (VLC) and high performance liquid chromatography (HPLC).

Following purification, the molecular formula of compound **2** (Figure 2) was deduced by high resolution mass measurement (HRESIMS m/z 265.1311 [$M+Na^+$]) to be $C_{15}H_{18}N_2O$ (calculated m/z 265.1316 [$M+Na^+$]). The ^{13}C NMR spectrum (Table 1) disclosed 15 resonances resulting from two methyl groups (δ_c 12.3 for C-1, δ_c 18.3 for C-4), two sp^3 methylene groups (δ_c 28.7 for C-2, δ_c 37.1 for C-9), six sp^2 methine groups (δ_c 123.5 for C-8, δ_c 129.9 for C-11, δ_c 129.9 for C-12, δ_c 128.2 for C-13, δ_c 129.9 for C-14, δ_c 129.9 for C-15), one sp^3 methine group (δ_c 37.6 for C-3) and four sp^2 quaternary carbon atoms (δ_c 161.5 for C-5, δ_c 158.5 for C-6,

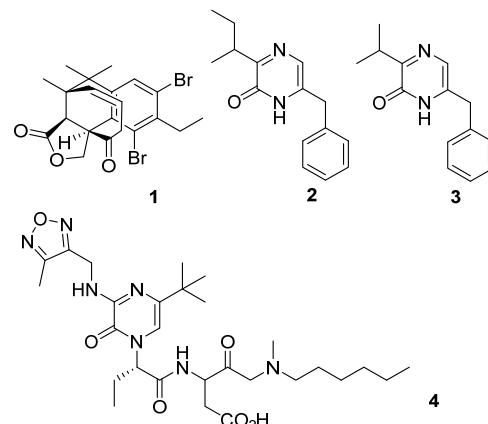
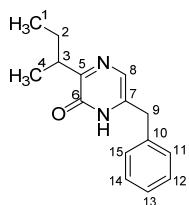


Figure 1: Structures of salimabromide (**1**), phileucin (**2**), phevalin (**3**) and M826 (**4**).

δ_c 139.9 for C-7, δ_c 138.1 for C-10). The 1H NMR spectrum (Table 1) exposed two signals for methyl groups (δ_H 0.90 for H₃-1, δ_H 1.20 for H₃-4), one for a methine group (δ_H 3.22 for H-3) and six protons resonating in the olefinic region (δ_H 7.17 for H-8, δ_H 7.37 for H-11, δ_H 7.31 for H-12, δ_H 7.30 for H-13, δ_H 7.31 for H-14, δ_H 7.37 for H-15). H-8 is incorporated into the double bond $\Delta^{7,8}$ and H-11, H-12, H-13, H-14 and H-15 are incorporated into an aromatic ring system. The aromatic ring system was indicated by the UV spectrum, revealing an absorption band with a maximum at 328 nm (Figure S1).

The planar structure of compound **2** as shown in Figure 1 was deduced from 1H - 1H COSY and 1H - ^{13}C HMBC measurements. The COSY spectrum showed two independent spin systems: (i) one aromatic part from H-11 to H-15, and (ii) one smaller alkyl part from H₃-1 to H₃-4. The HMBC spectrum further reveals the coupling between H-3 and both C-5 and C-6, indicating a branched alkyl moiety. HMBC correlations between H₂-9 and all C-11, C-15, C-7 and C-8 delineated the presence of a phenylalanine derived moiety. However, the connection between both moieties could not be easily observed. A closer look at the HMBC spectrum revealed a very weak coupling between H-8 and C-5, indicating their connection via a nitrogen atom, evidenced by the chemical shift of

**Figure 2:** Structure of phileucin.**Table 1:** ¹H and ¹³C NMR spectral data of compound 2 in chloroform-d₁.

No.	¹³ C [ppm]	mult.	¹ H [ppm]/J	¹ H- ¹ H COSY	¹ H- ¹³ C HMBC
1	12.3	CH ₃	0.9 (t 7.5) a: 1.81 (m)	2a;2b	2;3
2	28.7	CH ₂	b: 1.55 (m)	1;3	1;3;4;5
3	37.6	CH	3.22 (m)		1;2;4;5;6
4	18.3	CH ₃	1.2 (d 7.0)		2;3;5
5	161.5	C			
6	158.5	C			
7	139.9	C			
8	123.5	CH	7.17 (s)		5 (w)
9	37.1	CH ₂	3.88 (s)	8;11;15	7;8;11;15
10	138.1	C			
11	129.9	CH	7.37 (m)	12	13
12	129.9	CH	7.31 (m)	11;13	10
13	128.2	CH	7.30 (m)	12;14	11;15
14	129.9	CH	7.31 (m)	13;15	10
15	129.9	CH	7.37 (m)	14	13

carbon C-5 (Table 1). Taking the molecular formula and the chemical shifts of C-6 and C-7 into account, their connection via a peptide bond was deduced, completing the planar structure of compound 2. Due to the low amount of isolated substance, it was not possible to identify the stereochemistry of phileucin at C-3.

Unexpectedly, the isolated compound was not a polyketide, and thus not the product of the heterologously expressed PKSIII. Instead, structural analysis revealed a 3,6-disubstituted 2(1H)-pyrazinone, a compound which can be considered as dehydrated diketopiperazine, deduced from the amino acids phenylalanine and isoleucine. We suggest the name phileucin (2). The mechanism underlying the production of 2, due to the expression of the putative PKSIII gene from *E. salina* SWB007, is unknown. Compound 2 bears strong resemblance to phevalin (3, also known as aureusimine B), which was reported to have calpain inhibiting properties (IC_{50} 1.3 μ M) [5]. However, this finding was not confirmed with a synthetic sample of phevalin [6]. Phevalin and the structurally related tyrvalin were found in *Staphylococcus aureus* strains and are suspected to be involved in biofilm formation [7].

Moreover, the structurally related 2(1H)-pyrazinone derivative M926 (4) (Figure 1) potently inhibited caspase-3 [8]. In the light of the reported protease inhibiting properties of 3 and 4, compound 2 was evaluated for inhibiting activities towards clinically relevant proteases. The selected enzymes were human cathepsin L, a cysteine protease, as well as the serine proteases human leukocyte elastase, bovine trypsin and bovine chymotrypsin. However, phileucin (2) showed no significant inhibitory activity towards any of the enzymes tested; all IC_{50} values were above 100 μ M.

Experimental

Construction of strain *S. coelicolor* M1146 pGM1202-KS007: The putative ketosynthase gene *ks007* was amplified by colony-PCR, using *E. salina* SWB007 cells with the following primers: forward-primer containing an *Nde*I restriction-site (KS007-for: GCGCATATGTCAACCGTCATCCCCGTTGGTATT), and reverse-

primer containing a *Hind*III restriction-site (KS007-rev: GGCAAGCTTCAGCGGCTCGAAAGAAAAGTAG). PCR reactions were analysed by agarose gel electrophoresis on a 1% agarose gel. The expected band with a size of 1139 bp was excised and the DNA was purified using the Promega SV Gel and PCR clean up system, and ligated into the pGEM-T vector (Promega). The construct was transferred into chemically competent *E. coli* XL1 Blue cells by heat-shock transformation. Candidate colonies were grown in liquid LB_{amp} medium for 16 h at 37°C and plasmids were isolated using Promega Pure yield Miniprep Kit. Restriction of the plasmids with *Nde*I and *Hind*III verified the presence of the target gene *ks007*. A plasmid harbouring the *ks007* was chosen and the identity of the insert was verified by sequencing with T7 and SP6 primers (GATC Biotech). Subsequently, the *ks007* gene was transferred into pGM1202 (*Nde*I, *Hind*III) by ligation for 16 h at 4°C. The resulting plasmid pGM1202-*ks007* was transferred into competent *E. coli* ET12567 pUZ8002 cells using heat-shock transformation, resulting in the strain *E. coli* ET12567 pUZ8002 pGM1202-*ks007*.

The pGM1202-*ks007* plasmid was transferred to *S. coelicolor* M1146 by conjugation between *E. coli* ET12567 pUZ8002 pGM1202-*ks007* and *S. coelicolor* M1146. For the conjugation, ET12567 pUZ8002 pGM1202-*ks007* cells were grown in 100 mL LB_{apra/kan}-medium at 37°C up to an OD₆₀₀ of 0.5. The cells were then precipitated at 10000 rcf for 10 min and the resulting cell pellet washed twice in 10 mL LB-medium. The pellet was resuspended in 1 mL LB-medium. Meanwhile, 10 μ L of *S. coelicolor* M1146 spore solution was added to 500 μ L 2xYT broth (16 g/L tryptone, 10 g/L yeast extract, 5 g/L NaCl, pH 7.0) and heat shocked at 50°C for 10 min. After cooling to room temperature, 500 μ L of *E. coli* ET12567 pUZ8002 pGM1202-*ks007* suspension was added. The mixture was precipitated at 10000 rcf for 3 min and resuspended in the return flow. The resulting suspension was transferred to Cullum-medium agar plates (mannitol 20 g/L, soy-flour 20 g/L, agar 20 g/L, pH 7.3) containing 10 mM MgCl₂ and incubated at 30°C for 16 h. Subsequently, 1 mL nalidixic-acid solution (60 mg/mL in CH₃Cl) and 1 mL apramycin solution (100 mg/mL) were applied to the plates and allowed to set into the agar. The plates were incubated at 30°C for one week until colonies became visible. The visible colonies were transferred to Cullum-plates containing 60 μ g/mL nalidixic acid and 100 μ g/mL apramycin for selection, and were incubated for 5 days at 30°C. The presence of the *ks007* gene was verified by colony-PCR of the candidate colonies using the KS007-for and KS007-rev primers.

Isolation and purification of phileucin: *S. coelicolor* M1146 pGM1202-*ks007* was cultivated on a larger scale using 1 L of ISP2 medium (10 g/L malt extract, 4 g/L yeast extract, 4 g/L glucose, pH 7.5) containing 0.5 mg/L apramycin at 30°C, with a 120 rpm shaking speed. The expression of the gene was induced twice, once after 3 days and then after 5 days of incubation, by the addition of 0.05 mg/L thiostreptone. After 7 days the culture was extracted 2 times by liquid/liquid phase extraction using an equal amount of ethyl acetate. The ethyl acetate was evaporated and the crude extract dissolved in 10 mg/mL methanol/water (50/50 v/v). This extract was fractionated using Bakerbond™ SPE Octadecyl (C18) 2000 MG/6mL columns (Avantor Performance Materials, Center Valley, USA), resulting in a (50/50, v/v) MeOH/H₂O fraction and a 100% MeOH fraction. The 100% MeOH fraction was further processed by HPLC using a NUCLEOSHELL RP 18plus, 5 μ m column and 65%/35% methanol/water (v/v) as mobile phase, with a flow rate of 0.8 mL/min. The chromatogram showed a peak at 8 min with high absorption at 400 nm, which was roughly collected in multiple runs. For the final purification of the compound, the

composition of the mobile phase was altered to 63%/37% methanol/water (v/v) and the flow rate decreased to 0.7 mL/min.

Structure elucidation of phileucin: High-resolution mass spectrometry was performed using an ESI-MS micrOTOF spectrometer (Bruker Daltonik, Bremen, Germany). The structure elucidation was performed by NMR spectroscopic analysis of the purified compound. The measurements were performed using a Bruker Avance 500 DRX (Bruker, Billerica, USA) spectrometer operating at 500 MHz (^1H) and 125 MHz (^{13}C). Processing of the collected data was performed using Bruker XWIN-NMR 3.5 (Bruker). The signals were referenced to the respective solvent signal for CD_3OD ($\delta\text{H/C}$ 3.35/49.0). The molecular structure was elucidated, based on ^1H , ^{13}C , DEPT135, ^1H - ^1H COSY, ^1H - ^{13}C direct correlation (HSQC), and ^1H - ^{13}C long-range correlation (HMBC). The resulting signals were compared with values calculated by Advanced Chemistry Development (ACD/Labs) Software V11.02 (ACD/Labs).

Enzymatic assays: The proteases were assayed as described [9,10].

Phileucin

$[\alpha]_D$: +0.015 (*c* 1.00, acetone).

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IR (KBr): 2924.0, 1634.3, 1494.2, 1453.6, 960.0, 762.5, 717.6, 697.1 cm^{-1} .

UV/Vis λ_{max} (MeOH) nm (log ϵ): 206 (3.80), 328 (3.20)

^1H NMR (400 MHz, CDCl_3): 0.9 (3H, t, J = 7.5 Hz, H-1), 1.2 (1H, d, J = 7.0 Hz, H-4), 1.55 (1H, m, H-2b), 1.81 (1H, m, H-2a), 3.22 (1H, m, H-3), 3.88 (2H, s, H-9), 7.17 (1H, s, H-8), 7.30 (1H, m, H-13), 7.31 (1H, m, H-12), 7.31 (1H, m, H-14), 7.37 (1H, m, H-11), 7.37 (1H, m, H-11s)

^{13}C NMR (100 MHz CDCl_3): 12.3 (CH_3), 18.3 (CH_3) 28.7 (CH_2), 37.1 (CH_2), 37.6 (CH), 123.5 (CH), 128.2 (CH), 129.9 (CH), 129.9 (CH), 129.9 (CH), 129.9 (CH) 138.1 (C), 139.9 (C), 158.5 (C), 161.5 (C).

HRESI-MS, m/z : measured, 265.1311 [$\text{M} + \text{Na}^+$]; calcd. for $\text{C}_{15}\text{H}_{18}\text{N}_2\text{O}_\text{Na}$: 265.1316.

Supplementary data: The UV and IR spectra for phileucin are given.

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OM-X®, a Fermented Vegetables Extract, Facilitates Muscle Endurance Capacity in Swimming Exercise Mice

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The anti-fatigue effect was investigated of the probiotic supplement, OM-X®, on forced swimming capacity in mice. Mice were administered either vehicle (distilled water; DW) or OM-X® (85 mg/kg body weight) by gavage for 4 weeks. Forced swimming tests were conducted weekly using the Ishihara-modified Matsumoto swimming pool. The endurance swimming time of the final forced swimming exercise in mice fed with OM-X® group showed an approximately 2-fold increase compared with the vehicle control group. Biomedical parameters, including blood lactate, blood superoxide dismutase (SOD) activity, serum triacylglycerol (TG), hepatic total lipids (TL), TG and phospholipid (PL) were significantly lower in mice fed with OM-X® than those in the vehicle control group. Furthermore, the mRNA expression levels of carbamoyl phosphate synthetase 1 (Cps1) and arginase 1 (Arg1), in the urea cycle, were increased by OM-X® feeding. Thus, our findings suggest promotion of lipid metabolism and up-regulation of the urea cycle, at least in part, for the anti-fatigue effect mediated by OM-X®.

Keywords: OM-X®, Muscle endurance capacity, Lipid metabolism, Urea cycle, Anti-fatigue.

OM-X® is a probiotic supplement that was manufactured by Dr Ichiroh Ohhira using a unique fermentation method. It is strictly hand-made and is comprised of all natural components. OM-X® is a fermented mixture of vegetables, fruits, seaweeds, and mushrooms, using 12 strains of lactic acid bacteria (LAB) and bifidobacteria. After 5 years of fermentation at room temperature, the fermented mixture contains probiotics, prebiotics such as oligosaccharides and dietary fiber, and trace amounts of vitamins, minerals, short-chain fatty acids, and amino acids. In human clinical studies, OM-X® has shown beneficial effects on bone health [1], oral ulcerations [2], and colitis [3]. Recently we further reported the inhibitory effect of OM-X® on antigen-stimulated type I allergy [4]. The inhibitory mechanisms by OM-X® was due, at least in part, to the suppression of intracellular Ca²⁺ mobilization through inhibition of the translocation of cytosolic subunits of NADPH oxidase to membrane-bound subunits and direct scavenging of ROS that are produced by NADPH oxidase.

Most people are subjected to several stressors in their modern life, resulting in the accumulation of mental and physical fatigues. In the early stage of fatigue, favorable working conditions cannot be maintained, often resulting in impaired performances. Long-term accumulation of fatigue ultimately results in death or life-threatening diseases [5, 6]. Specifically, the recovery from fatigue is crucial for athletes to maintain their high performance in competitions. In order to avoid accumulating fatigue, athletes adopt several dietary and recovery methods that are individually matched. In recent studies, traditional plants [7], green tea [8, 9], astaxanthin [10], fucoidan [11] and deep seawater [12] have demonstrated beneficial effects in the suppression of fatigue accumulation. OM-X is a naturally-fermented plant-based probiotic containing herbs, seaweed, fruits and vegetables. Therefore, the aim of this study was

to evaluate the muscle endurance capacity in OM-X®-fed mice, and to prove that OM-X® is a beneficial naturally fermented extract against fatigue.

Changes in body weight, food consumption and water intake: The changes in body weight, food consumption, and water intake were similar between the vehicle control group and OM-X® group during the experimental period (Table 1).

Table 1: Body weight, food consumption, and water intakes in vehicle control mice and OM-X® fed mice for 4 weeks.

Group	water	OM-X
Body weight (g)		
0 w	32.1 ± 0.8	32.2 ± 1.6
1 w	35.1 ± 2.9	34.9 ± 2.1
2 w	40.0 ± 2.3	39.8 ± 2.3
3 w	42.1 ± 2.1	40.1 ± 3.0
4 w	43.5 ± 2.4	44.2 ± 2.4
Food consumption (g/day/mouse)		
0-1 w	3.0 ± 0.4	3.0 ± 0.5
1-2 w	5.8 ± 0.3	5.8 ± 0.4
2-3 w	5.0 ± 0.4	4.7 ± 0.7
3-4 w	5.3 ± 0.6	5.4 ± 0.3
Water intake (mL/day/mouse)		
0-1 w	5.9 ± 0.6	6.0 ± 0.4
1-2 w	7.0 ± 0.8	7.4 ± 0.6
2-3 w	7.0 ± 0.4	6.7 ± 0.4
3-4 w	7.4 ± 0.5	6.5 ± 0.8

Values are given as mean ± S. E. (each group n=17).

Effect of OM-X® on endurance capacity for swimming: To assess the effect of OM-X® on endurance capacity, we employed the swimming exercise using the Ishihara-modified Matsumoto swimming pool. As shown in Figure 1, intakes of OM-X® over 2 to 4 weeks gradually recovered the endurance swimming time up to the level of 1 week. The extended endurance swimming time in mice fed with OM-X® was significantly higher than that of the control group. Specifically, the endurance swimming time of the final forced swimming exercise in mice fed with OM-X® showed approximately a 2-fold increase compared with the vehicle control group.

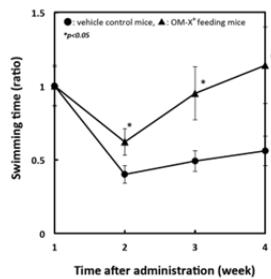


Figure 1: Effects of OM-X® on swimming exercise in mice. ●: vehicle control mice, ▲: OM-X® fed mice. Values are given as mean ± S.E. (each group n=17). $p < 0.05$ (*).

Changes in blood parameters and hepatic lipid profiles in mice fed with OM-X®: Serum components were analyzed after the last forced swimming test. Blood parameters and hepatic lipid profiles are shown in Table 2. The levels of blood lactate ($p < 0.01$), blood SOD activity ($p < 0.01$), serum TG ($p < 0.05$), hepatic TL ($p < 0.05$), TG ($p < 0.05$) and PL ($p < 0.05$) in mice fed with OM-X® were reduced to approximately 48, 25, 33, 26, 38, and 20%, respectively compared with those of vehicle control groups. The levels of BUN, creatinine and ketone bodies, which are the anti-fatigue and renal dysfunction indices, did not change between the vehicle control group and OM-X® group. Additionally, the activity of LDH, which is a myopathy index for over-exertion of muscle, was similar for both groups.

Table 2: Effects of OM-X® on serum biomedical parameters (blood glucose, BUN, creatinine, ketone bodies, LDH activity, lactate, SOD activity), serum lipid profiles, and hepatic lipid contents in mice.

Group	water	OM-X®
Blood glucose (mg/dL)	67.6 ± 4.7	64.0 ± 3.4
BUN (mg/100 mL)	71.9 ± 7.3	86.3 ± 6.3
Creatinine (mg/dL)	13.0 ± 0.9	12.1 ± 0.4
Ketone bodies (μmol/L)	132.6 ± 5.2	126.1 ± 8.3
LDH (Unit/L)	77.3 ± 12.9	82.6 ± 11.6
Lactate (mM)	5.3 ± 0.3	2.8 ± 0.2**
SOD activity (%)	55.6 ± 3.4	41.5 ± 2.1**
Serum lipid concentration		
TC (mg/100 mL)	219.3 ± 10.5	166.8 ± 9.2
TG (mg/100 mL)	366.7 ± 36.7	244.7 ± 13.2*
PL (mg/100 mL)	299.7 ± 20.2	298.8 ± 13.5
NEFA (mEq/L)	1.4 ± 0.1	1.6 ± 0.2
Hepatic lipid profiles		
Liver weight (g)	1.8 ± 0.6	1.7 ± 0.7
TL (mg/g liver)	59.7 ± 3.8	43.9 ± 5.5*
TC (mg/g liver)	8.8 ± 0.7	8.2 ± 0.5
TG (mg/g liver)	29.1 ± 3.1	17.9 ± 2.9*
PL (mg/g liver)	21.0 ± 2.0	17.3 ± 1.6*

Values are given as mean±S.E. (each group n=17).

(* $p < 0.05$, ** $p < 0.01$ vs. water)

Contents of liver and muscle glycogen: Stored glycogen in liver and muscle are very important resources for energy conservation and/or maintaining blood glucose in the physiologic range. During exercise, muscle glycogen is primarily consumed, and the levels of stored glycogen in liver and muscle are low. As shown in Figure 2, there was no change in liver and gastrocnemius muscle glycogen content between mice fed with OM-X® and the vehicle control mice.

The levels of 8-OHdG in liver and kidney: To evaluate the anti-oxidative effects of OM-X®, the levels of 8-OHdG in liver and kidney, as an oxidative damage index, were measured. There were also no differences in 8-OHdG levels between mice fed with OM-X® and the vehicle control mice (Figure 3).

The mRNA expression levels of the ornithine cycle enzyme in liver: The mRNA expression of Cps1, which acts as a rate-limiting enzyme in the ornithine cycle, and Arg1, which catalyzes the hydrolysis of arginine to ornithine and urea, were increased by OM-X® feeding (Figure 4). Specifically, the levels of Cps1 in mice fed with OM-X® significantly increased by approximately 3-fold

compared with the vehicle control mice. Furthermore, Arg1 levels in mice fed with OM-X® showed a trend towards an approximately 2-fold increase compared with vehicle control mice ($p < 0.1$), but this was not statistically significant.

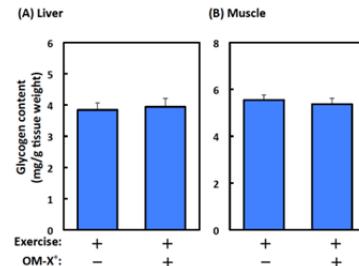


Figure 2: Effects of OM-X® on glycogen contents in exercise loaded-mice liver and muscle. (A) liver (B) muscle. Values are given as mean ± S.E. (each group n=17).

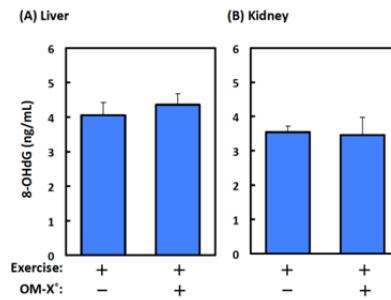


Figure 3: Effects of OM-X® on levels of 8-OHdG in exercise loaded-mice liver and kidney. (A) liver (B) kidney. Values are given as mean ± S.E. (each group n=17).

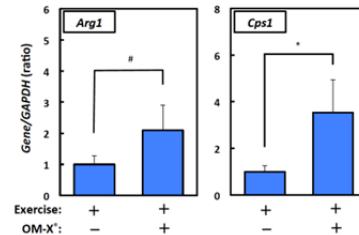


Figure 4: Effects of OM-X® on mRNA expression levels of enzymes of the urea cycle in swimming exercise loaded-mice liver. Values are given as mean ± S.E. (each group n=17). $p < 0.05$ (*), $p < 0.1$ (#), Student's t-test.

In this study, we examined the anti-fatigue effect (improvement of muscle endurance capacity) of OM-X® using a swimming test, and showed that OM-X® markedly prolonged the swimming time in mice (Figure 1). To delineate the mechanisms underlying this anti-fatigue effect, we further measured several biomedical parameters. The levels of blood lactate, serum TG, hepatic TL, TG, and PL in mice fed with OM-X® were significantly lower than those in the vehicle control group (Table 2). Blood lactate is a glycolysis by-product of carbohydrates under anaerobic conditions [16]. Brooks *et al.* reported that blood lactate, which is produced in fast-twitch oxidative-glycolytic fibers, is translocated to the heart and/or slow-twitch oxidative fibers, and then used as an energy substrate in mitochondria [17,18]. Therefore, the decrease in blood lactate levels in mice fed with OM-X® might be due to its utilization as a mitochondrial substrate for energy production. TG in blood, adipose tissue, and muscle fiber is one of the main building blocks of FFAs. During exercise, FFAs, which are produced from TG by lipoprotein lipase, are oxidized (by β -oxidation) in the mitochondria to supply energy [19, 20]. Thus, it was thought that the reduction of serum TG level in mice fed with OM-X® was due to the production of energy through β -oxidation of FFAs. Terao *et al.* also reported that the concentrations of cholesterol, TG, and PL of very low density lipoprotein (VLDL) in long-distance runners were maintained at

low levels compared with non-athletes, suggesting that VLDL is used as an energy source during running and exercise [21]. VLDL consists of TG (approx. 55%), cholesterol ester (approx. 12%), free cholesterol (approx. 7%), PL (approx. 18%), and protein (approx. 8%; apoprotein B100, -E, C1, -C2, and -C3) and is secreted mainly by the liver. The lowering of hepatic TL, TG and PL in mice fed with OM-X® was likely involved in VLDL formation for the production of energy.

Furthermore, exercise induces free radical formation in the liver and muscles [22]. These free radicals trigger oxidative damage such as lipid peroxidation. Despite the amount of swimming exercise in the OM-X® group being higher than that in the vehicle control group, the levels of 8-OHdG in the liver and kidney were similar between the two. As the almost 2-fold increase in swimming exercise in the OM-X® group did not result in a concomitant increase in oxidative damage, elevated SOD activity was presumed to contribute to the anti-oxidative effect of OM-X®.

The accumulation of ammonia in serum leads to fatigue during exercise. Since ammonia is a by-product of the metabolism of amino acids and other nitrogen compounds, the accumulated ammonia is metabolized via the ornithine cycle (urea cycle) in the liver [23]. Therefore, the up-regulation of mRNA expression and activities of hepatic enzymes involved in ammonia metabolism contributes to the reduction of fatigue. As shown in Figure 4, the expression level of Cps1, which catalyzes the first reaction in the ornithine cycle wherein ammonia and bicarbonate combine to form carbamoyl, in mice fed with OM-X® was approximately 3-fold higher than in vehicle control mice. Although a remarkable difference was not observed in expression of Arg1, which catalyzes the hydrolysis of L-arginine into ornithine and urea in the ornithine cycle, a trend to up-regulation was observed in OM-X® fed mice. These results suggest that intake of OM-X® is capable of regulating the expression of hepatic enzymes involved in ammonia metabolism, resulting in an inhibition of the accumulation of ammonia in blood, eventually leading to lesser muscle fatigue and prolongation of exercise.

Since OM-X® also includes a sugar (55.4%), it absolutely contributes to the prolongation of exercise. Recent studies indicated that the up-regulation of glucose transporter type 4 expression through the improvement of insulin sensitivity in muscle and 5' adenosine monophosphate-activated protein kinase activation contributes to prolongation of exercise [24, 25]. We further need to study glucose metabolism in OM-X® fed mice. Based on our results, the mechanisms of anti-fatigue effect by OM-X® were due, at least in part, to (i) catabolism of glucose in OM-X®, (ii) promotion of lipid metabolism, and (iii) up-regulation of the urea cycle. It should be noted that OM-X® is a beneficial natural extract for anti-fatigue.

Experimental

OM-X® preparation: The OM-X® sample was provided by BIOBANK Co., Ltd. (Okayama, Japan). All plant ingredients used in OM-X® are nutritionally rich and safe for human consumption. The general procedure involves loading each ingredient into the manufacturing vats one by one at different stages of fermentation. When each ingredient is added, it is fermented by LAB to produce various organic acids and amino acids. In addition, the solid ingredients are broken down to facilitate absorption of their components through fermentation. Thus, various ingredients that are beneficial to human health are fermented and matured for 5 years [1-4].

Animals and diet: Four-week-old male ddY mice (Japan SLC, Inc., Hamamatsu, Japan), weighing 20–22 g were housed in plastic cages in an air-conditioned room ($22 \pm 2^\circ\text{C}$) with a 12-h light and dark cycle (lighting from 0700 to 1900 h). All mice were fed commercial MF pellets (Oriental yeast Co., Ltd., Tokyo, Japan) and water *ad libitum* for 1 week to accustom them to the surroundings.

Swimming exercise: First, mice were subjected to a swimming exercise for 15 min to check their swimming performance. The mice that did not perform a self-motion were removed from this study. Thirty-four mice were then divided into 2 groups (n=17 per group) equally based on body weight and swimming activity. They were administrated either vehicle (distilled water; DW) or OM-X® at a dose of 85 mg/kg of body weight by gavage for 4 weeks. Forced swimming tests were then conducted weekly using the Ishihara-modified Matsumoto swimming pool [13-15]. The swimming pool (length×width×depth=90×45×45 (cm³)) was filled with water to a depth of 38 cm (Figure 5A). A current was generated in the pool using a pump (type C-P60H; Hitachi, Tokyo, Japan), and the current strength was adjusted by a valve. The current speed at the surface was measured with a digital current meter (model SV-101-25S; Sankou, Tokyo, Japan) at the start of every swimming session (10 L/min) and was maintained at a constant speed (14 L/min). The water was maintained at a temperature of $27 \pm 0.5^\circ\text{C}$ using an electric heater. The end point of the swimming test was defined as the time at which the mice could not resist the current and failed to rise to the surface of the pool within 7 s to breathe (Figure 5B). After swimming until the end point, mice were immediately rescued and their bodies wiped with a towel. Upon completion of the test on the final day of the swimming exercise, mice were sacrificed by isoflurane anesthesia, and blood was immediately collected using a 1 mL syringe without heparin to measure blood glucose level, blood urea nitrogen (BUN), creatinine, ketone bodies, lactate, total cholesterol (TC), TG, PL, non-esterified fatty acid (NEFA), transforming growth factor (TGF)-β, lactate dehydrogenase (LDH), and SOD activity. The liver, kidney, and gastrocnemius muscle were then excised immediately, and kept at -80°C until analysis. This animal experiment was approved by the Kindai University animal use committee, and the animals were maintained according to the guidelines of Kindai University for the care of laboratory animals.

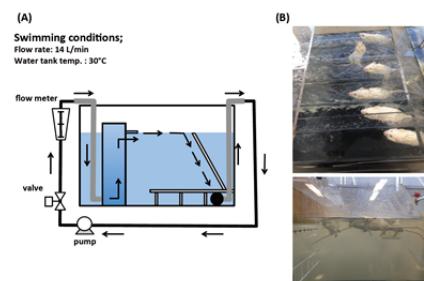


Figure 5: Equipment for forced swimming exercise. (A) Schematic diagram of the Ishihara-modified Matsumoto swimming pool, (B) appearance of swimming exercise.

Analytical procedures: The profile of serum TC, TG, PL, NEFA, blood glucose, BUN, creatinine, and ketone bodies was measured with clinical analysis kits {cholesterol CII, triglyceride G, phospholipid B, non-esterified fatty acid C test, glucose CII test, BUN-B test, creatinine test, and total ketone bodies-test, (Wako Pure Chemical Industries, Ltd., Osaka, Japan)} and lipid contents in the liver, extracted by the method of Folch *et al.*, were measured in the same way. The concentration of lactate in serum was measured using a Lactate Colorimetric assay kit (BioVision Inc., CA, USA).

8-Hydroxy-2'-deoxyguanosine (8-OHdG) determination: To determine the anti-oxidative effects of OM-X® in exercised mice, we measured the 8-OHdG levels in either liver or kidney as an oxidative damage index. DNA was extracted from each tissue lysate, which was homogenized using a bead beater-type homogenizer µT-01 (TAITEC, Koshigaya, Saitama, Japan, φ5 mm SUS bead, 3000 rpm/min for 1 min) using the DNA Extractor kit (Wako). The extracted DNA from each tissue was hydrolyzed with 8-OHdG assay preparation reagent (Wako), and then measured using the 8-OHdG enzyme-linked immunosorbent assay kit (Japan Institute for the Control Aging, NIKKEN SEIL Co., Ltd, Shizuoka, Japan), and normalized to DNA concentration.

Contents of liver and muscle glycogen: The contents of liver or muscle were measured with a glycogen colorimetric assay kit (BioVision Inc.). Frozen liver or gastrocnemius muscle (10 mg) was homogenized with glycogen development buffer in a 2 mL tube using a bead beater-type homogenizer µT-01 (TAITEC, φ5 mm SUS bead, 3000 rpm/min for 1 min). The homogenized sample was centrifuged at 8,000×g, 4°C for 5 min, and the supernatant was assayed using the glycogen colorimetric assay kit.

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Quantitative real time-PCR: Total RNA was extracted from cells using a RNeasy Mini Kit (QIAGEN Inc, Valencia, CA, USA). cDNA was synthesized from 0.25 mg of total RNA using a PrimeScript reagent kit (Takara Bio, Ohtsu, Japan). cDNA was subjected to quantitative RT-PCR using an Applied Biosystems StepOne™ Real-Time PCR system (Foster City, CA, USA). Primers for arginase 1 (Arg1), carbamoyl phosphate synthetase 1 (Cps1) and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were purchased from Takara Bio. The expression level of each gene was determined using the comparative Ct method and normalized to that of GAPDH, which was used as an internal control. The PCR reaction consisted of 45 cycles (95°C for 10 s and 60°C for 40 s) after an initial denaturation step (95°C for 10 min).

Statistical analysis: All data were analyzed using the Mac statistical analysis software package for Macintosh, version 2.0 (Esumi Co., Tokyo, Japan). All data are expressed as means ± standard errors (SEs). Statistical analysis was performed using the Student's t-test, and differences were considered significant when p-values were less than 0.05.

Essential Oils of *Morus alba* and *M. nigra* Leaves: Effect of Drying on the Chemical Composition

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Detailed GC and GC/MS analyses of essential-oil samples obtained by hydrodistillation of *Morus alba* L. and *M. nigra* L. leaves (four samples) allowed the identification of 131 constituents representing 95.1 – 96.4% of the total GC peak areas. The most abundant compounds classes were alkanes, diterpenoids, carotenoid-derived compounds and fatty acid-related constituents with *trans*-phytol (7.9 – 71.2%), (*E,E*)-geranyl linalool (0.2 – 8.0%), (*Z*)-bovolide (8.1%) and *n*-chain alkanes (in total, 17.5 – 52.4%) as the major constituents of the analyzed samples. In general, there were only quantitative differences noted between *M. nigra* essential oils from fresh and dry leaves. The most discernable changes included a variation in the content of the major constituents (e.g. the relative amount of *trans*-phytol and the total amount of alkanes decreased by 5.8% and ≈ 2%, respectively, while that of (*E,E*)-geranyl linalool increased by 7.8%). On the other hand, the composition of *M. alba* essential oil was much more significantly affected by the drying process. The highest quantitative differences were noted for *trans*-phytol, geranyl acetone, and all isomers of 4,6,8-megastigmatrien-3-one. Also, a rare plant metabolite, (*Z*)-bovolide, characteristic of leaf senescence, was only identified (8.1%) in the essential oil of *M. alba* dried leaves.

Keywords: *Morus alba*, *Morus nigra*, Metabolomics, Apocarotenoids, (*Z*)-Bovolide, Storage.

The genus *Morus* L. (Moraceae), commonly known as mulberries, comprises about 16 species of deciduous, wild-growing/cultivated trees that are spread across many temperate world regions, especially in the Northern Hemisphere [1]. Species of this genus have a long history of use in folk medicine, with a number of interesting biological/pharmacological properties already scientifically proven [2]. Specifically, *Morus alba* L. and *M. nigra* L. were shown to possess analgesic, sedative, diaphoretic, antihypertensive, antityrosinase and antidiabetic activities [2]. Only three *Morus* species (*M. alba*, *M. nigra* and *M. rubra*) are part of the Serbian flora [1]. *M. alba* (white mulberry or ‘beli dud’ in Serbian) is a tree native to northern China. Nowadays, white mulberry is a worldwide naturalized species cultivated to feed silkworms in the commercial production of silk [1]. A SciFinder search revealed more than 11,000 reports dealing with *M. alba*. However, only seven reports dealt with the essential-oil analysis of this particular taxon. These studies showed that the essential oil from *M. alba* leaves contained high amounts of alkanes, diterpenes and sesquiterpenes [3], while the major constituents of the essential oil isolated from dry fruits were 1,8-cineole, geraniol, linalyl acetate, α -pinene and limonene [4]. *M. nigra*, known as black mulberry or ‘crni dud’ in Serbian, is a deciduous fruit-bearing tree, native to Iran and Afghanistan [1]. Black mulberries are much less frequently cultivated than white mulberry trees [1]. Interestingly, based on archives from the 13th century, a black mulberry tree, planted in the courtyard of the Patriarchate of Peć, is one of the oldest plant organisms in Serbia [1]. Up to now, non-volatile secondary metabolites of *M. nigra* were the subject of several studies [5]. However, there are no previous studies on the chemical composition of the essential oil of this plant species.

The drying process of plant material has a crucial role in preventing the growth of microorganisms and, also, makes storage and transportation of plant material much easier [6]. During the drying process, evaporation of volatile phytochemicals and/or the formation of artefacts, as a result of external (e.g. oxidation by

atmospheric oxygen) and internal ((non)enzymatic) degradation/transformation, can cause changes in the aroma and medicinal potential of plant material [6]. Several studies reported significant changes in the biological/pharmacological properties (e.g. antioxidant capacity), and changes in color and chemical composition of the essential oil as a consequence of plant material drying [6]. *Morus* spp. leaves can be used, for medicinal purposes, either fresh or dried [2]. However, *Morus* leaves, like other herbal teas or medicinal plants, are usually stored and consumed in dried form. For that reason, the current work is aimed to analyze the composition of the essential oil from both fresh and dried leaves of two *Morus* species (*M. alba* and *M. nigra*) cultivated in Serbia and to evaluate the influence of the drying process on the volatiles of the leaves.

Detailed GC and GC/MS analyses of four essential-oil samples obtained by hydrodistillation of both fresh and dried *M. alba* and *M. nigra* leaves, collected from mulberry trees cultivated in SE Serbia, allowed, in total, the identification of 131 constituents that represented 95.1 – 96.4% of the total GC peak areas (Table 1). The most abundant classes of compounds identified were alkanes and diterpenoids that together constituted more than 60% of sample MA-2, and more than 88% of the samples MN-1, MN-2, and MA-1. The remaining identified volatiles were carotenoid derivatives (1.4 – 24.8%) and fatty acid related constituents (tr – 8.1%). The major contributors of *M. nigra* essential oils were: *trans*-phytol (65.4 – 71.2%), (*E,E*)-geranyl linalool (0.2 – 8.0%) and normal chain alkanes (tricosane (2.0 – 2.8%), pentacosane (2.4 – 3.8%), heptacosane (1.9 – 2.7%), nonacosane (3.4 – 4.1%) and hentriacontane (2.4 – 3.5%)). Similarly, the bulk, by percentage, of both *M. alba* essential oils was comprised of alkanes, diterpenoids and carotenoid derivatives. *trans*-Phytol (7.9 – 61.6%) was the predominant detected constituent, followed by *cis*-bovolide (8.1%), pentacosane (8.2 – 10.9%), heptacosane (3.7 – 7.9%), nonacosane (4.2 – 12.4%) and hentriacontane (3.2 – 12.4%).

Table 1: Chemical composition of the essential oils isolated from fresh and dried leaves of *Morus alba* and *M. nigra* from Serbia.

Compound	Class ^{a)}	RI ^{b)}	<i>Morus nigra</i> ^{c)}		<i>Morus alba</i> ^{d)}		ID ^{e)}
			MN-1	MN-2	MA-1	MA-2	
Relative content [%] ^{f)}							
Furfural	FA	828	-	-	tr	-	f, g, h
(E)-3-Hexen-1-ol	FA	844	-	-	tr	-	f, g
Tiglic acid	FA	903	-	-	tr	-	f, g, h
4-Methylnonane	A	957	tr	-	-	-	f, g
Benzaldehyde	O	959	-	-	tr	-	f, g, h
Decane	A	1000	tr	-	-	-	f, g, h
Benzyl alcohol	O	1034	-	tr	tr	-	f, g, h
2-Phenylethanal	O	1051	tr	-	0.1	-	f, g
4-Methyldecane	A	1057	tr	-	-	-	f, g
2-Methyldecane	A	1068	tr	-	-	-	f, g
p-Cresol	O	1069	-	-	tr	-	f, g
Undecane	A	1100	1.0	-	-	-	f, g
Dehydroabina ketone	M	1123	-	-	tr	-	f, g
Iosphorone	C	1125	tr	-	-	-	f, g
3,7-Dimethyldecane ^{k)}	A	1125	tr	-	-	-	f, g
4-Ketoisophorone	C	1145	-	-	tr	-	f, g
Benzoic acid	O	1155	-	-	tr	-	f, g, h
Octanoic acid	FA	1162	-	-	tr	-	f, g, h
2-Methylundecane	A	1163	0.2	-	-	-	f, g
3-Methylundecane	A	1170	tr	-	-	-	f, g
Dodecane	A	1200	2.0	-	-	-	f, g, h
Safranal	C	1202	tr	tr	tr	f, g	
2,6-Dimethylundecane	A	1213	0.4	-	-	-	f, g
2-Methyldodecane	A	1264	0.3	-	-	-	f, g
3-Methyldodecane	A	1270	0.3	-	-	-	f, g
Dihydrodulau I	C	1298	-	-	tr	-	f, g
Tridecane	A	1300	1.8	tr	-	-	f, g, h
2,6-Dimethyldodecane	A	1311	tr	-	-	-	f, g
4-Vinylguaiacol	O	1311	-	tr	tr	f, g	
1-Methylnaphthalene	O	1315	tr	-	-	-	f, g
3,7-Dimethyldodecane	A	1317	tr	-	-	-	f, g
Eugenol	O	1353	-	-	tr	-	f, g, h
2-Methyltridecane	A	1363	tr	-	-	-	f, g
3-Methyltridecane	A	1370	tr	-	-	-	f, g
2,6,10-Trimethyldodecane ^{k)}	S	1375	0.2	-	-	-	f, g
(Z)-β-Damascenone	C	1380	-	-	tr	-	f, g
1-Undecanol	FA	1385	tr	-	-	-	f, g
1-Tetradecene	FA	1391	tr	-	-	-	f, g
Tetradecane	A	1400	0.8	-	tr	-	f, g, h
(E)-β-Damascenone	C	1409	-	-	tr	f, g	
2,6-Dimethyltridecane	A	1413	tr	-	-	-	f, g
Geranyl acetone	C	1447	-	-	-	9.8	f, g, h
Unidentified constituent ^{j)}	A	1461	tr	tr	0.1	3.2	
2-Methyltetradecane	A	1463	tr	-	-	-	f, g
3-Methyltetradecane	A	1470	tr	-	-	-	f, g
3,4-Dehydro-β-ionone	C	1476	0.2	-	-	tr	f, g
(E)-β-Ionone	C	1479	tr	-	tr	2.6	f, g
Pentadecane	A	1500	0.3	-	tr	-	f, g, h
2-Tridecanone	FA	1502	tr	-	tr	-	f, g
(E,E)-o-Farnesene	S	1503	tr	-	-	-	f, g
3,4-Dimethyl-5-pentyl-							
2(5H)-furanone (syn. dihydrobovolide)	FA	1512	-	-	-	tr	f, g
(Z)-3,4-Dimethyl-5- pentylidene-2(5H)- furanone (syn. cis- bovolide, (Z)-bovolide) ^{l,m)}	FA	1518	-	-	-	8.1	f, g
Caparatriene ^{k)}	S	1523	-	-	-	tr	f, g
(6Z,8E)-4,6,8- Megastigmatrien-3-one	C	1559	-	tr	0.1	tr	f, g
(E)-Nerolidol	S	1559	-	-	tr	-	f, g
2-Methylpentadecane	A	1563	tr	-	-	-	f, g
3-Methylpentadecane	A	1570	tr	-	-	-	f, g
(Z)-3-Hexenyl benzoate	FA	1571	-	-	tr	-	f, g
(6Z,8Z)-4,6,8- Megastigmatrien-3-one	C	1577	0.3	1.2	0.3	4.3	f, g
2-Methyl-4-(2,6,6- trimethyl-1-cyclohexen-1- yl)-2-butenal (syn. boronia butenala) ^{k)}	C	1582	-	-	tr	-	f, g
Hexadecane	A	1600	tr	-	-	-	f, g, h
(6E,8E)-4,6,8- Megastigmatrien-3-one	C	1610	tr	tr	tr	tr	f, g
Tetradecanal	FA	1613	-	-	tr	-	f, g, h
(6E,8Z)-4,6,8- Megastigmatrien-3-one	C	1623	0.2	0.6	0.2	4.3	f, g
2-Methylhexadecane	A	1663	-	-	tr	-	f, g
1-Tetradecanol	FA	1678	-	-	tr	-	f, g, h
2-Pentadecanone	FA	1696	tr	-	tr	-	f, g, h
Heptadecane	A	1700	tr	tr	tr	tr	f, g, h
Pentadecanal	FA	1714	tr	-	tr	tr	f, g, h
Tetradecanoic acid	FA	1758	-	-	tr	-	f, g, h
Benzyl benzoate	FA	1768	-	-	tr	-	f, g
3-Methylheptadecane	A	1771	-	-	tr	-	f, g
Octyl octanoate	FA	1776	-	-	tr	-	f, g
Phenanthrene	O	1784	-	-	tr	-	f, g
1-Octadecene	FA	1793	-	-	tr	-	f, g
Octadecane	A	1800	tr	tr	tr	tr	f, g, h
Hexadecanal	FA	1816	-	-	tr	-	f, g

Neophytadiene (isomer I)	D	1835	tr	-	tr	-	f, g
Hexahydrofarnesyl acetone	C	1840	2.0	3.0	0.8	3.8	f, g
1-Hexadecanol	FA	1882	-	-	tr	-	f, g
1-Nonadecene	FA	1893	-	-	tr	-	f, g
Nonadecane	A	1900	tr	tr	0.1	tr	f, g, h
(E,E)-Farnesyl acetone	C	1909	-	-	tr	tr	f, g
Methyl palmitate	FA	1924	tr	tr	0.1	tr	f, g
Isophytol	D	1945	tr	tr	0.1	-	f, g
Hexadecanoic acid	FA	1969	-	-	2.0	-	f, g, h
Ethyl hexadecanoate	FA	1992	tr	-	-	-	f, g
1-Eicosene	FA	1994	-	-	tr	-	f, g
Eicosane	A	2000	tr	tr	0.1	tr	f, g, h
(E,E)-Geranyl linalool	D	2022	0.2	8.0	0.6	tr	f, g
Methyl (Z,Z)-9,12,15- octadecatrienoate	FA	2096	0.4	tr	0.3	-	f, g
Heneicosane	A	2100	0.3	tr	0.7	tr	f, g, h
trans-Phytol	D	2113	71.2	65.4	61.6	7.9	f, g
Octadecanoic acid	FA	2160	-	-	0.3	-	f, g
(Z,Z)-9,12,15- Octadecatrienoic acid	FA	2164	0.2	-	0.3	-	f, g
1-Docosene	FA	2192	-	-	0.2	-	f, g
Docosane	A	2200	0.2	tr	0.5	tr	f, g, h
Eicosano	FA	2213	-	-	tr	-	f, g
2-Methylhexadecane	A	2265	-	tr	tr	-	f, g
1-Tricosene	FA	2297	-	-	0.1	-	f, g
Tricosane	A	2300	2.0	2.8	2.8	3.0	f, g, h
2-Heneicosane	FA	2307	-	-	tr	-	f, g
δ-Octadecalactone	FA	2347	-	-	tr	0.2	f, g
3-Methyltricosane	A	2374	-	-	tr	-	f, g
1-Tetracosene	FA	2396	tr	tr	0.7	tr	f, g
Tetracosane	A	2400	0.5	0.6	1.4	3.0	f, g, h
2-Docosanone	FA	2410	-	-	tr	-	f, g
2-Methyltetracosane	A	2465	-	-	tr	-	f, g
1-Pentacosene	FA	2497	-	-	0.8	-	f, g
Pentacosane	A	2500	2.4	3.8	8.2	10.9	f, g, h
1-Hexacosene	FA	2597	tr	tr	0.5	tr	f, g
Hexacosane	A	2600	tr	tr	0.7	2.8	f, g, h
2-Methylhexacosane	A	2665	-	-	tr	-	f, g
1-Tetracosanol	FA	2677	-	-	tr	-	f, g
1-Heptacosene	FA	2697	-	-	tr	-	f, g
Heptacosane	A	2700	1.9	2.7	3.7	7.9	f, g, h
Pentacosanal	FA	2714	-	-	tr	-	f, g
3-Methylheptacosane	A	2775	-	-	tr	-	f, g
1-Octacosene	FA	2797	tr	-	0.2	tr	f, g
Octacosane	A	2800	tr	tr	0.2	tr	f, g, h
2-Methyloctacosane	A	2864	-	-	tr	-	f, g
Nonacosane	A	2900	3.4	4.1	4.2	12.4	f, g, h
3-Methylnonacosane	A	2975	-	-	tr	-	f, g
1-Triacotene	FA	2998	-	-	0.1	tr	f, g
Triacotane	A	3000	tr	tr	0.1	tr	f, g, h
1-Hexacosyl acetate	FA	3013	-	-	tr	0.2	f, g
2-Methyltriacontane	A	3065	-	-	tr	-	f, g
Henriciacontane	A	3100	2.4	3.5	3.2	12.4	f, g, h
1-Dotriacontene	FA	3198	-	-	tr	-	f, g
Dotriacontane	A	3200	tr	-	tr	tr	f, g, h
1-Octacosyl acetate	FA	3213	-	-	tr	-	f, g
Triotriacontane	A	3300	tr	tr	0.2	tr	f, g, h

^{a)} For compound class abbreviations see bottom of this Table. ^{b)} RI = retention indices determined relative to a homologous series of *n*-alkanes (C_8 - C_{33}) on a DB-5MS column. ^{c)} MN-1 = Essential-oil sample obtained by a hydrodistillation of *Morus nigra* fresh leaves, i.e. immediately after the collection of plant material, MN-2 = Essential-oil sample obtained by a hydrodistillation of *M. nigra* leaves dried for one month at room temperature. ^{d)} MA-1 = Essential-oil sample obtained by a hydrodistillation of *M. nigra* fresh leaves, i.e. immediately after the collection of plant material, MA-2 = Essential-oil sample obtained by a hydrodistillation of *M. alba* leaves dried for one month at room temperature. ^{e)} ID = Identification method, f = constituent identified by a mass spectral comparison with those given in the literature [7], g = constituent identified by experimental retention index (DB-5MS column) matching with literature data [7], h = constituent identity confirmed by GC co-injection of an authentic standard. ^{f)} tr = trace (< 0.05%); - = not detected. ⁱ⁾ MS (EI, 70 eV), m/z (rel. int., %): 178 (21, [M⁺]), 163 (42), 151 (6), 150 (47), 136 (12), 135 (100), 122 (7), 121 (30), 109 (13), 108 (11), 107 (47), 105 (7), 95 (19), 94 (7), 93 (31), 91 (25), 85 (7), 81 (25), 80 (10), 79 (65), 78 (9), 77 (34), 71 (16), 70 (8), 69 (47), 67 (15), 66 (9), 65 (16), 57 (17), 55 (25), 53 (17), 52 (6), 51 (10), 43 (18), 42 (6), 41 (41), 40 (5), 39 (24). ^{k)} Relative stereochemistry was not determined. ^{l)} Mass spectrum and retention index were reported by G. MacLeod [8]. ^{m)} MS (EI, 70 eV), m/z (rel. int., %): 181 (4), 180 (30, [M⁺]), 165 (5), 138 (15), 137 (87), 125 (29), 124 (100), 123 (8), 110 (5), 109 (12), 96 (10), 95 (6), 83 (7), 82 (14), 81 (17), 79 (6), 67 (5), 55 (59), 54 (16), 53 (10), 41 (11), 39 (12).

In general, there were only quantitative variations noted among *M. nigra* essential-oil constituents from fresh leaves (MN-1;

hydrodistilled immediately after the collection of plant material) and dried leaves (**MN-2**; the essential oil was isolated after a one-month drying period at room temperature). The most discernable changes included an alteration in the content of the major constituents: the content of *trans*-phytol and the total amount of alkanes decreased by 5.8 and \approx 2%, respectively, while the relative amount of (*E,E*)-geranyl linalool increased by 7.8%. Contrary to this mostly quantitative compositional change in the *M. nigra* essential-oil samples during the drying of leaves, the composition of the corresponding *M. alba* essential oils (**MA-1** and **MA-2**) differed not only quantitatively, but also qualitatively. Again, *trans*-phytol was the constituent that displayed the most significant drop in its relative content; this time its relative amount was lower by *ca.* 8 times; the percentage of geranyl acetone and all isomers of 4,6,8-megastigmatrien-3-one increased after the period of drying (Table 1). Also, a γ -lactone, *cis*-bovolide, appeared in the essential oil of *M. alba* dry leaves among the major contributors (8.1%).

Carotenoid-derived compounds (apocarotenoids or norisoprenoids) reached 24.8% of the total essential oil from sample **MA-2**, while compounds of this class formed only 1.4% of the oil sample from fresh *M. alba* leaves (**MA-1**). Thus, not only the relative content of the carotenoid fraction, but also its qualitative composition, distinguished the samples of the essential oil from fresh and dried *M. alba* leaves (**MA-1** and **MA-2**; Table 1). In general, the flavor of plant material is one of the main sensory properties used in its selection, acceptance and ingestion [9]. The obtained differences, i.e. the increased amount of carotenoid derivatives, suggested that the drying process has a crucial rule for the known flavor of tea (water infusion) prepared from dried *M. alba* leaves. For example, β -ionone (violet-like, woody, floral, fruity flavor), present in both *M. alba* essential-oil samples (**MA-1** and **MA-2**; Table 1), is the main contributor to the aroma of roses despite its relatively low concentration in rose oil [10].

The increased amount and the large structural diversity of apocarotenoids found in *M. alba* dried leaves can be attributed mainly to the catalytic versatility of a family of carotenoid cleavage dioxygenases (CCDs) that catalyze the synthesis of apocarotenoid volatiles only at the leaf senescence stage, even though the CCD enzymes are present throughout the development of this plant organ [10]. A proposed biosynthetic pathway, linking most of the identified apocarotenoids from the dried *M. alba* leaves with their common precursor (phytoene), is depicted in Figure 1. Based on the structures of the biosynthesized apocarotenoids (C_{13} -norisoprenoids - β -ionone, 3,4-dihydro- β -ionone, safranal, β -damascenone and all diastereoisomers of megastigma-4,6,8-trien-3-one) it seems that the bio-oxidative cleavage reactions that involved 9,10 (9',10')-dioxygenase (CCD1, CCD7 and/or AtCCD1 enzyme, which cleave the 9,10 double bonds of their respective carotenoid substrates [10]), were the dominant ones (Figure 1).

The sample of essential oil from dried *M. alba* leaves contained a compound eluting at RI 1518 that accounted for 8.1% of the oil; this was completely absent from all other oils. A direct comparison of the mass spectrum of the compound in question (molecular ion at *m/z* 180, base peak at *m/z* 124; Figure 2) with those from the available commercial mass spectral libraries (Adams [7], Wiley 6, NIST11 and MassFinder 2.3) did not result in a positive hit. From the experience of the authors, the very same compound is known to have been the constituent of a number of other essential oils, but remained unidentified [11]. The first clues to its identity came from a consideration of another constituent present in that essential oil sample, namely, dihydrobovolide, identified based on a comparison with a mass spectrum and retention index from MassFinder 2.3, eluting (RI 1512) just before the compound of interest.

Dihydrobovolide possesses a mass spectrum that displayed ions at *m/z* values that were 2 amu higher than those of the yet unidentified constituent, including its molecular ion (at *m/z* 182). This led us to consider the possibility that the compound at RI 1518 represents a dehydro derivative of dihydrobovolide. A literature search revealed that such a compound exists: (*E/Z*)-3,4-dimethyl-5-pentylidene-2(5*H*)-furanone, known as (*cis/trans*)-bovolide (Figure 2). A comparison of the reported [8] mass spectrum and RI value, for the column of the same polarity, of bovolide with those that correspond to the component from our oil corroborated the tentative identification. Thus, the component at RI 1518 was identified as one of the geometric isomers of bovolide.

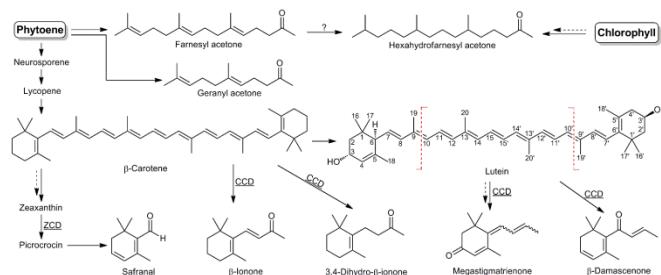


Figure 1: A proposed biosynthetic pathway for the formation of apocarotenoids in the essential oil of *M. alba* dried leaves starting from a single substrate - phytoene. ZCCD is zeaxanthin cleavage dioxygenase and CCD is carotenoid cleavage dioxygenase.

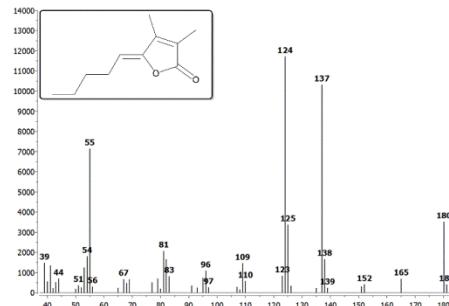


Figure 2: Mass spectrum and structure of *cis*-bovolide ((*Z*)-3,4-dimethyl-5-pentylidene-2(5*H*)-furanone).

Bovolide was first isolated and identified from butter, and was so-named because of its bovine origin [12]. Previously, Horita *et al.* confirmed that bovolide and dihydrobovolide were formed by light-induced post-harvesting biochemical transformations of native plant metabolites, usually accompanying the drying process of plant material [12c]. Although two diastereoisomers of bovolide could exist ((*Z*)- and (*E*)-3,4-dimethyl-5-pentylidene-2(5*H*)-furanone), only one is known to form as a result of such transformations of native fatty acid related plant metabolites. The dominant formation of the *Z* isomer was previously noted in the literature [12d], and that fact could be understood in terms of the apparent large difference in the minimal potential energies of the two diastereoisomers. To verify this, we calculated, using ChemBio 3D Ultra 12.0 software, at the MM2 level of theory, the ΔE mentioned between the conformations with the minimal energy of the two diastereoisomers. It amounted to \approx 4 kcal/mol (9.4635 and 13.6868 kcal/mol for (*Z*)-3,4-dimethyl-5-pentylidene-2(5*H*)-furanone and (*E*)-3,4-dimethyl-5-pentylidene-2(5*H*)-furanone, respectively). For comparison sake, the *E* isomer of 3-hexene was only 0.21 kcal/mol more stable than the *Z* (the difference was calculated in the same manner). Such a difference in stability suggests that, in the equilibrium between the isomers, at room temperature, one would expect a domination of (*Z*)-bovolide (more than 99.99% in equilibrium). Thus, we believe that the isomer of bovolide that we detected is most probably also the thermodynamically most stable one, i.e. *Z*-isomer. It appears to

us that the trivial names *cis*-bovolide, or only bovolide (CAS No. 774-64-1), are rather confusing with regard to the stereochemistry of this compound. It is difficult to understand the meaning of *cis* in this particular case (four different groups attached to the double bond in question). We believe that the stereodesignator Z is much more appropriate for use, and even better to also state the nomenclature name along with the trivial one. Bovolide and dihydrobovolide, which are not connected by the regular metabolic pathway of fatty acid biosynthesis [12a,12b], were previously identified in a range of different teas [12c] and as constituents of tobacco, peppermint oil, *Lycium chinese* and *Carphephorus corymbosus* essential oils [13]. Both compounds possess a strong aroma described as celery-like and were patented for improving the flavor of tobacco [12a,12b]. Interestingly, their presence was used as a marker for exposure of plant material to light [12c].

Experimental

Plant material: Leaves of *Morus nigra* and *M. alba*, collected in July, 2015 (village Gornji Barbeš, near Gadžin Han, Serbia), were identified by Marija Marković; voucher specimens (VM0012 and VM0013) were deposited in the Herbarium of the Faculty of Science and Mathematics, University of Niš, Serbia. A portion of the leaves was subjected to hydrodistillation immediately after collection, while another portion was dried for 1 month at room temperature, without direct exposure to sunlight.

Isolation of essential oils: Fresh (3 x 300 g batches) and dried leaves (3 x 100 g batches) were subjected to hydrodistillation for 4.5 h using the original Clevenger-type apparatus [6b]. The yields of essential oils were 0.0011 and 0.0012% from the dried plant

material (**MN-2** and **MA-2**, respectively) and 0.0080% (w/w) from both samples of fresh leaves (**MN-1** and **MA-1**).

GC-MS analysis: GC-MS analyses (3 repetitions) were carried out using a Hewlett-Packard 6890N gas chromatograph equipped with a fused silica capillary column DB-5MS (5% phenylmethylsiloxane, 30 m × 0.25 mm, film thickness 0.25 µm, Agilent Technologies, USA) and coupled with a 5975B mass selective detector from the same company. The injector and interface were operated at 250 and 300°C, respectively. Oven temperature was raised from 70 to 290°C at a heating rate of 5°C/min and the program ended with an isothermal period of 10 min. As a carrier gas helium at 1.0 mL/min was used. The samples, 1.0 µL of essential oil solutions in diethyl ether (10.0 mg of an essential oil sample per 1.0 mL of solvent), were injected in a pulsed split mode (the flow was 1.5 mL/min for the first 0.5 min and then set to 1.0 mL/min throughout the remainder of the analysis; split ratio 40 : 1). MS conditions were as follows: ionization voltage 70 eV, acquisition mass range *m/z* 35-650, scan time 0.32 s. Percentage composition of the essential oils was computed from the GC peak areas without any corrections. Constituents were identified by comparison of their linear retention indices (relative to C₈-C₃₃ *n*-alkanes [7] on a DB-5MS column) with literature values [7] and their mass spectra with those of authentic standards, as well as those from Wiley 6, NIST11, MassFinder 2.3, and a homemade MS library with the spectra corresponding to pure substances and components of known oils, and wherever possible, by co-injection with an authentic sample.

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Vibrational and Thermal Studies of Essential Oils Derived from *Cistus ladanifer* and *Erica arborea* Shrubs

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Essential oils from the two most representative shrub species from the Iberian Peninsula (namely *Cistus ladanifer* L. and *Erica arborea* L.) have been characterized by Fourier transform infrared spectroscopy (FTIR) and thermoanalytical techniques (TG/DTG and DSC). Vibrational spectra have been compared with those of components of the plants, and with those of oils, gums and resins from other species. The different content in terpenoids of *C. ladanifer* oil (mainly mono- and sesquiterpenoids) and *E. arborea* oil (mainly triterpenoids) is reflected in the ATR-FTIR by the position of the bands at around 2873 cm⁻¹, 1730 cm⁻¹ and 1678 cm⁻¹. As regards their thermal behavior, *C. ladanifer*-derived oil evinced higher thermal stability than that of obtained from *E. arborea*: the pyrolysis of the former was sensitized at 210°C, whereas for the latter it occurred at 143°C. These temperatures are high enough to state that thermolabile constituents such as terpenoids are conserved in the hydrodistillation and that this extraction process ensures the recovery of the main constituents of both essential oils.

Keywords: *Cistus ladanifer*, *Erica arborea*, Oils, TG/DTG, DSC, FTIR.

Pyrophytes are plants which have adapted to tolerate fire. "Passive pyrophytes" (e.g., *Asphodelus albus* Willd.) are able to resist the effects of fire, particularly when it passes over quickly, and hence can out-compete less resistant plants, which are damaged. "Active pyrophytes" (such as *Cistus ladanifer* L. and *Erica arborea* L.) have a similar competing advantage, but also contain volatile oils and hence encourage the incidence of fires which are beneficial to them. *C. ladanifer* (gum rockrose), native to the Western Mediterranean, is known in Spanish as 'Jara pringosa' (meaning 'sticky shrub') because the whole plant is covered with a sticky exudate of a fragrant resin. This resin, known as 'labdanum', 'ladanum', 'laudanum' or 'ladanon', is produced by boiling the leaves and twigs. It presents a relatively high content of ambrox (amber odor) and, consequently, it is valued in perfumery as a substitute for ambergris (from the sperm whale, which is an endangered species). *Cistus* essential oil is obtained by either hydrodistillation or steam distillation of the leaves and stems, reaching a price of over 200 €/L.

Cistus oil is relatively rich in monoterpenes (67%) and sesquiterpenes (20%) and poor in diterpenes (3%). The major constituents are α-pinene (a bronchodilator), *trans*-pinocarveol, bornyl acetate, terpinen-4-ol and camphene (monoterpenes); viridiflorol (estrogen mimic, also reported in propolis), ledol and globulol (sesquiterpene alcohols); and 15-nor-labdan-8-ol (diterpene) [1-3].

Erica arborea (tree heath) is in the heather family, *Ericaceae*. It is native to the maquis shrublands (garigue biome) of the Mediterranean Basin, Portugal and the Canary and Madeira Islands. The leaves and flowers of *E. arborea* have been popularly used as an anti-rheumatic, a diuretic, an astringent and in the treatment of urinary infections, while its wood (briar root or *bruyère* in French) is used for making smoking pipes and jewelry.

E. arborea contains many active compounds such as flavonoids, monoterpenes, triterpenoids, phenylpropanoid glucosides and condensed tannins [4]. Pharmacological activities of the extracts from this plant have been reported to be anti-inflammatory, antioxidant and analgesic. In fact, the composition of its bark (with leaves) has been reported to include triterpenoids such as friedelin, lupeol, betulin and ursolic acid [5, 6].

According to Bessah and Benyoussef [4], the water-distilled essential oil from leaves of *E. arborea* of Algerian origin ($d=0.8587$; $[\alpha] +2^{\circ}44'$) contains 75 components, amongst which palmitic acid (33.3%), (Z,Z,Z)-9,12,15-octadecatrien-1-ol, a fatty alcohol (6.6%), and nonacosane, a straight-chain hydrocarbon (6.1%), are the major constituents [4]. Other components, present in lower proportions, are β-fenchyl alcohol, β-caryophyllene, β-bourbonene, eugenol, ionol, geranylacetone and germacrene D [4].

To the best of the authors' knowledge, no vibrational or thermal studies have been reported in the literature for any of the two aforementioned oils. The thermal behavior of *C. ladanifer* has been studied by TG and triple shot pyrolysis [7] of the entire plant, and the studies on *E. arborea* have been aimed at the assessment of the thermal behavior of the entire plant for its integration in wildland fire spread models [8] or to measure its ignitability, combustibility and sustainability as a forest fuel [9].

In the work presented herein, the physicochemical properties of the essential oils obtained from these two pyrophytic Mediterranean shrubs have been investigated by ATR-FTIR vibrational spectroscopy and by thermogravimetric (TG) and differential scanning calorimetric (DSC) thermoanalytical techniques. These data are of interest in extraction processes for the cosmetics industry, provided that they reveal differentiation patterns of the vegetable oil fingerprints.

ATR-FTIR spectra: The wavenumbers for the main bands in the ATR-FTIR spectra of the oils of the two plants under study, together with those from leaves, roots, capsules, fine and coarse-texture components for each of the species, are summarized in Table 1. A close similarity can be observed, on the one hand, between the leaves and roots spectra and, on the other hand, between the fine and coarse components spectra. The capsules vibrational pattern for *E. arborea* exactly matched that of the fine components and is listed in the same column.

Table 1: Main bands in the ATR-FTIR vibrational spectra of *C. ladanifer* and *E. arborea* oils and various plant components. All wavenumber values are in cm^{-1} .

<i>Cistus ladanifer</i>					<i>Erica arborea</i>				
Oil	Leaves	Roots	Fine	Coarse	Oil	Leaves	Roots	Capsules / Fine	Coarse
3465	3412	3416	3419	3421	3360	3396	3405	3421	3412
2953	x	x	x	x	2970	x	x	x	x
2918	2919	2920	2922	2920	2931	2923	2929	2918	2919
2871	2850	x	x	x	2874	2852	x	2850	
x	2360	2361	2355	2361	x	2360	2361	2360	2360
x	#	2344	2339	2343	x	#	2344	2342	2343
1737	1733	1735	1732	1735	1721	1733	1736	1739	1740
1681	x	x	x	x	1674	x	x	x	x
x	1615	1617	1621	1617	1614	1616	1617	1616	1616
x	1515	1508	1505	1508	x	1516	1513	1508	1510
1448	1455	1452	1455	1457	1454	1455	1451	1464	1459
1375	1367	1375	1373	1375	1374	1373	1374	1375	1375
1328	#	1320	1320	1320	1329	#		1331	
1245	1232	1243	1242	1243	1249	1243	1247	1246	1247
1164	1164				1152		1151		
1125									
1111					1085		1081		
x	1035	1035	1031	1035	1046	1035	1031	1050	1039
1015					1007				
952					939				
886					860				
815					821				
787					758				
635					665				
618	617	559	602	610	559		579	608	607

The oils spectra differed from those of the leaves, roots and other plant components from which they were obtained by the absence of the absorption bands at 2360 cm^{-1} (absorbed CO₂), 2344 cm^{-1} (glucose ring stretching), 1515 cm^{-1} (benzene ring stretching in lignin) and 1035 cm^{-1} (lignin), present in the latter components. Nonetheless,

they all shared the presence of bands at 1455 cm^{-1} (CH_2 cellulose, lignin) and 1375 cm^{-1} (C-H cellulose, hemicellulose) [10]. Bands specific to the oils are those which appeared at 2960 , 2873 , 1678 , 945 and 818 cm^{-1} . The bands at 2960 and 2873 cm^{-1} are assigned to asymmetrically and symmetrically stretching vibration of C-H of aliphatic CH_3 groups, due to the alkyl rest of the triglycerides present in large quantities in vegetable oils, while the band at around 1678 cm^{-1} can either be attributed to C=C stretching vibration of *cis* disubstituted olefins (RHC=CHR) or to an oxo group ($\alpha;\beta$ unsaturated) from terpenoids. The C-H out-of-plane deformation band observed between 952 and 939 cm^{-1} is highly characteristic of *trans* double bonds, and the band at around 818 cm^{-1} is related to an isopropylidene group ($\text{R}_2\text{C=CHR}$), usual in terpenoids [11-14].

The fact that the labdanum and tree heath oils spectra (Figure 1), as is the case for other oils, showed bands in common with resins and gums has led us to compare these spectra with those of rosehip and palm oils, myrrh and mastic resins and tragacanth gum (see Table 2). It is worth noting that, in the $2920-1160\text{ cm}^{-1}$ region, there is a high correspondence with the bands of *R. rubiginosa* oil, a moderate correspondence with those of other oils and resins, and low agreement with those of tragacanth gum.

In the case of *C. ladanifer* oil, whose terpene composition is favorable to low-number isoprenic units (monoterpene and sesquiterpenes), the band at 1737 cm^{-1} (attributable to $\nu(\text{CO})$ ester carbonyl or to terpenic oxo groups) is shared with rosehip and palm oils, whereas the composition rich in terpenes with a high number of isoprenic units (triterpenoids) exhibited by *E. arborea* oil leads to a shift of this band towards lower wavenumbers (1721 cm^{-1}). As regards the band at around 2875 cm^{-1} (attributed to symmetrical $\nu(\text{C-H})$ from CH_3) in the *E. arborea* spectrum, it should be noted that it agrees with that of mastic resin. Another useful band in terms of differential identification of the oils is that which appears at 1681 cm^{-1} in *C. ladanifer* oil and shifted to lower wavenumbers in tree heath oil (1674 cm^{-1}) and rosehip oil (1653 cm^{-1}). This band does not appear in palm pulp oil or in tragacanth gum.

Table 2: Comparison of the vibrational spectra of *C. ladanifer* and *E. arborea* oils with rosehip and palm oils, myrrh and mastic resins and tragacanth gum.

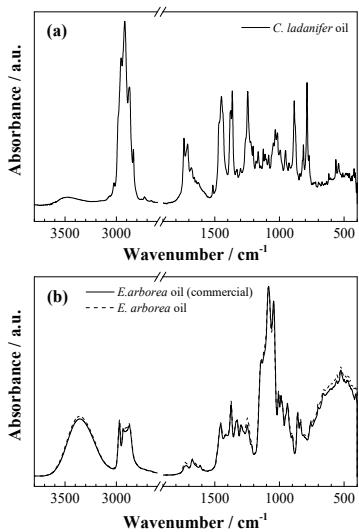


Figure 1: FTIR spectra of (a) *Cistus ladanifer* oil and (b) *Erica arborea* oil.

Thermal analysis: TG plots of gum rockrose and tree heath oils showed mass loss in the 50–240°C and 100–220°C temperature range, respectively, with a more abrupt pattern for *E. arborea* (Figure 2). In both cases this mass loss corresponded to pyrolysis, which was sensitized in the DTG curves by endotherms at 151°C (*C. ladanifer*) and at 208°C (*C. arborea*). From these temperatures, it can be seen that *E. arborea* oil presents higher thermal stability than that of *C. ladanifer*. In any case, the decomposition point of both oils is high enough to ensure the preservation of terpenoids as main constituents. The chief products of pyrolysis were straight-chain alkanes and alkenes [16]. It should be noted that heating of isopropanol extracts (not shown) resulted, after solvent evaporation, in decomposition of the oils extending up to 400 °C.

The low-temperature DSC thermograms of the two oils under study are depicted in Figure 3. Both *C. ladanifer* and *E. arborea* oils vitrify upon cooling. Upon heating, only devitrification is observed. For *C. ladanifer* oil the glass transition is observed at $T_g = (-113 \pm 1)$ °C and for *E. arborea* oil at $T_g = (-79 \pm 1)$ °C.

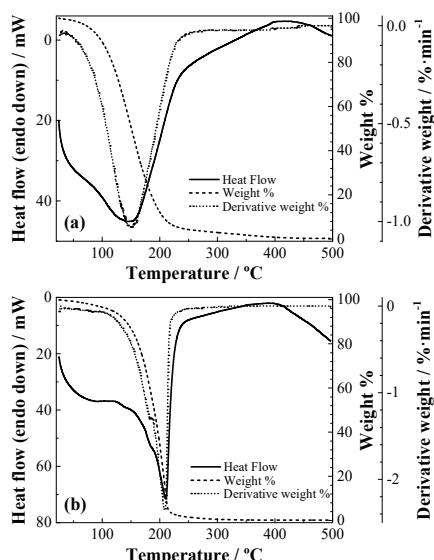


Figure 2: Thermograms for (a) *C. ladanifer* oil, $m = 21.14$ mg; and (b) *E. arborea* oil, $m = 17.77$ mg. The left y axis corresponds to the DSC curve, while the two y axes on the right correspond to TG/DTG curves.

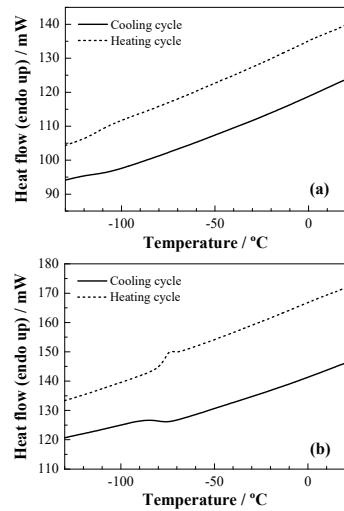


Figure 3: Low-temperature DSC thermograms for (a) *C. ladanifer* oil, $m = 4.1$ mg; and (b) *E. arborea* oil, $m = 5.7$ mg. The cooling/heating rate was $|\beta| = 50\text{°C}\cdot\text{min}^{-1}$.

As regards the FTIR results, this technique has proven useful to identify the different components of various parts of the plants and their derived products, serving as a fingerprint technique which provides insight into the biochemical composition of the samples. As noted by Huck [17], this method is particularly suitable for the fast and simultaneous qualitative and quantitative characterization of natural products and their constituents. Thus, the data presented herein can be used for quality control in the cosmetics industry.

With regard to the thermal stability of the oils under study, it is high enough to ensure that their thermolabile constituents would be preserved during hydro-distillation, which is the method recommended by the French Pharmacopoeia for the extraction of essential oils from dried spices and the quality control of essential oils in the laboratory [18].

On another topic, provided that the two shrub species under study are active pyrophytes, the results should also be put in relation to their impact on forest fires: the flammability of the shrubs under study (and, by extension, those of their resins and oils) must be referred (apart from ignition time and moisture) to their contents of terpenoids. The high flammability of *E. arborea* can be ascribed to its high emission of terpenes throughout all the year (which can reach up to $40\text{ }\mu\text{g}\cdot\text{g}_{\text{DM}}^{-1}\cdot\text{h}^{-1}$), in spite of the fact that its composition is relatively rich in low-volatile terpenoids, such as triterpenoids. Conversely, although *C. ladanifer* emits into the air modest amounts of terpenoids ($<7\text{ }\mu\text{g}\cdot\text{g}_{\text{DM}}^{-1}\cdot\text{h}^{-1}$), it has a very high flammability in the summer due to the potentiated emission of α -pinene, a particularly volatile monoterpene, which, as noted above, is an important component of its essential oil [19].

The essential oils from two Western Mediterranean pyrophytes, namely *C. ladanifer* and *E. arborea*, have been characterized by ATR-FTIR spectroscopy and thermal analytical (TG/DTG and DSC) techniques. Their vibrational spectra have been compared with those of other parts of the plants (leaves, roots, capsules, etc.) and with those of oils, gums and resins from other species. The specific location of the bands from unsaturated and $\alpha:\beta$ unsaturated oxogroups has been related to the different content of terpenoids of *C. ladanifer* oil (mono- and sesquiterpenoids) and *E. arborea* (triterpenoids). As regards the thermal behavior, *E. arborea* oil showed higher thermal stability than that of *C. ladanifer*, as evinced by the effects in the TG and DSC thermograms (at 210°C and

143°C, respectively). Thus, recognition of the TG and DSC characteristic patterns of both essential oils can also be helpful in identifying the type of oil. On the other hand, the delayed thermal decomposition of the oils under study, together with the characterization of terpenoids by ATR-FTIR, provide evidence that oil constituents that are thermolabile, such as terpenoids, are conserved in the extraction process, thus ensuring that hydrodistillation is a valuable method to recover the main constituents of these essential oils.

Experimental

Samples: *C. ladanifer* and *E. arborea* samples were collected from wild plants growing in the municipality of Ayoó de Vidriales (in the province of Zamora, Castilla y Leon, Spain) during the flowering period (Spring 2015). Oils from *Rosa rubiginosa* L. and *Elaeis guineensis* Jacq., myrrh from *Commiphora*, mastic from *Pistacia lentiscus* L. and tragacanth gum from *Astragalus* samples used for comparison purposes were of commercial origin.

Methods: Hydro-distillation of 100 g of fresh and whole leaves of *C. ladanifer* and *E. arborea* was carried out in a Clevenger-type apparatus for 3 h. The essential oils were collected, dried under anhydrous sodium sulfate and stored at 4°C until used. To confirm their purity, essential oil composition was determined by gas chromatography coupled to mass spectrometric (GC-MS) analysis, according to the experimental conditions specified by Bessah and Benyoussef [4]. A commercial sample *E. arborea* essential oil, supplied by Radhe Shyam (Barcelona, Spain), was also tested. An alternative extraction method consisting of suspending the vegetal dry matter in amyl alcohol in a 1:2 (w/v) ratio for 30 min, under constant

shaking, at room temperature, was also used to isolate high-purity oil from both plants.

Apparatus: GC/MS analysis was conducted with an Agilent Technologies 7890A apparatus (Santa Clara, CA, USA). The vibrational spectra of the materials in the 400-4000 cm⁻¹ spectral range were measured using a Thermo Scientific (Waltham, MA, USA) Nicolet iS50 FT-IR spectrometer, equipped with an in-built diamond attenuated total reflection (ATR) system. Spectra of the oils were recorded at room temperature with a 1 cm⁻¹ spectral resolution, and 64 scans. TG and high-temperature DSC analyses were conducted with a Perkin-Elmer (Waltham, MA, USA) STA6000 simultaneous thermal analyser by heating the samples in a slow stream of N₂ (20 mL·min⁻¹) from room temperature up to 500°C, with a heating rate of 20°C·min⁻¹. Pyris v.11 software was used for data analysis. The low-temperature DSC experiments were performed in a Perkin-Elmer Pyris 1 power compensation calorimeter. A liquid nitrogen Cryofill cooling unit was used, and helium at a 20 mL·min⁻¹ flux was employed as the purge gas. Samples were cooled to -170°C and then heated to room temperature at a 50°C·min⁻¹ rate. The obtained data were analysed using TA Instruments Universal Analysis V4.1D software.

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Chemical Composition and Antibacterial Activity of the Essential Oil of *Phellodendron lavallei*

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The essential oils from inflorescences, fruits and leaves of *Phellodendron lavallei* Dode were analyzed by GC, GC/MS and ¹H NMR spectroscopy. About 100 compounds were identified. The principal components in the fruit oil were myrcene (47.7-52.0%) and limonene (38.4-40.9%), followed by germacrene D (1.2-3.2%) and (E)-β-caryophyllene (2.4-2.9%). The inflorescence oil contained mainly myrcene (44.8%), limonene (43.2%) and (E)-β-ocimene (4.1%). The main constituents of the leaf oil were limonene (26.7%) myrcene (22.1%), (E,E)-α-farnesene (5.6%) and α-pinene (5.0%). The antibacterial activity was examined towards both Gram-positive and Gram-negative bacteria. The fruit oil showed weak activity against the bacteria tested.

Keywords: *Phellodendron lavallei*, Essential oil composition, Myrcene, Limonene, Antibacterial activity.

Phellodendron lavallei Dode (Rutaceae) is a medium sized, deciduous tree with thick, corky bark, leathery pinnate leaves, small yellowish-green flowers and pea-sized black, aromatic fruits. The plant is native to east Asia. In Europe and North America it is cultivated in parks and along the roads because of its attractive appearance, high resistance to frost, drought, insects and pollution [1a-c]. Bark of the tree is used in Eastern medicine, due to anti-inflammatory, antibacterial, antifungal, antipyretic and hypoglycemic activities. Its leaves are effective antihepatotoxic and antiviral agent. Only few papers have described biologically active substances of this species. Alkaloids (berberine, jatrorrhizine, magnoflorine, palmatine, phellodendrine) and limonoids (limonin, obacunone) were isolated from bark [2a-d]. In leaves, flavonoids (norcariside, phellatin, phellavin) were found [3a,b]. Seeds contained 16% of lipids, including fatty acids (lauric, myristic, isopalmitic, eicosatrienoic), phospholipids and sterols (ergosterol, stigmasterol, β-sitosterol) [4].

In the literature there is no data on the essential oil of this species, and for that reason we decided to study the chemical composition of the essential oil from different parts of *P. lavallei* and the antibacterial activity of its fruit oil. The yields and chemical composition of the oils from fruits, inflorescences and leaves of the plant are presented in Table 1. The fruits were rich in essential oil (1.0-1.2%). Inflorescences and leaves contained only negligible amounts of oil. About one hundred compounds were identified, representing 98-99% of the oils. The monoterpene hydrocarbons myrcene and limonene were the principal constituents of the oils.

The fruit oil contained mainly myrcene (47.7-52.0%) and limonene (38.4-40.9%), followed by germacrene D (1.2-3.2%) and (E)-β-caryophyllene (2.4-2.9%). Other components occurred in amounts less than 1%. The enantiomeric ratio of limonene was: (R)-(+)limonene (97.3%) : (S)-(+)limonene (2.7%). The content of essential oil increased during the fruit's development, however, the oil composition varied only slightly at the same time. Drying of

fruits had no influence on the yield and composition of the oil. The fruit oil of *P. lavallei* resembles that of *P. sachalinense* oil with high amounts of myrcene (50-54%) and limonene (37-40%) [5]. It differs from *P. amurense*, *P. chinense*, *P. japonicum* and *P. piritiforme* oils in that the prominent component is myrcene (62-86%), whereas limonene occurs in concentration of 0.6-2% [6a-d].

The main constituents of the inflorescence oil were myrcene (44.8%), limonene (43.2%) and (E)-β-ocimene (4.1%). The leaf oil contained mainly limonene (26.7%), myrcene (22.1%), (E,E)-α-farnesene (5.6%) and α-pinene (5.0%).

P. lavallei fruit oil inhibited the growth of all tested Gram-positive and Gram-negative standard strains at concentrations between 53.0 μL/mL – 67.0 μL/mL. The lowest values of MIC for *Enterococcus faecalis* ATCC 51299 and *E. faecium* ATCC 35667 were obtained – 54.0 μL/mL and 56 μL/mL, respectively. Slightly more resistant to the oil was *Staphylococcus aureus* ATCC 433000 with a MIC of 59.0 μL/mL. A similar MIC value (59.0 μL/mL) was found for one of the Gram-negative standard strains – *Escherichia coli* ATCC 25922. The oil possessed significantly lower antibacterial activity against other Gram-negative bacteria – *Klebsiella pneumoniae* ATCC 700603 and *Acinetobacter baumannii* ATCC 19606; the MIC values were 64.0 μL/mL and 67.0 μL/mL, respectively.

The oil of *P. lavallei* contains large amounts of terpene hydrocarbons, which have rather low antimicrobial activity [7]; this is why the oil showed weak activity against the bacteria tested.

This is the first report on the essential oil of *P. lavallei*. The fruits of this species, due to their high yield of essential oil and its high content of myrcene and limonene, can be used as a source of natural myrcene and (R)-(+)limonene.

Table 1: Chemical composition of the essential oils from fruits (A – fresh unripe, B – fresh ripe, C – air-dried ripe), inflorescences (D) and leaves (E) of *Phellodendron lavallei*.

Compound	RI	LRI	A	B	C	D	E	Identification methods
Hex-3-en-1-ol*	844	851					0.3	RI, MS
Hex-2-en-1-ol*	857	861					0.4	RI, MS
Heptanal	882	882				tr	0.2	RI, MS
α -Pinene	933	936	0.4	0.6	0.5	2.4	5.0	RI, MS
Camphepane	947	950	tr	tr	tr	0.1	0.1	RI, MS
Sabinene	969	973	0.1	0.1	0.1	0.1	0.1	RI, MS
β -Pinene	974	978	tr	tr	tr	0.1	0.1	RI, MS
Myrcene	985	987	52.0	49.0	47.7	44.8	22.1	RI, MS, $^1\text{H-NMR}$
Hex-3-enyl acetate	999	1002					0.5	RI, MS
α -Phellandrene	1001	1002	0.3	0.3	0.2	0.1	0.3	RI, MS
α -Terpinene	1008	1013				tr	tr	RI, MS
p -Cymene	1015	1015	tr	tr	tr	tr	tr	RI, MS
Limonene	1026	1025	38.4	40.9	40.0	43.2	26.7	RI, MS, $^1\text{H-NMR}$
(Z)- β -Ocimene	1028	1029	tr	tr	tr	tr	4.2	RI, MS
(E)- β -Ocimene	1041	1041	0.8	0.7	0.4	4.1	3.2	RI, MS
Nonanal	1075	1076	tr	tr	tr	tr	0.4	RI, MS
Terpinolene	1084	1082	tr	tr	tr	tr	tr	RI, MS
Linalool	1087	1086	0.5	tr	tr	0.2	0.5	RI, MS
Perillene	1092	1090	tr	tr	tr	tr	1.9	RI, MS
<i>allo</i> -Ocimene	1116	1113	0.3	0.1	0.1	1.3	1.0	RI, MS
<i>cis</i> -Limonene oxide	1125	1126			0.1			RI, MS
<i>trans</i> -Limonene oxide	1130	1130			tr			RI, MS
Ethyl benzoate	1151	1150					0.1	RI, MS
Cryptone	1160	1160	tr	tr	tr			RI, MS
Terpinen-4-ol	1166	1164	tr	tr	0.1	tr	0.1	RI, MS
Decan-2-one	1174	1176	tr	tr	tr			RI, MS
α -Terpineol	1178	1176	tr	tr	tr	tr	tr	RI, MS
β -Cyclocitral	1201	1195	tr	tr	tr	tr	0.1	RI, MS
<i>trans</i> -Carveol	1203	1200	tr	tr	tr			RI, MS
<i>cis</i> -Carveol	1210	1210	tr	tr	tr			RI, MS
Citronellol	1212	1213	tr	tr	tr	tr	tr	RI, MS
Carvone	1216	1214	tr	tr	tr			RI, MS
(Z)-Hex-3-enyl 2-methylbutanoate	1218	1229 ^a					0.1	RI, MS
(Z)-Hex-3-enyl 3-methylbutanoate	1221	1232 ^a					0.2	RI, MS
Geraniol	1238	1235	tr	tr	tr	0.1	0.1	RI, MS
Methyl citronellate	1244	1245	tr	tr	tr	tr		RI, MS
Methyl nerolate	1264	1265	tr	tr	tr			RI, MS
Bornyl acetate	1270	1270	tr	tr	tr			RI, MS
Undecan-2-one	1275	1273	0.2	0.4	0.8	0.1		RI, MS, $^1\text{H-NMR}$
Perilla alcohol	1280	1280	tr	tr	tr			RI, MS
Undecan-2-ol	1284	1284	tr	tr	0.1	tr		RI, MS
Methyl geranate	1305	1306	tr	tr	tr	tr		RI, MS
Citronellyl acetate	1334	1337	0.1	tr	0.1	0.1	0.1	RI, MS
Bicycloelemene	1337	1338	tr	tr	tr	tr	tr	RI, MS
δ -Elemene	1341	1341	0.2	0.1	0.2	tr	0.3	RI, MS, $^1\text{H-NMR}$
Neryl acetate	1344	1342	tr	tr	tr	tr		RI, MS
α -Cubebene	1355	1355	tr	tr	tr	tr	0.1	RI, MS
Geranyl acetate	1362	1362	0.5	0.2	0.2	0.4	0.1	RI, MS, $^1\text{H-NMR}$
N-Methyl methyl anthranilate	1372	1372				tr		RI, MS
α -Copaene	1382	1379	tr	0.1	0.1	tr	0.1	RI, MS, $^1\text{H-NMR}$
Methyl perillate	1384	1381	tr	tr	tr			RI, MS
α -Elemene	1386	1381	tr	tr	tr			RI, MS
β -Bourbonene	1388	1386	tr	tr	tr			RI, MS
β -Elemene	1392	1389	0.1	0.1	0.2	tr	tr	RI, MS, $^1\text{H-NMR}$
β -Cubebene	1394	1390	tr	tr	tr	tr	0.3	RI, MS
α -Gurjunene	1415	1413	tr	tr	tr			RI, MS
β -Ylangene	1421	1420	tr	tr	tr		0.3	RI, MS
β -Caryophyllene	1424	1421	2.8	2.4	2.9	0.4	1.1	RI, MS, $^1\text{H-NMR}$
β -Copaene	1435	1430	0.2	0.2	0.3	tr	0.2	RI, MS
β -Gurjunene	1437	1437	tr	tr	tr	tr	tr	RI, MS
Isogermacrene	1447	1446	0.1	tr	0.1	tr	0.2	RI, MS
(E)- β -Farnesene	1450	1446	tr	tr	tr	tr	0.1	RI, MS
ζ -Muurolene	1453	1455	tr	tr	0.1	tr	0.1	RI, MS
α -Humulene	1459	1455	0.2	0.1	0.2	tr	0.2	RI, MS, $^1\text{H-NMR}$
<i>cis</i> -Muurola-4(15),5-diene	1462	1462	tr	tr	0.1	tr	0.2	RI, MS
<i>allo</i> -Aromadendrene	1465	1463	tr	0.1	tr	tr		RI, MS
(E)- β -Ionone	1472	1468					0.1	RI, MS
γ -Muurolene	1476	1474	tr	tr	tr	tr	0.4	RI, MS
Tridecan-2-one	1478	1477	0.1	0.1	0.4	tr		RI, MS, $^1\text{H-NMR}$
Germane D	1483	1479	1.2	2.4	3.2	0.2	3.4	RI, MS, $^1\text{H-NMR}$
<i>epi</i> -Cubebol	1488	1490	tr	0.1	0.1	tr		RI, MS
Bicyclogermacrene	1492	1494	tr	0.2	0.2	0.1	0.1	RI, MS, $^1\text{H-NMR}$
α -Muurolene	1495	1496	tr	tr	0.1	tr	1.0	RI, MS
(E,E)- α -Farnesene	1499	1498	tr	tr	tr	tr	5.6	RI, MS
β -Himachalene	1503	1500					0.2	RI, MS
γ -Cadinene	1510	1507	tr	tr	tr	tr	0.9	RI, MS
Cubebol	1517	1514	tr	0.1	0.1	tr		RI, MS
δ -Cadinene	1522	1520	tr	0.2	0.4	0.2	2.6	RI, MS, $^1\text{H-NMR}$
Cadina-1,4-diene	1527	1523	tr	tr	tr	tr	0.1	RI, MS
α -Cadinene	1534	1530	tr	tr	tr	tr	0.3	RI, MS
β -Elemol	1545	1541	0.1	tr	tr	tr		RI, MS
Germane B	1550	1552	tr	tr	tr	tr		RI, MS
(E)-Nerolidol	1553	1553	0.2	tr	tr	0.1	1.2	RI, MS, $^1\text{H-NMR}$
Germacrene D-4-ol	1573	1571	0.2	0.3	0.3	0.1	1.7	RI, MS, $^1\text{H-NMR}$
Spathulenol	1576	1572	tr	0.1	0.1	tr	tr	RI, MS, $^1\text{H-NMR}$

Table 1 (continued)

Compound	RI	LRI	A	B	C	D	E	Identification methods
Caryophyllene oxide	1580	1578	0.1	tr	tr	0.1	0.2	RI, MS, ¹ H-NMR
Globulol	1584	1589	tr	tr	tr	tr	tr	RI, MS
Salvia-4(14)-en-1-one	1592	1592	tr	tr	tr	tr	tr	RI, MS
Viridiflorol	1596	1592				tr	0.1	RI, MS
1,10-di- <i>epi</i> -Cubenol	1617	1615	tr	tr	tr	tr	tr	RI, MS
1- <i>epi</i> -Cubenol	1625	1623	tr	tr	tr	tr	0.1	RI, MS
<i>epi</i> - α -Murolool	1633	1633	tr	0.1	0.1	0.3	1.7	RI, MS, ¹ H-NMR
<i>epi</i> - α -Cadinol	1635	1633	tr	tr	tr	tr	0.1	RI, MS
α -Cadinol	1645	1643	tr	0.2	0.2	0.5	2.7	RI, MS, ¹ H-NMR
(E,E)-Farnesol	1717	1718	0.4	0.1	0.1	0.2	0.5	RI, MS, ¹ H-NMR
(E,E)-Farnesyl acetate	1840	1843 ^a	0.1	0.1	0.1	tr	1.1	RI, MS, ¹ H-NMR
Cembrene	1941	1938	tr	tr	tr	tr	tr	RI, MS
Isophytol	1944	1949					0.1	RI, MS
Phytol	2108	2114					3.2	RI, MS
Aliphatic hydrocarbons C ₁₉ -C ₂₇						tr	0.3	RI, MS
Total		99.6	99.3	99.9	99.3	99.3	98.8	
Monoterpene hydrocarbons		92.3	91.7	89.0	96.2	96.2	62.8	
Oxygenated monoterpenes		1.1	0.2	0.5	0.8	0.8	2.9	
Sesquiterpene hydrocarbons		4.8	5.9	8.1	0.9	0.9	17.8	
Oxygenated sesquiterpenes		1.1	1.0	1.0	1.3	1.3	9.4	
Aliphatic ketones		0.3	0.5	1.2	0.1	0.1		
Oil yield in fresh source (v/w, average of several replicates)		1.0	1.2	2.6	0.06	0.06	0.01	
Oil yield in dry weight (v/w)		2.7	3.0	2.9	0.2	0.2	0.04	

RI – retention indices on Rtx-1 column; LRI – retention indices of literature on DB-1 column according to MassFinder 3.1; ^a – retention indices of literature on DB-5 column [8]; tr – trace (<0.05%); * – correct isomeric form not identified.

Experimental

Plant material: *Phellodendron lavallei* Dode was collected in the Forest Experimental Station, Arboretum of Warsaw Agriculture University in Rogow, Poland. Inflorescences were harvested at the beginning of June, leaves and unripe fruits in July, and ripe fruits in October 2013. Voucher specimens (No. 2013/Pla1-2013/Pla4) were deposited in the Herbarium of the Institute of General Food Chemistry, Lodz University of Technology.

The fresh inflorescences (moisture 75.0%) and fresh leaves (moisture 72.8%) were cut into small pieces, and fresh unripe fruits (moisture 61.3%), fresh ripe fruits (moisture 62.0%) and air-dried for a month ripe fruits (moisture 10.6%) were ground. Each sample of plant material (200 g) was separately hydrodistilled in a Clevenger-type apparatus for 3 h to obtain the essential oil. After decanting and drying over anhydrous MgSO₄, the oils were stored at low temperature (5°C) before analysis. The oils had a pale yellow color and intensive terpenic aroma.

Analysis of the essential oils: The chemical composition of the oils was determined by chromatographic and spectroscopic methods.

The gas chromatographic (GC) analyses were carried out on a Carlo Erba Instruments HRGC 5300 gas chromatograph operated with a split/splitless injector, a flame ionization detector using an apolar capillary column Rtx-1 (dimethylpolysiloxane), 60 m × 0.25 mm × 0.25 μm film thickness (Restek, Bellafonte, PA, USA), temperature program 50–300°C at 4°C/min, injector temperature 320°C, detector temperature 310°C. Injection volume was 0.08 μL, split ratio 1:13, carrier gas helium with a flow 4 mL/min.

The fruit oil was also analyzed by GC with a capillary column Rtx-β-Dexst, 30 m × 0.32 mm × 0.25 μm film thickness (Restek, Bellafonte, PA, USA), temperature program 50–220°C at 4°C/min.

The gas chromatography/mass spectrometry (GC/MS) analyses were performed on a Thermo Electron Corporation Trace GC Ultra gas chromatograph equipped with a mass detector DSQ II. The MS operating parameters were: ionization voltage 70 eV, ion source temperature 200°C, mass range 33–420 m/z. Other conditions of the analyses were the same as described under GC analyses.

¹H NMR spectra were obtained on a Bruker DPX 250 Avance spectrophotometer using CDCl₃ as solvent and TMS as a standard.

Additionally, the fruit oil (15.0 g) was vacuum distilled in a Vigreux column to obtain fraction 1 (12.6 g) containing monoterpene hydrocarbons and fraction 2 (1.8 g, residue). Fraction 2 was separated by flash chromatography (FC) on Kieselgel 60, particle size 0.040–0.063 mm (Merck) with n-hexane and increasing amounts of diethyl ether to obtain fractions 2.1–2.6. Fraction 2.1 (835 mg), containing sesquiterpene hydrocarbons and fractions: 2.2 (310 mg), 2.3 (170 mg), 2.4 (190 mg), 2.5 (170 mg) and 2.6 (150 mg), containing oxygenated compounds, were separated on Kieselgel 60 impregnated with AgNO₃ to obtain fractions 2.1.1–2.1.6, 2.2.1–2.2.4, 2.3.1–2.3.4, 2.4.1–2.4.3, 2.5.1–2.5.4 and 2.6.1–2.6.4. All fractions were analyzed by GC/MS, additionally fractions: 1, 2.1.1–2.1.6, 2.2.2, 2.2.3, 2.3.2, 2.3.3, 2.4.2, 2.5.2, 2.5.3, 2.6.2 and 2.6.3 by ¹H NMR spectroscopy. Fraction 1 contained myrcene (57%) and limonene (42%), fraction 2.1.1 (65 mg) δ-cadinene (63%) and α-copaene (18%), fraction 2.1.2 (390 mg) (E)-β-caryophyllene (54%) and germacrene D (38%), fraction 2.1.3 (220 mg) germacrene D (46%) and (E)-β-caryophyllene (45%), fraction 2.1.4 (25 mg) β-elemene (56%), fraction 2.1.5 (58 mg) bicyclogermacrene (24%) and δ-elemene (22%), fraction 2.1.6 (33 mg) α-humulene (63%), fraction 2.2.2 (235 mg) undecan-2-one (37%), tridecan-2-one (27%) and geranyl acetate (24%), fraction 2.2.3 (35 mg) (E,E)-farnesyl acetate (36%), fraction 2.3.2 (52 mg) undecan-2-one (55%) and caryophyllene oxide (21%), fraction 2.3.3 (75 mg) germacrene D-4-ol (47%), fraction 2.4.2 (84 mg) (E)-nerolidol (53%), fraction 2.5.2 (102 mg) *epi*-α-murolol (68%), fraction 2.5.3 (40 mg) spathulenol (25%), fraction 2.6.2 (70 mg) α-cadinol (62%) and fraction 2.6.3 (65 mg) (E,E)-farnesol (83%).

Identification of components of the oils and components of the fractions was based on comparison of their retention indices relative to n-alkanes (C₈-C₂₆) and their MS with those of commercial libraries (NIST 98.1, Wiley Registry of Mass Spectral Data 8th edn, MassFinder 3.1) and literature [8]. Major components of the oil, separated by FC, were additionally identified by comparison of their ¹H NMR spectra with literature data [9a-j].

A quantitative analysis (expressed as percentage of each component) was carried out by peak area normalization measurements without correction factors.

Antibacterial activity of the essential oil: The Gram-positive standard strains: *Staphylococcus aureus* ATCC 433000, *Enterococcus faecalis* ATCC 51299, *E. faecium* ATCC 35667, and Gram-negative: *Escherichia coli* ATCC 25922, *Klebsiella pneumoniae* ATCC 700603, and *Acinetobacter baumannii* ATCC 19606 came from the collection of the Environmental Biology Department, Medical University of Lodz. The bacteria were cultivated on Columbia Agar (bioMerieux) at 37°C for 24 h under aerobic conditions. The suspensions of bacterial standard strains with an optical density of 0.5MF scale were prepared with a bioMerieux densitometer.

The antibacterial activity of the essential oil from fruits of *P. lavallei* was investigated by the micro-dilution broth method. The essential oil was diluted in ethanol and used as a stock solution. This stock solution was mixed with a 100 µL Mueller-Hinton Broth (Graso, Poland) to obtain concentrations from 52.0 µL/mL to 68.0 µL/mL. An inoculum containing $1.5 \cdot 10^8$ CFU (10 µL) per well was added to broth with various essential oil concentrations; as well as to broth with no oil added containing only ethanol at concentrations used in the dilutions of oil and were transferred to 96-well microtiter plates. The Minimal Inhibitory Concentration (MIC) was determined as the lowest concentration of oil which inhibits the visible growth of bacteria after 24 h of incubation at 37°C under aerobic conditions.

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Antimicrobial Potential of *Rosmarinus officinalis* Commercial Essential Oil in the Treatment of Vaginal Infections in Pregnant Women

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Treating inflammatory conditions such as vaginosis, vaginitis, and vulvovaginitis in pregnancy is a special problem due to limitations of available drugs. However, possible treatment options can be found also in naturally originated products, such as essential oils (EOs) of different plants. The aim of this study was to evaluate *in vitro* antimicrobial and toxic activities of the commercial EO of *Rosmarinus officinalis* L. (*Lamiaceae*) against five Gram-positive and Gram-negative bacterial strains and two *Candida* strains obtained from pregnant women with vaginal infection. Gas chromatography-mass spectrometry of the tested EO revealed oxygenated monoterpenes to be the major ingredients, while microdilution assay showed the highest activity on *Staphylococcus aureus* II strain at 6.2 mg/mL. After 24 hours, toxicity was determined at 19.4 mg/mL on *Artemia salina* nauplii. The obtained results show this EO to be a promising alternative therapy for vaginal infections, although further toxicity and safety research is required.

Keywords: *Rosmarinus officinalis*, Essential oil, Pregnancy, Inflammation, Antimicrobial activity, Toxicity.

Species of *Lamiaceae* are the most known sources of essential oils (EOs), among which the EO of *Rosmarinus officinalis* L. (commonly known as rosemary) is already well known for its antimicrobial activity. While carnosic and rosmarinic acid may be the main bioactive antimicrobial compounds present in rosemary extracts [1a], antibacterial efficacy of rosemary essential oil is associated with the presence of compounds such as α-pinene, camphor and (-)-bornyl acetate [1b]. Furthermore, this EO expresses additional beneficial properties, such as improving circulation and relieving pain [1c], as well as having a potential anticarcinogenic activity [1d]. There is a rising need to investigate plants used in alternative therapies to determine if their use is based on the principles of evidence-based medicine [1e]. Novel natural agents will be useful in the upcoming post-antibiotic era, with the evident need for isolation of new compounds from natural products to which antimicrobial resistance has not been detected [1a].

The genital inflammatory conditions, vaginosis, vaginitis, and vulvovaginitis, occur in more than 30% of the population of women in their reproductive years [1f]. Vaginosis is a genital inflammatory condition that occurs in women of all ages [1g] and is characterized by dysbiosis of the vaginal microbiome [1h]. The etiology of vaginosis is still not fully understood but it is most likely a result of changes in the vaginal pH, caused by substitution of the peroxide-producing *Lactobacillus* species with infective agents, both aerobic and anaerobic bacteria, and yeasts [1i]. One of the major problems with treating these conditions in pregnancy is that pregnancy itself limits the number of drugs that can be used, because of their potential teratogenic effect. The most commonly used antibiotics, such as penicillin-based drugs and metronidazole, as well as many antifungals, are increasingly less effective due to the increase of antimicrobial resistance to infectious agents [1e,1j]. However,

possible treatment options can be found in various plant products, such as EOs, extracts, and tinctures. Some traditional medical practices have used such products because of their healing properties for centuries, but it is only recently that we have started to determine, isolate and understand the mechanism of action of active compounds in these products. By knowing which molecule or, even better, what specific combination of molecules, contributes the most to the antimicrobial effect of certain natural products, we can test and recommend their use in accordance with the principles of evidence-based medicine [1k]. The aim of this research was to evaluate *in vitro* possible antimicrobial and toxic activities of the commercial EO of *R. officinalis* against bacterial and *Candida* strains isolated from pregnant women with vaginal infection and to assess its possible application as an alternative treatment for inflammatory gynecological diseases.

The chemical composition of the EO determined with GC/MS is given in Table 1. Oxygenated monoterpenes were the major compounds. This finding is in agreement with the study of Hussain *et al.* [2a] in which the tested *R. officinalis* EO from Pakistan contained a high concentration of oxygenated monoterpenes (61.7%). Similarly, Teixeira *et al.* [2b] reported that the rosemary EO they analyzed contained 59.7% of oxygenated monoterpenes, which is also consistent with our findings. The major components were eucalyptol (39.0% GC/MS; 42.2% GC/FID), camphor (13.7% GC/MS; 11.6% GC/FID), α-pinene (10.3% GC/MS; 12.1% GC/FID), (1S)-(-)-β-pinene (7.2% GC/MS; 8.2% GC/FID) and endo-borneol (4.7% GC/MS; 4.3% GC/FID). Similarly, several published papers reported that the major components of *R. officinalis* EO were eucalyptol (29.2%), camphor (17.2%) and α-pinene (11.5%) [2c-2e]. The variations in the main constituent (eucalyptol and camphor) concentrations could be explained by

Table 1: Chemical composition of the rosemary essential oil (GC/MS analysis).

No of Peak	^a Rt (mins)	^b Chemical constituents	^c RI	Composition (Area %; Relative %)	Formulas	MW
1	12.6	Tricyclene	933	0.8	C ₁₀ H ₁₆	136.1
2	12.9	α-Pinene	944	10.3	C ₁₀ H ₁₆	136.1
3	13.4	Camphene	963	4.0	C ₁₀ H ₁₆	136.1
4	14.1	(1S)-(-)-β-Pinene	989	7.2	C ₁₀ H ₁₆	136.1
5	14.2	β-Myrcene	992	1.1	C ₁₀ H ₁₆	136.1
6	14.8	Pseudolimonene	1013	0.1	C ₁₀ H ₁₆	136.1
7	14.9	(+)-3-Carene	1018	0.2	C ₁₀ H ₁₆	136.1
8	15.3	p-Cymene	1035	1.9	C ₁₀ H ₁₆	136.1
9	15.4	(+)-Limonene	1040	3.5	C ₁₀ H ₁₆	136.1
10	15.6	Eucalyptol	1047	39.0	C ₁₀ H ₁₈ O	154.1
11	17.0	Linalool	1104	1.5	C ₁₀ H ₁₈ O	154.1
12	18.2	trans-Sabinol	1161	0.2	C ₁₀ H ₁₈ O	152.1
13	18.3	(-)Camphor	1168	13.7	C ₁₀ H ₁₀ O	152.1
14	18.6	Pinocarvone	1181	0.3	C ₁₀ H ₁₈ O	150.1
15	18.8	endo-Borneol*	1191	4.7	C ₁₀ H ₁₈ O	154.1
16	18.9	Terpinen-4-ol	1195	1.8	C ₁₀ H ₁₈ O	154.1
17	19.2	α-Terpineol	1210	3.1	C ₁₀ H ₁₈ O	154.1
18	19.3	(1R)-(-)-Myrtenal	1215	0.6	C ₁₀ H ₁₈ O	150.1
19	19.6	Fenchyl acetate	1231	0.3	C ₁₂ H ₂₀ O ₂	196.1
20	20.2	(+)-Carvone	1267	0.1	C ₁₀ H ₁₄ O	150.1
		Acetic acid, 1,7,7-trimethylbicyclo[2.2.1]hept-2-yl ester				
21	20.8		1298	1.6	C ₁₂ H ₂₀ O ₂	196.1
22	23.2	Caryophyllene	1449	3.8	C ₁₅ H ₂₄	204.2
23	23.9	Humulene	1486	0.3	C ₁₅ H ₂₄	204.2

^aRetention time. Compounds listed in order of elution; ^bChemical constituents measured on the non-polar capillary column (HP-5MS, 60 m fused silica) and GC-MS detection; ^cRI (retention indices) measured relative to *n*-alkanes (C9–C18) on the non-polar column (HP-5 column) and GC-MS detection.

* endo-Borneol is (-)-Borneol

environmental and local climatic conditions, the time of harvest, the stage of plant development, or the method of extraction [2f,2g].

The chemical composition of EOs is crucial for their bioactivities [3a]. Objeda-Sana *et al.* [3b] tested individual major components of rosemary EO against Gram-positive and Gram-negative bacteria. They noticed that 1,8-cineole (eucalyptol) and α-pinene were active against Gram-negative bacteria; camphor and borneol inhibited only Gram-positive bacteria; while myrcene had no inhibitory activity. Klančík *et al.* [3c] investigated pure rosmarinic acid against Gram-positive bacteria and found it much less effective than crude EO. Our examined rosemary EO exhibited a higher and selective antibacterial activity against the Gram-positive bacteria (Table 2), with the MIC and MBC values ranging from 6.2 to 50 mg/mL. The strongest antibacterial activity of the EO was obtained against *S. aureus* II, with MIC/MBC values of 6.2 mg/mL, while its resistance to common antibiotics was higher than that of *S. aureus* I.

In another study, Hammer *et al.* [3a] determined that rosemary EO possesses a considerable antimicrobial activity against *E. coli*, *P. aeruginosa*, and *S. aureus*, while our EO showed significant inhibition of growth of *S. aureus* and only a modest activity against *E. coli*. In our research, we did not detect any antimicrobial effect of rosemary EO on the *P. aeruginosa* isolate, whereas MIC/MBC activity was recorded against *P. mirabilis* (25 mg/mL). This difference compared with the latter study may be due to different origins of the isolates. Hammer *et al.* [3a] used attested isolates, while we used clinical isolates. According to some other studies, EOs exhibit a stronger activity to Gram-positive bacteria [2a,2b].

In our study the analyzed rosemary EO reached MIC/MBC values for the *S. aureus* I isolate of 25 mg/mL, while in other studies rosemary EOs achieved MIC/MBC against attested *S. aureus* at 0.3 mg/mL [2c], indicating that the clinical isolates used in our study had strong antimicrobial resistance. Furthermore, the fact that the isolated clinical strains of *E. coli* I and *E. coli* II were resistant to the commercial antibiotics used in our study but sensitive to the EO, which showed MIC and MBC/MFC values of 12.5 and 25 mg/mL,

respectively, is of particular significance. This may be due to synergistic effects of the mixture of various chemical compounds, or it may be supposed that the effectiveness of EOs depends not only on their chemical composition but also on other conditions, such as pH of broth medium and strain-specific characteristics of the microorganisms [3d].

Table 2: MIC and MBC (MFC) values of the EO and common antibiotics and antimycotics (mg/mL)

Bacterial strain	Essential oil*				Antibiotics*							
	<i>R. officinalis</i>		STR		AMP		TET		CXM			
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC		
<i>E. coli</i> I	25	25	nd**	nd	nd	nd	nd	nd	nd	nd		
<i>E. coli</i> II	12.5	12.5	nd	nd	nd	nd	nd	nd	0.5	nd		
<i>S. aureus</i> I	25	25	0.1	0.1	0.1	0.1	0.1	0.1	0.5	0.3		
<i>S. aureus</i> II	6.2	6.2	0.1	0.5	nd	nd	0.1	0.5	nd	nd		
<i>P. mirabilis</i>	25	25	0.1	0.5	0.1	0.3	0.1	0.5	0.3	nd		
<i>P. aeruginosa</i>	50	50	nd	nd	nd	nd	nd	nd	nd	nd		
Fungal strain												
	<i>R. officinalis</i>				Nystatin							
	MIC	MFC			MIC				MIC	MFC		
<i>C. albicans</i> I	5	5			0.1				0.1	0.1		
<i>C. albicans</i> II	1.2	1.2			0.1				0.1	0.1		

*values are expressed in mg/mL; **nd- not detected, Tested antibiotics: STR = Streptomycin; AMP = Ampicillin; TET = Tetracycline; CXM = Cefuroxime

Our rosemary EO also showed an antifungal activity against both *Candida* isolates (1.2 – 5 mg/mL). The lowest MIC/MFC values were at 1.2 mg/mL for *C. albicans* II (Table 2). The EO antifungal activity was weaker than that of the tested antifungal drugs, which is similar to recent data showing that rosemary EO expressed antimicrobial activity on *Candida* species at 1 mg/mL [2e]. Likewise, strain-specific effects were observed for the *Candida* isolates, similarly to the research of Pintore *et al.* [2h].

LC₅₀ values in the toxicity assay after 24 and 48 hours were 19.4 mg/mL and 5.4 mg/mL, respectively. Prolonged exposure to the EO (from 24 to 48 hours) resulted in increased brine shrimp mortality. There was a high linear correlation between the EO concentration and mortality after 24 ($r^2 = 0.9$) and 48 hours ($r^2 = 0.8$). These results indicate that increased EO concentrations (3.1 – 50 mg/mL) lead to an increase in EO toxicity (from 43% to 95% after 24-hour incubation and from 65% to 97% after 48-hour incubation). The brine shrimp bioassay is a quick and cheap test for establishing acute toxicity of EO. Parra *et al.* [3e] found a high correlation between *in vivo* and *in vitro* tests ($r = 0.8$, $p < 0.05$). Considering that EO toxicity increased with higher concentrations and a longer exposure time, it is necessary to be careful with EO dosage and application.

Our results are in agreement with the results of a previous study reporting eucalyptol and camphor to be the most prevalent constituents [3f]. However, the fact that they were the most prevalent does not necessarily mean that these compounds were responsible for the recorded antimicrobial activity. Actually, some studies have shown that carnosol, carnosic acid, and rosmarinic acid possess major antibacterial [3g], antioxidant and anti-inflammatory properties [3h 3i]. In the study of Jordan *et al.* [3j] carnosic acid was the major component credited for bacteriostatic effects. Hence, any study of EOs should take into account that each EO is specific and its composition depends on the plant it was derived from and on the type of extraction procedure applied [3a,3l].

Furthermore, the antimicrobial effect of EOs cannot be explained by a single mechanism of action or with a single active compound. It is considered that all the antiviral, antifungal and antimicrobial effects of EOs are due to synergistic effects of their numerous bioactive constituents [3d].

Promising results were seen in studies comparing the standard antimicrobial drug ciprofloxacin and rosemary oil [3e]. Moreover, some studies evaluating the synergistic effect of antibiotics and rosemary EO found that the combination of the EO with aminoglycoside antibiotics could achieve a better effect against multidrug-resistant *S. aureus* and *E. coli* [3k,3l] than antibiotics alone. Further research which would clarify the underlying mechanism of the synergy is therefore recommended.

Our results showed a significant *in vitro* antimicrobial activity of the *R. officinalis* EO on clinically isolated vaginal bacterial and *Candida* strains. We may conclude that the EO represents a potential alternative therapy for vaginal infections, when used in appropriate doses. Future research should include a wider range of bacteria and *Candida* strains, and explore the synergistic effects with common antimicrobial drugs, as well as cytotoxic activity of rosemary EO, preferably in *in vivo* studies.

Experimental

Essential oils and GC/MS analysis: The analyzed commercial EO of *R. officinalis* was purchased from MeLab (Belgrade, Serbia, serial number ER010514) and obtained by steam distillation of the leaves and buds and stored in a refrigerator at +4°C prior to the analysis. The chemical composition was analyzed using a gas chromatography-mass spectrometry (GC/MS) after preparation by using dilution of oil with dichloromethane (1:10, w/v) and the trials were performed on an Agilent capillary gas chromatograph coupled with a mass spectrometer (MSD) (model GC Agilent 7890A; MS 5975C). A non-polar capillary column (HP-5MS, fused silica) was used with these parameters: oven temperature was changed during a 30-min run time in the following procedure: from 50°C (for 1 min) to 100°C at 5°C/min and to 200°C at 9°C/min with the final temperature kept for 7.89 min; temperature of the injector was 250°C, with helium as carrier gas (constant flow mode, flow rate was 0.8 mL/min), split injection technique (50 : 1); ionization energy 70 eV, in the EI ionization mode; ion source temperature was 230°C; scan mass range of *m/z* 30–550 and interface line temperature 280°C. Identification of the EO components was made from their retention indexes (RI) and by comparing their mass spectra, using the Wiley Registry of Mass Spectral Data, 7th edition, NIST8 Mass Spectral Library. The other components were identified by comparison of their spectral data and RI with the literature data (our own laboratory database and mass spectra generated by the instrument software - NIST mass spectrometry and MSD ChemStation E.02.00.493), after which all identified components were confirmed by the AMDIS software program (version 2.66). The proportion of the components was obtained from the electronic integration of the MSD signals of specific ion representation.

Essential oil and GC/FID analysis: Samples of the same EO were analyzed by gas chromatography (GC) with the use of a flame ionization detector (model GC Hewlett Packard 6890; FID). Retention indices were determined with a mixture of *n*-alkanes (C9 - C18) under the same experimental conditions. The preparation of the EO for the GC/FID analysis involved dilution of dichloromethane (1:10, w/v), and the trials were performed on a Hewlett Packard capillary gas chromatograph coupled with FID non-polar capillary column (HP-5MS 60 m, 320 m ID, 0.2 µm film thickness, fused silica) used with these parameters: oven temperature was changed during the 35-min run time in the following procedure: from 50°C (for 1 min) to 100°C at 5°C/min and to 200°C at 9°C/min with the final temperature kept for 12.9 min; temperature of the injector was 250°C, with helium as the carrier gas (constant pressure mode; pressure was 13.8 psi), split

injection technique (10 : 1); detector temperature 300°C, hydrogen flow 40.0 mL/min, air flow 450.0 mL/min, constant makeup flow 45.0 mL/min. Validation (and relative composition in %) of identified components was performed by GC with FID (GC/FID) and compared with experimentally acquired retention indices.

Bacterial and Candida strains: Six bacterial isolates: *Escherichia coli* I, *E. coli* II, *Staphylococcus aureus* I, *S. aureus* II, *Pseudomonas aeruginosa*, *Proteus mirabilis* and two *Candida* isolates, *C. albicans* I and *C. albicans* II, were obtained from the Faculty of Medicine, Clinical Centre of Vojvodina, Department of Obstetrics and Gynecology and derived from pregnant women at their regular appointments. The institutional Ethical Board of the same institution approved the study protocol.

Antimicrobial assay: The antibacterial activity, minimum inhibitory concentrations (MICs) and minimum bactericidal/fungicidal concentrations (MBCs/MFCs) of the rosemary EO were evaluated by a double-microdilution method according to the standard CLSI procedure [3m]. The tested strains were obtained from overnight cultures, grown at 37°C on nutrient agar (Müller-Hinton, MHA, Torlak, Belgrade, Serbia) for bacteria, and on malt agar (MA, Torlak, Belgrade, Serbia) for yeasts. McFarland inocula of bacteria and yeasts were prepared in sterile saline solution; reaching the final number of approximately 1.5×10^6 CFU/mL for bacteria and 1.5×10^5 for yeasts. Müller Hinton broth (MHB, Torlak, Beograd, Serbia) and Sabouraud broth (SB, Torlak, Belgrade, Serbia) were used for the antimicrobial screening. A double-dilution test was performed in either MHB or SB with 1% dimethyl sulfoxide (DMSO) in a 96-well microtitre plate (Spektar, Čačak, Serbia). The final concentrations of the EO ranged from 1.2–50 mg/mL. After 24-h incubation at 37°C for the bacterial strains and 48 h for yeasts, MIC values were determined visually. MBCs/MFCs were confirmed after inoculation of MHA and MA plates with 100 µL of broth where turbidity was absent (MIC point). Nystatin (Hemofarm, Vršac, Serbia) and commercial fluconazole solution (Hemofarm, Vršac, Serbia) (2 mg/mL) were used as positive controls for yeasts (in final concentrations 0.03, 0.06, 0.13 mg/mL), while antibiotics: streptomycin, ampicillin, tetracycline, and cefuroxime (Himedia, Mumbai, India) were positive controls for bacteria (0.5 mg/mL). As a negative control, 1% DMSO was used. The test was performed in 3 replicates and the average value was taken as the final result.

Brine shrimp toxicity assay: Toxicity of the investigated rosemary EO was made by *Artemia salina* bioassay according to Meyer *et al.* [3n]. In each microwell of the plate, 10 larvae were added. The wells already contained 25 µL of specific EO dilution in 1% (DMSO) and 225 µL of artificial sea water (ASW). LC₅₀ values of EO were determined by serial double dilutions in 6 repetitions (3.1 – 50 mg/mL). A mixture of 25 µL 1% DMSO (oil emulsifier) and 225 µL ASW with larvae was used as a control. For toxin control, microcystin standard MCYST-LR (SigmaAldrich, Germany) was used. Microplates were incubated under a lamp, at 30°C, without feeding the shrimps. Acute toxicity was calculated after 24 and 48 h, by counting dead nauplii. Surviving shrimps were counted under a stereomicroscope (Zeiss, Germany) and afterwards, 50 µL of methanol was added to each well. All the tests were repeated twice and performed in triplicate. Abbott's equation was applied to calculate the mortality rate: Mortality (%) = ((*n* test – *n* control)/*n* control) x 100, where *n* control and *n* test were the numbers of dead larvae in the first test and control tests. Toxicity was expressed as LC₅₀ (mg/mL). LC₅₀ represented the lethal concentration required to kill 50% of the population of test brine shrimps, *Artemia salina*.

Statistical analyses: Statistical analyses of the data were performed by analysis of variance (ANOVA), using Microsoft Excel XP. Duncan's multiple range test was applied for the determination of significant differences between results ($p < 0.05$) in the antimicrobial testing using Analytical Software Statistica 8.1.

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Chemical Profile and *in vitro* Biological Activities of Essential Oils of *Nectandra puberula* and *N. cuspidata* from the Amazon

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Essential oils (EO) from leaves and branches of *Nectandra puberula* Schott (Nees) and from leaves of *N. cuspidata* Nees & Mart. were obtained by hydrodistillation and their chemical compositions determined by gas chromatography - mass spectrometry (GC-MS). The main compounds identified from *N. puberula* EO were apiole (22.2%), β -caryophyllene (15.1%), β -pinene (13.3%), germacrene D (8.3%), pogostol (6.6%) and bicyclogermacrene (6.4%) in the leaves; and apiole (28.1%), pogostol (19.8%) and guaiol (11.2%) in the branches. The EO of *N. cuspidata* leaves showed β -caryophyllene (26.9%), bicyclogermacrene (16.0%) and spathulenol (5.2%) as the main compounds. The EOs were subjected to antibacterial screening and displayed promising activity against *Escherichia coli* ($MIC = 19.5 \mu\text{g.mL}^{-1}$). In addition, the EOs were tested for cytotoxic activity against MCF-7 breast tumor cells and the IC_{50} values were 64.5 ± 1.6 and $117.1 \pm 11.9 \mu\text{g.mL}^{-1}$ for the leaf EOs of *N. puberula* and *N. cuspidata*, respectively.

Keywords: Lauraceae, Sesquiterpenes, Phenylpropanoids, Breast cancer, *Escherichia coli*.

The Lauraceae is a pantropic family, containing about 50 genera and probably 2500-3500 species, including trees and shrubs. Southeastern Asia and tropical America are the main centers for diversity of the Lauraceae. These species frequently occur in lowland rain forests to tropical montane forests with an altitude up to ca. 4000 m [1, 2]. Many species are highlighted due to their economic importance in cooking, in carpentry and construction, papermaking, in the perfume industry, the chemical industry, and folk medicine [3]. The genus *Nectandra* belongs to the *Ocotea* complex and comprises about 114 species. It is endemic to the Neotropics with restricted distribution in tropical and subtropical Americas, with an estimated 43 species occurring in Brazil [4, 5, 6]. This genus has reported folk medicinal uses as an antifungal, anti-inflammatory, antimalarial, anticancer, analgesic, and febrifuge, as well as being used for the treatment of ulcers, among others [5].

N. puberula Schott (Nees) is commonly known in the Amazon region as "louro amarelo" and "yellow cinnamon" and is presently restricted in distribution to the north, southeast, and southern regions of Brazil [7]. It is a tree up to 20 m in height. Neolignans have been found in the extract of the trunk wood [8]. *N. cuspidata* is a shrub or tree, 5–30 m in height. The plant is not endemic to Brazil, but occurs in most states [2, 7]. A phytochemical investigation of the leaf extract resulted in the isolation and characterization of megastigmane, aporphinoid alkaloids, sesquiterpenes, polyprenols and sterols [9].

Despite the high aromatic potential, there have been few studies on the chemical composition of essential oils (EOs) of *Nectandra* species. To our knowledge, there are no previous reports on either the phytochemistry or the biological activities of the EOs of these two *Nectandra* species.

Fifty-nine volatile components were identified in the EOs from *Nectandra* species, comprising approximately 97.7% of the total composition of the oils (Table 1). The most representative class of compounds in the leaf EOs of both species was oxygenated terpenoids; about 42.3% in *N. puberula* and 76.2% in *N. cuspidata*. Comparison of the chemical profiles of the samples is shown in Figure 1. *N. puberula* oils showed a significant difference in terpene accumulation between leaves and branches. The branch oil displayed a higher concentration of oxygenated sesquiterpenoids (44.7%). The main compounds identified in the EO from *N. puberula* were apiole (22.2%), β -caryophyllene (15.1%), β -pinene (13.3%), germacrene D (8.3%), pogostol (6.6%) and bicyclogermacrene (6.4%) in the leaves, and apiole (28.1%), pogostol (19.8%) and guaiol (11.2%) in the branches. The EO of *N. cuspidata* leaves showed β -caryophyllene (26.9%), bicyclogermacrene (16.0%) and spathulenol (5.2%) as the main compounds.

The occurrence of sesquiterpene hydrocarbons and oxygenated sesquiterpenoids in *Nectandra* species has been reported. The main compounds of the EO from *N. megapotamica* leaves were α -bisabolol (62.3–69.4%) and δ -elemene (8.2–22.6%) [10]. The leaf oil of *N. salicina* growing wild in Costa Rica was dominated by atractylone (14.6%), viridiflorene (10.1%), α -pinene (9.4%), β -caryophyllene (7.2%), α -humulene (7.0%), δ -cadinene (6.1%), β -pinene (6.0%) and germacrene D (5.8%) [11]. Similarly, the leaf oil of *N. membranacea* from Costa Rica was rich in δ -cadinene (14.1%), α -copaene (12.5%), germacrene D (6.9%), α -humulene (5.6%), β -caryophyllene (5.5%), and bicyclogermacrene (4.8%), as well as the monoterpenes α -pinene (22.4%) and β -pinene (12.6%) [12]. The EO of *N. coriacea* showed valerenol (16.7%) and γ -eudesmol (12.3%) [13]. In addition, the occurrence of monoterpenoids in *Nectandra* oils was reported such as mentha-

1(7),8-diene, α -terpinolene, α -pinene, β -pinene and α -terpineol in *N. falcifolia* and *N. elaiophora* [14, 15]. However, the occurrence of phenylpropanoid derivatives such as dillapiole has not been previously reported in volatile oils from *Nectandra* species.

The antiproliferative effect of metabolites isolated from *Nectandra* species against tumor cell lines have been reported [16]. The neolignans from *N. megapotamica* displayed cytotoxic activity and induced apoptosis in leukemia cells (HL-60) [17]. The crude extract of *N. rigida* was rich in dehydrodiisoeugenol, which is reported to be a cytotoxic agent [18]. In this study, the EOs showed good activity against the MCF-7 breast tumor cell line. The IC₅₀ values were 64.5 ± 1.6 and 117.1 ± 11.9 $\mu\text{g}\cdot\text{mL}^{-1}$ for EOs from leaves of *N. puberula* and *N. cuspidata*, respectively.

The most activity was observed for *N. puberula* oil, which is characterized by high contents of apiole (22.2%), a phenylpropanoid, followed by β -caryophyllene (15.1%) and β -pinene (13.3%). The cytotoxic effects against MCF-7 cell lines have been observed for phenylpropanoids such as anethole, cinnamaldehyde and eugenol [19]. Apiole was administered to mice at 1-30 mg/kg body weight through intraperitoneal injection and displayed a promising antitumor effect against colon tumors (COLO 205) in an *in vivo* xenograft model [20]. The IC₅₀ values for α -pinene and β -caryophyllene have been found to be 20.6 and 19.7 $\mu\text{g}\cdot\text{mL}^{-1}$, respectively, against MCF-7 cells [21]. In addition, β -caryophyllene potentiated the anticancer activity of paclitaxel on MCF-7 at a concentration of 10 $\mu\text{g}\cdot\text{mL}^{-1}$ [22]. The main compounds in *N. cuspidata* oil were β -caryophyllene (26.9%), bicyclogermacrene (16.0%) and spathulenol (5.2%). The EO from *N. leucantha*, rich in bicyclogermacrene (28.4%) and spathulenol (5.8%), displayed significant cytotoxic activity against murine melanoma (B16F10-Nex2), human glioblastoma (U-87) and human cervical carcinoma (HeLa) cell lines. However, the IC₅₀ values against MCF-7 were 193.8 $\mu\text{g}\cdot\text{mL}^{-1}$ and 19.0 $\mu\text{g}\cdot\text{mL}^{-1}$ for the EO and bicyclogermacrene, respectively [23].

The antimicrobial property is of great importance in the applications of EO against certain human or animal pathogens [24]. In bacteria, the permeabilization of the membranes is associated with loss of ions and reduction of membrane potential, collapse of the proton pump and depletion of the ATP pool [25]. *Nectandra* leaf oils showed notable antibacterial activity against *E. coli* (MIC = 19.5 $\mu\text{g}\cdot\text{mL}^{-1}$) (Table 2).

The antibacterial activities of the main compounds, apiole, β -caryophyllene, β -pinene, germacrene D, bicyclogermacrene and spathulenol have been reported previously [26]. The EO from *Psammogeton canescens*, rich in α -pinene (20.0%) and apiole (15.3%), exhibited strong antibacterial activity against *Escherichia coli* (MIC, 14.0 ± 0.8 $\mu\text{g}\cdot\text{mL}^{-1}$) [27]. *Eugenia uniflora* oil, rich in apiole (11.1%) and oxygenated sesquiterpenes, was active towards two Gram-positive bacteria, *Streptococcus equi* and *Staphylococcus epidermidis* [28]. The oil of the trunk bark of *Onychopetalum amazonicum*, rich in β -caryophyllene and spathulenol, exhibited good activity against *Staphylococcus epidermidis* (ATCC 12228), *Escherichia coli* (ATCC 10538) and *Kocuria rhizophila* (ATCC 9341), with MIC values of 62.5 $\mu\text{g}\cdot\text{mL}^{-1}$ [29]. In addition, β -caryophyllene exhibited strong antibacterial effect against *E. coli* (MTCC 732), with a MIC value of 9.0 ± 2.2 μM [30].

In summary, our results indicate that *Nectandra* oils could be used as natural antimicrobial agents against *E. coli* and tested in future studies for the treatment of breast cancer.

Table 1: Chemical composition of EOs from *Nectandra* species from the Amazon.

Constituents	RI ^{calc.}	RI ^{Lit.}	<i>N. puberula</i> Leaves	<i>N. cuspidata</i> Leaves
α -Pinene	934	932	5.0	
β -Pinene	978	974	13.3	1.4
Myrcene	992	988	0.8	
Limonene	1028	1024	2.7	3.0
Linalool	1102	1095		0.8
δ -Elemene	1336	1335		4.4
α -Copaene	1374	1374		4.6
β -Bourbonene	1386	1387	0.5	
β -Elemene	1391	1389	2.0	2.1
β -Caryophyllene	1421	1417	15.1	2.8
β -Copaene	1427	1430		0.5
β -Gurjunene	1432	1431		0.4
α -trans-Bergamotene	1435	1432		3.1
γ -Elemene	1435	1434		0.9
α -Guaiene	1440	1437	1.3	0.8
Aromadendrene	1443	1439		0.4
α -Humulene	1452	1452	1.8	1.2
(E)- β -Farnesene	1458	1454		1.5
γ -Murolene	1476	1478		0.5
Amorpha-4,7(11)-diene	1480	1479		4.5
Germacrene D	1483	1484	8.3	4.5
γ -Himachalene	1484	1481		2.5
cis-Eudesma-6,11-diene	1488	1489	2.3	1.7
unidentified	1490		1.1	
epi-Cubebol	1496	1493		1.5
Bicyclogermacrene	1498	1500	6.4	16.0
4-epi-cis-Dihydroagarofuran	1501	1499		1.0
α -Bulnesene	1508	1509	2.3	1.7
β -Bisabolene	1509	1505		1.4
γ -Cadinene	1512	1513		1.2
7-epi- α -Selinene	1516	1520		0.3
δ -Cadinene	1523	1522	1.1	3.0
trans-Cadina-1,4-diene	1531	1533		0.5
Elemol	1551	1548		0.9
Germacrene B	1558	1559	1.3	
Murol-5-en-4- α -ol	1558	1559		1.2
Maaliol	1564	1566		0.6
Caryophyllenyl alcohol	1567	1570		0.3
Spathulenol	1578	1577	1.7	5.2
Unidentified	1583			4.7
Caryophyllene oxide	1584	1582	2.1	1.3
Viridiflorol	1590	1592		1.6
Guaiol	1597	1600		11.2
Rosifolol	1600	1600		0.9
6-Methoxy-elemicin	1601	1595	1.7	
Rosifolol	1610	1600		0.7
unidentified	1621			0.6
1-epi-Cubenol	1626	1627		0.7
Murola-4,10(14)-dien-1- β -ol	1630	1630		1.1
Camphoric acid	1633	1634		0.6
allo-Aromadendrene epoxide	1637	1639		1.6
epi- α -Murolol (= α -Murolol)	1640	1640		1.5
α -Murolol (=Torreyol)	1644	1644		0.3
α -Eudesmol	1652	1652		1.3
α -Cadinol	1653	1652		1.7
Selin-11-en-4- α -ol	1655	1658		0.4
Pogostol	1656	1651	6.6	19.8
Unidentified	1660		0.7	
Bulnesol	1669	1670		1.4
epi- β -Bisabolol	1670	1670		1.4
α -Bisabolol	1682	1685		1.0
Apiole	1687	1677	22.2	28.1
Monoterpene hydrocarbons		21.7	4.4	
Oxygenated monoterpenoids			0.8	
Sesquiterpene hydrocarbons		42.3	19.5	76.2
Oxygenated sesquiterpenoids		10.3	44.7	18.0
Phenylpropanoids		23.9	28.1	
Others			0.6	
Total identified		98.3	100.0	94.8

RI^{Calc.} = based on DB-5ms capillary column and alkane standards (C8-C32).

RI^{Lit.} = based on Adams [31].

Table 2: Antimicrobial activity of *Nectandra* essential oils from the Amazon.

	Minimum inhibitory concentration ($\mu\text{g}\cdot\text{mL}^{-1}$)	<i>N. puberula</i>	<i>N. cuspidata</i>
Gentamicin (positive control)			
<i>P. aeruginosa</i>	<19.5	1250	1250
<i>E. coli</i>	<19.5	19.5	19.5
<i>S. epidermidis</i>	<19.5	1250	1250
<i>S. aureus</i>	<19.5	625	625
<i>B. cereus</i>	<19.5	625	312.5

Experimental

Plant material: *Nectandra puberula* Nees was collected in Santarém (S 02° 25.0' 10.2" W 54° 44' 26.9") and a voucher (HSTM000092) was deposited in the herbarium of Universidade Federal do Oeste do Pará, Santarém, Pará State, Brazil. *N. cuspidata* was collected in Caxiuanã National Forest, Marajó Island (S 01° 44' 18.8" W 51° 27' 27.4"), and a voucher (MG 104948) was deposited in the Herbarium of Museum Paraense Emílio Goeldi, Belém, Pará state, Brazil.

Leaves and branches from several mature plants were air-dried, pulverized, and subjected to hydrodistillation using a Clevenger-type apparatus (100 g, 3 h). The essential oils were dried over anhydrous sodium sulfate, and their percentage contents were calculated on the basis of the dry weight of plant material. The moisture contents of the samples were calculated after phase separation using a Dean–Stark trap (5 g, 60 min) using toluene as the solvent phase.

Gas chromatographic – mass spectral analysis: The essential oils were analyzed by GC-MS using an Agilent 6890 GC with an Agilent 5973 mass selective detector (MSD) [operated in the EI mode (electron energy = 70 eV), scan range = 40–400 amu, and scan rate = 3.99 scans/sec], and an Agilent ChemStation data system. The GC column was an HP-5ms fused silica capillary with a (5% phenyl)-polymethylsiloxane stationary phase, film thickness of 0.25 µm, length of 30 m, and internal diameter of 0.25 mm. The carrier gas was helium with a column head pressure of 48.7 kPa and a flow rate of 1.0 mL/min. Injector temperature was 200°C and detector temperature was 280°C. The GC oven temperature program was used as follows: 40°C initial temperature, held for 10 min; increased at 3°C/min to 200°C; increased at 2°C/min to 220°C. A 0.2%, w/v, solution of the sample in CH₂Cl₂ was prepared and 1 µL was injected using a splitless injection technique. The percentages of each component are reported as raw percentages based on total ion current without standardization. Individual components were identified by comparison of both mass spectrum and GC retention data with authentic compounds present in commercial libraries [31].

Antibacterial assay: The essential oils were screened for antimicrobial activity against *Escherichia coli* (ATCC No. 10798), *Pseudomonas aeruginosa* (ATCC No. 27853), *Bacillus cereus* (ATCC No. 14579), *Staphylococcus aureus* (ATCC No. 29213), and *Staphylococcus epidermidis* (ATCC No. 12228), using the microbroth dilution technique, as previously reported [32]. Thus, 50 µL of 1%, w/v, solution of the samples in DMSO was placed in a

well of 96 well plates and 50 µL of cation-adjusted Mueller Hinton broth (CAMHB) was added. The sample solutions were then serially diluted (1:1) by transferring 50 µL of sample-CAMHB mixture to the next lane and adding 50 µL fresh CAMHB to obtain concentration from 2500 µg.mL⁻¹ to 12.5 µg.mL⁻¹. The bacteria were harvested from a fresh culture and added to each well at a concentration of approximately 1.5 × 10⁸ colony forming units (CFU)/mL. The plates were incubated at 37°C for 24 h and the final minimum inhibitory concentration (MIC) was determined as the lowest concentration with no turbidity. Gentamicin was used as positive antibiotic control and DMSO was used as negative control.

Cytotoxic assay: MCF-7 human breast adenocarcinoma cells (ATCC No. HTB-22) were grown in RPMI 1640 supplemented with 10% fetal bovine serum (FBS), 30 mM HEPES, NaHCO₃, and penicillin streptomycin. *In vitro* cytotoxic activity of essential oils on MCF-7 cells was performed using the 96-well MTT assay, as previously reported [33]. Cells were plated into 96-well cell culture plates at a concentration of 1.2 × 10⁴ cells/well and a volume of 100 µL in each well. The plate was then labeled and incubated at 37°C and 5% CO₂ for 48 h. By then, the cells had reached 70–80% confluent growth. The supernatant fluid was gently aspirated (without touching the bottom of the well to avoid removing cells) and replaced with 100 µL growth medium containing 1.0 µL and 0.5 µL of essential oil (1% in DMSO), giving a final concentration of 100 and 50 µg.mL⁻¹. The plate was then incubated at 37°C and 5% CO₂ for 48 h. Then, the liquid was gently aspirated from each well. In a tube, 10 mL feeding medium was mixed with 2 mL of MTT stock solution (and was protected from light). Into each well, 100 µL of the MTT solution was added and the pre-read absorbance was immediately measured spectrophotometrically at 570 nm (using a Molecular Devices SpectraMax Plus 384 microplate reader). Formazan crystals were formed over the course of 4 h at 37°C and 5% CO₂. After 4 h, DMSO was used to dissolve the purple crystals. The amount of MTT-formazan produced was determined spectrophotometrically at 570 nm. Growing medium, DMSO, and tingenone (100 µg.mL⁻¹) served as negative, compound, and positive controls, respectively. Solutions were added to wells in 8 replicates. Average absorbances, standard deviations, and percent kill ratios (% kill_{oil} / % kill_{control}) were calculated. Median inhibitory concentrations (IC₅₀) were determined using the Reed-Muench method [34].

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Chemical Composition and Acaricidal Activity of *Thymus algeriensis* Essential Oil against *Varroa destructor*

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The aim of the present study was to determine the chemical composition and evaluate the acaricidal activity of *Thymus algeriensis* essential oil (TAEQ) against *Varroa destructor*. This ectoparasitic mite is a pest of the honey bee *Apis mellifera*. The essential oil from the aerial parts of *T. algeriensis*, obtained by hydrodistillation, was obtained in a yield of 2.8± 0.2%, w/w. The TAEQ was analyzed by GC and GC/MS. Thirty-four compounds were identified, representing 99.3% of the oil. The main constituents were carvacrol (48.4%), γ -terpinene (14.9%), *p*-cymene (14.7%), and thymol (5.6%). Four lots were constituted at the level of an apiary in order to study the dynamics of the *Varroa destructor* and its host, *Apis mellifera*. After diagnosis by the biological method "install of diapers", the lots were treated at different doses of TAEQ (0.1, 0.3 and 0.5%). TAEQ was sprayed on top of the hives. The results show that TAEQ at 0.5% resulted in a decrease in the rate of infestation of *Varroa destructor*, causing a mortality rate of 32.6% without negative effect on the nesting of the queen. The essential oil of *T. algeriensis* could be used as a bioacaricidal agent.

Keywords: *Thymus algeriensis* L., *Varroa destructor*, *Apis mellifera*, Essential oils, Acaricidal activity.

The honey bee, *Apis mellifera* L., is critical for crop pollination and honey production. The ectoparasitic mite *Varroa destructor* is a pest of the honey bee. This mite, which feeds on haemolymph of brood and adult bees, causes colony disorder, weakness, decrease in brood and deformation of bees [1a]. It also reduces the ability of the colony to pollinate plants [1b]. The parasite destroys the mechanical protective barriers of the integument and impairs the immune system of the bees. The disease caused by the *Varroa* mites is called varroasis and is one of the main causes of economic damage to the beekeeping industry. Varroasis was reported for the first time in Umm Teboul (East of Algeria) in June 1981, in an apiary of a beekeeping cooperative. Currently, this parasite has spread rapidly throughout the country. Natural products having components with various modes of action might provide an effective solution to the problem of varroasis [1c]. Essential oils might be an option to control *V. destructor*. Several authors report that these oils might be useful in reducing both mite infestation rates and hive contamination [1d-1e].

The genus *Thymus* L. (Lamiaceae) comprises more than 250 species growing wild throughout the world. This genus is represented by eleven species in the flora of Algeria, one of which is *T. algeriensis* Boiss. et Reut. *T. algeriensis* is the most endemic widespread North African species and well known as "Zaatar". The chemical composition of *T. algeriensis* essential oil has been extensively studied in different Maghreb countries [2a-2o], and the existence of several chemotypes has been revealed (Table 1). Therefore, the aims of this study were to analyze further the composition of the essential oil of wild *T. algeriensis* from Ain-Defla (Northern Algeria), and to determine its acaricidal activity

against *V. destructor* and its host, *A. mellifera*. To the best of our knowledge, this study represents the first report on acaricidal activity of *T. algeriensis* essential oil.

Extraction afforded a yellow-red oil with a very strong and persistent odor of *Thymus*. The essential oil yield was 2.8 ± 0.2%, w/w, (n= 03). This plant can be assigned to one of the oil-rich species of the Lamiaceae. The percentages and retention indices of the identified oil components are listed in Table 2 in the order of their elution on the HP-5MS column. Thirty-four components were identified, accounting for 99.3% of the total oil. This was found to be rich in monoterpene phenols (carvacrol: 48.4% and thymol: 5.6%) and their corresponding monoterpene hydrocarbon precursors: *p*-cymene (14.7%) and γ -terpinene (14.9 %). Oxygenated monoterpenes were the predominant chemical group (58.4%), followed by monoterpenes (38.2%), whereas the sesquiterpenoid content was very low (2.7%). Chemical profiling of the TAEQ sample revealed that it could belong to the carvacrol chemotype, as is quite common for Algerian samples of this species. The compositions of samples from North African countries showed that carvacrol and thymol were individually or jointly the predominant components. Thus, thymol (36.8%) and myrcene (20.2%) were reported as the major components for *T. algeriensis* from Libya [2c], while other authors cited thymol (14.4–65.1%) and carvacrol (22.8–70.3%) as the major compounds for Moroccan samples [2b].

According to the main compounds, several chemotypes were identified in wild plants growing in Tunisia [2h]. In Algeria, two chemotypes were found: thymol/ *p*-cymene/ γ -terpinene (i) and

Table 1: Major compounds of *T. algeriensis* essential oils (%) from various countries.

Country	Ref ^{a)}	Major compounds ^{b)} (%)								
		1	2	3	4	5	6	7	8	9
Libya	[2c]		36.8	20.2						
	[2a]	4.7	38.5	8.9	7.2					
	[2i]		38.5	8.9	7.1					
	[2b]	22.8-70.3	14.4-65.1							
Morocco	[2m]	8.1	37.8	11.7	15.1					
	[2k]	80.4	3.4	5.0	2.0	1.8				
	[2k]	49.3	0.8	2.6	0.9	0.8				
	[2j]						20.5	27.7		
Algeria	[2h]						17.7		15.5	8.2
	[2g]		29.5	13	6.9					
	[2e]			9.9			6.5	10.6	6.5	10.1
	[2l]	29.2		6.8		43.3				
	[2l]	4	71		3	0.5				
	[2d]		62.7							
			78.8							

^{a)} Ref: References; ^{b)} Compound: 1. Carvacrol; 2. Thymol; 3. p-Cymene; 4. γ -Terpinene, 5. Linalool; 6. 1,8 -Cineole; 7. 4-Terpineol; 8. α -Pinene; 9. Camphor.

Table 2: Chemical composition of *Thymus algeriensis* essential oil from Northern Algeria (mean of triplicates).

Compound	RI ^{a)}	RI ^{b)}	Area (%)
α -Thujene	930	923	1.4 \pm 0.6
α -Pinene	939	930	1.1 \pm 0.4
Camphepane	944	945	tr
Sabinene	975	970	tr
β -Pinene	979	973	tr
1-Octen-3 ol	977	977	0.2 \pm 0.1
β -Myrcene	991	988	2.7 \pm 0.9
α -Phellandrene	1003	1002	0.3 \pm 0.1
α -Terpinene	1017	1015	2.2 \pm 0.9
p-Cymene	1025	1023	14.7\pm 2.6
Limonene	1029	1028	0.8 \pm 0.3
<i>trans</i> β -Ocimene	1050	1048	tr
γ -Terpinene	1060	1059	14.9\pm 2.8
<i>cis</i> -Sabinene hydrate	1070	1066	0.1 \pm 0.1
α -Terpinolene	1089	1088	0.1 \pm 0.1
Linalool	1097	1098	1.2 \pm 0.6
1-Octen-3-yl-acetate	1116	1113	0.4 \pm 0.1
Camphor	1146	1145	tr
Borneol	1175	1167	0.1 \pm 0.1
Terpinen-4-ol	1177	1177	1.5 \pm 0.7
<i>trans</i> -Dihydrocarvone	1201	1197	0.1 \pm 0.1
Carvacrol methyl ether	1245	1244	1 \pm 0.6
Thymol	1290	1293	5.6\pm1.8
Carvacrol	1299	1306	48.4\pm4.2
Terpinyl acetate	1347	1340	0.1 \pm 0.1
α -Gurjunene	1411	1413	0.7 \pm 0.2
β -Caryophyllene	1419	1423	0.1 \pm 0.1
Aromadendrene	1441	1443	0.2 \pm 0.1
Alloaromadendrene	1460	1465	0.1 \pm 0.1
Bicyclosesquiphellandrene	1473	1479	0.1 \pm 0.1
Ledene	1493	1499	0.2 \pm 0.1
γ -Cadinene	1506	1517	0.1 \pm 0.1
δ -Cadinene	1523	1526	1 \pm 0.4
Caryophyllene oxide	1583	1582	0.2 \pm 0.1

tr: traces (<0.1); RI^{a)}, retention indices [4a]; RI^{b)}, retention indices relative to C7-C30 on HP-5MS capillary column. Monoterpene hydrocarbons: 38.2%. Oxygen-containing monoterpenes: 58.4%; Sesquiterpene hydrocarbons: 2.5%; Oxygen-containing sesquiterpenes: 0.2%; Total identified: 99.3%

terpinyl acetate / nerolidol / α -pinene / borneol / bornyl acetate (ii) [2g]. The oil composition of two *T. algeriensis* samples collected from the same location was different with one thymol-rich and the other linalool-rich [2d]. TAEAO showed a large variability and displayed different chemical profiles. This great variability and diversity observed in the chemical composition of the essential oils of *Thymus* can be attributed to many factors, including climatic and soil variations, stage of the vegetative cycle, seasonal variation, and the method of preservation and extraction.

Diagnosis of the presence of *Varroa destructor* mite in an apiary revealed the existence of the mite in all the hives. All colonies of the hive were parasitized by the *Varroa* mite and presented a degree of infestation variable from 1.4 to 34%. There is a correlation between the number of *Varroa* collected daily and the total population within the colony. The mite population can be estimated throughout the year by multiplying the daily mite drop by: 250 - 500 or 20 - 40 when the brood is either absent or present, respectively [3a].

In order to ensure the survival of the colony the following year, anti-*Varroa* treatment is imperative, and some authors propose critical periods of intervention [3b] to the apiculture industry. To reduce the infestation threshold to 50 mites, it is necessary that the acaricidal treatment is conducted in August before the breeding of the winter bee brood [3c].

In order to determine the effect of treatment doses on the nesting of the queen, calculation of the brood area is necessary. The essential oils were diluted with analytical grade ethanol to the following concentrations: 0.1, 0.3, and 0.5%, v/v. The results of the treatments are represented in Figure 1. Lot D presents a very high brood area (1147 cm²), followed by lot B (913 cm²), A (903 cm²), and C (447.5 cm²), respectively. The hives treated by TAEAO (0.5%) represent the best brood area with an evolution from 925 - 1147 cm² corresponding to an increase of 225 cm². This means that this treatment had no adverse effect on the activity of the bee colony and the laying of the queen. As for the duration of application of treatment, the difference is not significant ($p = 0.4123$). Finally, the treatment did not disrupt the population of the hive; the brood remained compact and hatching continued normally, indicating the safety of the essential oils to the colony bees. The treatment with TAEAO had no influence on the activity of the colony bees and the laying of the queen. Application of TAEAO, in winter (from 14/01 to 16/03) reduced the infestation of various lots infected by *V. destructor*. The results of the treatments at different concentrations (0.1, 0.3, and 0.5%) are represented in Figure 2. Mortality rates obtained were: A (4.1%), B (24.0%), C (32.4%) and D (32.6%), an average number of dead mites: A (173), B (435) C (1274) and D (1366). These results are consistent with those reported in the literature [3d]. Concerning the factor treatment dose, there is a clear significant difference ($p < 0.05$) between the four lots. Statistical analysis shows that treatment of lots A and B had no effect on the mortality of *Varroa*; then, it is very significant for lots C and D, which shows the effectiveness of treatment by the TAEAO concentration of 0.5%. There is a strong correlation between the number of mites killed and the concentrations of the oils tested. One possible explanation for this result is the presence of carvacrol and thymol as major components and the synergic effect with other monoterpenes, such as *p*-cymene, and γ -terpinene. Indeed, previous studies have found that carvacrol was acaricidal against several species of ticks, while carvacrol and γ -terpinene acted as miticides [3e]. Thymol and structurally related compounds like *p*-cymene are effective as acaricides; it has an acaricidal activity against *Varroa*, but while decreasing the laying of the queen [3f]. γ -Terpinene, another active substance of *Thymus* oil, has a very good varroacidal effect. However, the use of *Thymus* oil containing γ -terpinene at a concentration of 250 μ g/L of air has proved to be very toxic for both *V. destructor* and bees [1c]. On the other hand, the third application of TAEAO (0.5%) presented the most important rate of mortality (340 mites) during the treatment. The statistical analysis also reveals that this application ($p = 0.012$) corresponds to the stadium phoretic of the *Varroa* mite (Figure 2). During the period of development of the laying and nesting of the queen, the natural

mortality of *Varroa* decreases. This result is consistent with the literature [3g]. Also, it is slightly higher than that obtained by spraying oxalic acid at 0.56 g per colony on adult worker bees, where a mite mortality rate in the phoretic stage of 25.9% was observed [3h]. So, it is much higher than that obtained by applying, for 12 days, an extract of *Lantana camara*, when the infestation rate was reduced to 0.20% [3i]. Treatment with *Citrus aurantium* and *Cymbopagon flexuosus* reduced the average percentage of *Varroa* infestation by 100% after the fourth week [3j]. However, the spray of thyme oils resulted in the death of 65.9% of *Varroa* [3k]. In late autumn, the *Varroa* mite is very sensitive to essential oils due to the formation of a cluster and the absence of brood [3l]. When the external temperature range is from 15-20°C, some authors recommend the application of products containing thymol, because, beyond this range, outside temperature affects negatively the efficiency of the thymol and bee activity [1c]. These conditions are for our study and in the climatic conditions of Algeria in late winter with a temperature of 12-19° C. It follows that the application, in winter, of *T. algeriensis* essential oil by spraying inside the hives is very effective against infestation by *V. destructor* and for maintaining the activity of colony bees and the laying of the queen.

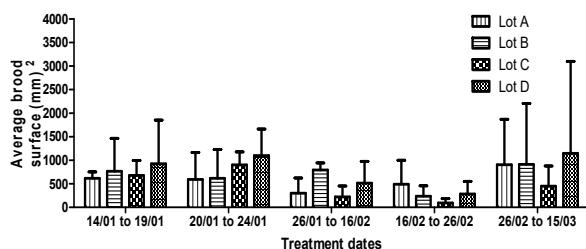


Figure 1: Average brood area during the spraying period of *T. algeriensis* essential oils (TAEAO) at different doses. Lot A: control (untreated colonies), Lot B: TAEAO (0.1%), Lot C: TAEAO (0.3%), Lot D: TAEAO (0.5%).

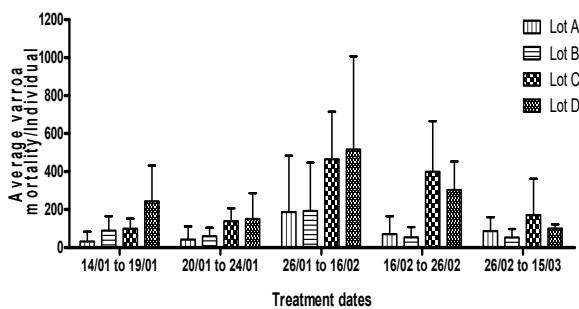


Figure 2: Average mortality of *Varroa* during the spraying period of *T. algeriensis* essential oils (TAEAO) at different doses. Lot A: control (untreated colonies), Lot B: TAEAO (0.1%), Lot C: TAEAO (0.3%), Lot D: TAEAO (0.5%).

The use of natural products as acaricides may represent an important alternative for the control of *V. destructor* since they are rich sources of bioactive compounds that are biodegradable. The present findings demonstrate that *T. algeriensis* essential oil may be used as a natural acaricide. Carvacrol, thymol, *p*-cymene, and γ -terpinene are the major compounds in *T. algeriensis* essential oil and may act synergistically to produce the observed acaricidal action against *V. destructor*. Treatment with TAEAO (concentration 0.5%) applied in the climatic conditions of Algeria in late winter, is effective against the mite *V. destructor* and has no influence on the activity of the colony bees and the laying of the queen.

Experimental

Biological material: This study has been carried out in an apiary installed at Ain Defla (Algeria). Twenty hives were randomly

distributed in 4 blocs (A, B, C, and D), each block containing 5 hives. Block A (hives witnesses) was left without applications. The other blocs were treated with different doses (0.1, 0.3, and 0.5%) of *T. algeriensis* essential oil in 96% ethanol.

Plant material and essential oil hydrodistillation: Aerial parts of wild *T. algeriensis* were collected in the first week of June 2013 in Mekhatria within the region of Ain-Defla located in northern Algeria (at 140 Km Northwest of Algiers - latitude: 36°25' N; longitude: 2° 21'7" E; Altitude: 365m). A voucher specimen was deposited in the Herbarium of the Agronomic Department, Djilali Bouaama University of Khemis Miliana. Air-dried plants (50 g) with 600 mL distilled water (1:12, w/v) were separately subjected to hydrodistillation for 2 h using a Clevenger-type apparatus. All experiments were conducted in triplicate and results were expressed on the basis of dry weight.

Essential oil analysis: Ten mg of essential oil was dissolved in 5 mL of diethyl ether. The essential oils were analyzed by gas chromatography coupled to a flame ionization detector (GC-FID) and by gas chromatography coupled to a mass spectrometer (GC-MS).

GC-FID analysis: The analysis of the oil was carried out by means of an Agilent technology HP GC 6890 system with a flame ionization detector (FID), using a capillary column coated with 5 % phenyl-methyl siloxane (30 m x 0.25 mm x 0.25 μ m film thickness Agilent Technologies, Hewlett-Packard, CA, USA). The temperature program was as follows: 40°C during 1 min, then raised in a first ramp to 200°C at 6°C/min, followed by a second ramp to 280°C at 30°C/min, and finally kept at 280°C during 2 min. Injection was realized in splitless mode at 280°C; the volume injected was 1 μ L of diluted oil (10 mg of oil/5 mL diethyl ether). Detector temperature was fixed at 300°C; Carrier gas was helium at 1 mL/min.

GC-MS analysis: GC/MS was performed with an Agilent HP 6890 GC system coupled with an Agilent HP 5973 Network Mass Selective Detector operated by HP Enhanced ChemStation software. Analytical conditions were fixed as follows: Agilent HP-5MS capillary column (30 m x 0.25 mm, df = 0.25 μ m), a split-splitless injector at 250°C (splitless mode), temperature program: from 40°-250°C at 6°C/min, mobile phase: carrier gas helium at 1 mL/min. The mass spectra were recorded in EI mode (70 eV), scanned mass range: from 35 to 500 amu. Source and quadrupole temperatures were fixed at 230°C and 150°C, respectively. The identification of the components was performed on the basis of chromatographic retention indices and by comparison of the recorded spectra with computed spectral library (Wiley 275. L) [4a]. For sesquiterpene hydrocarbons, further confirmations were obtained by comparing the mass spectra with data from the literature [4a-4b]. Retention indices (RI) were calculated by means of a mixture of homologue *n*-alkanes (C_7-C_{30}) analyzed under the same chromatographic conditions used for the analysis of essential oils [4a].

Diagnosis of the presence of the Varroa mite before treatment by spraying: A diagnosis using the biological method "Installing the swaddling clothes" was performed before the treatment with *T. algeriensis* essential oil during the summer period. It allows the detection of the presence of the parasite and then to confirm and to assess the degree of infestation. Also, the diagnosis allowed us to establish a procedure to follow in order to preserve the bees in the best possible conditions [4c]. Among the biological methods for the diagnosis, the utilization of diapers or cover-bottoms was chosen. These swaddling clothes were smeared with fat (Vaseline)

supported by a grid on the floor of the hive. The swaddling clothes were then removed and examined carefully using a magnifying glass to detect dead *Varroa* mites. This method lasted for 30 days during which the swaddling clothes were replaced in the morning once every 3 days.

Treatment of the Varroa mites by spraying: The technique used was the spraying of TAE0 at different doses (0.1, 0.3, and 0.5% in

96% ethanol). It was applied to the top of the frames of the hive using sprayers to ensure contact of the treatment with the *Varroa* mites. The treatments were applied all 6 days during the winter period from 14/01/2012 to 26/02/2012. Then, dead *Varroa* mites were counted on the dashboard greased all 3 days with the aid of a magnifying glass. The surface of the brood was calculated using the equation of an ellipse.

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Cereal Lignans, Natural Compounds of Interest for Human Health?

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Cereals are suggested to be the most important sources of lignan in the diets of western populations. Recent epidemiological studies show that European subpopulations in which the major source of lignans are cereals, display lower disease frequency regarding metabolic and cardiovascular diseases. The biological mechanisms of lignan are several. Beyond their antioxidant and anti-inflammatory actions at nutritional doses some lignans regulate the activity of specific nuclear receptors (NRs), such as the estrogen receptors (ERs), and also NRs that are central switches in glucose and fatty acid metabolism such as PPAR α , PPAR γ and LXRs, highlighting them as selective nuclear receptor modulators (SNRMs). These include enterodiol (END) and enterolactone (ENL), the metabolites produced by the gut microbiota from food lignans. The available knowledge suggests that given some additional research it should be possible to make 'function' claims for a regular intake of lignans-rich foods related to maintaining a healthy metabolism.

Keywords: Lignan, Cereal bran, Obesity, Metabolic syndrome, Health claims.

Introduction: Lignans are a group of compounds found in fiber-rich foods such as cereals, oilseed, nuts, vegetables (brassica), and fruits (berries) typically regarded as components of healthy diets [1-3]. They are expected to have antioxidant, anti-inflammatory and hormone-mimicking effects in humans both as native compounds and after fermentative conversion in the colon [4-7].

Intakes of lignan-rich foods have been shown to be correlated with a lower incidence and progression of diseases / health conditions related to the metabolic syndrome, type 2 diabetes, cardiometabolic risk factors, specific types of cancers and overall mortality [8-12]. The most studied food sources rich in lignans, such as oilseeds (flaxseed and sesame seeds), cereals, and in particular cereal fibers, may contain sufficient concentrations of specific lignans to exert functional effects in humans. Amongst cereals, wheat and rye contain the highest concentration of lignans, mostly in their bran fraction. The most abundant lignans here are 7-hydroxymatairesinol and syringaresinol [14, 15].

Recent studies have highlighted some of the possible mechanisms of action of lignans. Almost all are endowed with antioxidant and anti-inflammatory activities. However, some of these compounds have also shown the ability to bind and activate/inhibit nuclear receptors (NRs) such as the estrogen receptors (ERs) [7], the peroxisome proliferator activate receptor gamma (PPAR γ) [16], and the liver X receptor alpha (LXR α) [17]. The regulation of gene expression through these receptors may require the metabolization of most plant lignans to the mammalian lignans enterodiol (END) and enterolactone (ENL) [18], while some lignans like 7-hydroxymatairesinol are shown to engage NRs also in their unmetabolized form [19]. Metabolization of lignans requires gut microbial processes to release them from the food matrix and undergo deglycosylation in the intestine and conversion in the liver to active metabolites. Several of the lignans that have been studied so far have shown interesting roles in promoting human health. Some of them were identified in plants to be of pharmaceutical and cosmetic interest although the diet remains the major route of intake of these chemicals [5, 13, 20, 21]. In particular cereal lignans are

present in high amounts in the daily diet of western populations and are likely to be the dietary plants that provide most of the beneficial bioactive lignans in Europe, the US and Canada [5, 13, 20, 21]. Here we review the recent and ongoing research on the characterization of the health effects of cereal lignans and their mechanisms.

Methods: A comprehensive computer literature search of the PubMed/MEDLINE, Embase and Scopus databases was conducted to find relevant published articles about the role of lignans in health. No beginning date limit was used; the search was updated until January 2016. Only articles in the English language were selected. To expand our search, references of the retrieved articles were also screened for additional studies.

Lignan chemistry and biochemistry: The chemistry of lignans has been studied for the past 65 years. They possess very different structural composition and can be found in nature in vegetables, mostly in exudates and resins, although they can be present in different parts of the plant. Lignans are polyphenolic substances derived from phenylalanine via dimerization of substituted cinnamic alcohols, known as monolignols, to a dibenzylbutane skeleton and they are found mostly in plants as aglycones, esters or glycosides. These dimeric phenylpropanoids show two C6-C3 units attached to the central carbon C8 while, when two phenylpropanoid units are found coupled in a different manner, like C5-C5', the new term "neolignan" has been adopted (Figure 1) [22].

The lignans can be classified into eight chemical groups and several subgroups based upon the way in which oxygen is incorporated into the skeleton and on the cyclization pattern [22]. Lignan subgroups arise via the shikimic acid pathway from the reduction of cinnamic acid derivatives produced from aromatic aminoacids (L-phenylalanine and L-tyrosine) and subsequently reduced via coenzyme A to alcohols that are the main precursors of all lignans (*p*-coumaryl alcohol, coniferyl alcohol and sinapyl alcohol) in plants. The enantioselective dimerization of two coniferyl alcohol monomers via intermolecular 8,8' oxidative coupling, generate pinoresinol, a major lignan in food seeds and grains [23].

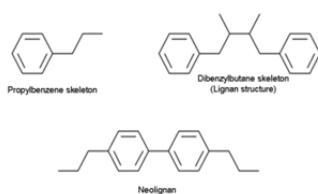


Figure 1: Basic structure of lignans. a and b) The propylbenzene skeleton and the dibenzylbutane skeleton of lignan attached C6-C3 to the central carbon C8. c) Two two phenylpropanoid units of a neolignan coupled C5-C5'.

Subsequent metabolism steps generate lariciresinol and secoisolariciresinol through pinoresinol/lariciresinol reductase and secoisolariciresinol dehydrogenase respectively. Secoisolariciresinol is the common precursor of all dibenzylbutyrolactol lignans including matairesinol, yatein and aryltetralin lignans (Figure 2). These subclasses of lignans includes some important bioactive chemicals from which several structural analogues have been derived, studied and employed as pharmaceuticals [24].

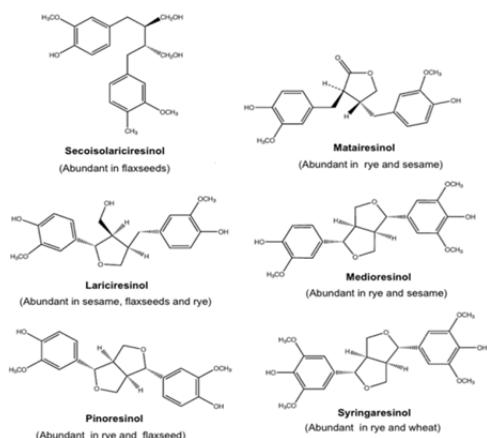


Figure 2: Major lignans in oilseeds and cereals. Secoisolariciresinol is the common precursor of all dibenzylbutyrolactol lignans. These subclasses of lignans include some important bioactive chemicals from which several structural analogues have been derived, studied and employed as pharmaceuticals [24].

Lignans can be converted by intestinal microbiota, under strictly anaerobic condition, to the mammalian lignans, enterodiol (END) and enterolactone (ENL). These compounds, also called enterolignan, following metabolism steps in the gut acquire specific bioactivities and healthy effects, some of which have been shown to occur through the interaction with nuclear receptors [18]. The biotransformation of plant lignans to enterolignan has been extensively studied with secoisolariciresinol (SDG) through the pathway consisting of glycoside hydrolysis, demethylation and dehydroxylation and intermediates [25]. The metabolization steps will be further discussed in the last paragraph of this review in a more detailed manner.

Lignans in nutrition, availability and food sources: Several of the lignans under study are present in plants of a very different kind throughout the whole plant kingdom. Some of them are highly concentrated in plants of medical interest, i.e. *Podophyllum* species containing podophyllotoxin, which is particularly interesting for its anticancer properties [24, 26]. High concentrations of lignans are also found in plants of nutritional interest. The richest are by far oilseeds such as flaxseed and sesame, with concentrations up to 1% of the seed weight. These seeds are consumed in functional amounts

in defined geographical regions of the world, in particular the Middle East, North Africa and Asia.

Cereals, nuts, broccoli (*Brassica* species), fruits, berries and beverages (tea, coffee, beer wine) are suggested to be the most important sources of lignans in the diets of Europeans. According to a Dutch lignan database, the average daily intake of lignans in the Netherlands was 1000 to 2000 µg/day and in the Finnish population 1081 µg/day, ranging from 1136 µg/day for men to 1036 µg/day for women) [13]. In another study in the Finnish population the average lignan intake based on urinary excretion of lignan was 1224 µg/day [27]. Sweden registered the highest intake in Europe with 1773 µg/day total lignans for women and 1947 µg/day for men. Intakes between 1062 and 1188 µg/day have been reported in south Europe (Italy) [13] and 600 µg/day in the USA (Breast Cancer and Environment Research Project Puberty Study) [28]. Higher levels of intake (up to 13.5 mg/day) have been reported in previous studies in the USA and Canada [21]. In the east, a Chinese study reported intakes measured through the urinary excretion of mammalian lignans giving values in the range of 0.30 and 1.18 nmol/mg creatinine of enterodiol and enterolactone, respectively. The levels of enterodiol were two-fold higher in the Chinese population compared with US women in a similar age range, while enterolactone was observed at similar levels in both populations [29].

Cereal lignans: Up to date, the most investigated lignans have been extracted from food sources such as flaxseed and sesame seeds, the first being known as the richest food source of the lignan secoisolariciresinol and the latter being the most consumed lignan-rich food in the world with a unique lignan composition. However, in the western countries cereals are the major source of lignans (because of their regular daily consumption) with an impact that may influence human health (Table 1).

Wheat: Wheat is the most consumed cereal in the diet in Europe, the USA, and Canada. It is well accepted and has an attractive low price as food, but also as a lignan source. By wheat milling the grain is split into two major parts, the flour (about 70-80%), and the bran fraction, which is approximately 15%. This last fraction contains most of the grain lignans (70-85%), which are mostly situated within the aleurone cells, the single cell layer at the inner site of the bran.

Novel milling and dry-fractionation techniques and new extraction and analytical methods have recently allowed full-scale separation of aleurone cells from the other layers of wheat bran, yielding a fiber rich concentrate which contains many nutrients and well-known bioactive lignans, as well as several new ones with potent bioactivity (Figure 3) [31, 32].

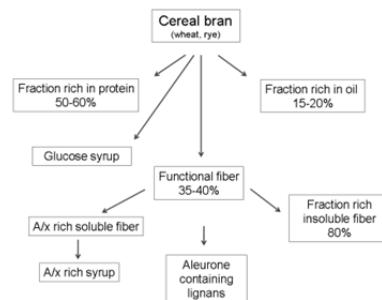


Figure 3: A schematic representation of the fractionation of cereal bran (approximately 15% of the grain). Most of the grain lignan (70-85%) is situated within the aleurone, the single cell layer at the inner site of the bran. Separation of the aleurone cells from the other layers of the cereal bran yields a fiber rich concentrate which, beyond well-known and new lignans, contains many other nutrients.

Table 1: Concentration of main lignan in different food sources.

Food sources	Lignan ($\mu\text{g}/100 \text{ g}$)						
	Pinoresinol	Syringaresinol	Lariciresinol	Secoisolariciresinol	Matairesinol	7-Hydroxymatairesinol	Sesamine
Flaxseeds	871	48	1780	165759	529	35	ND
Sesame	47136	205	13060	240	1137	7209	62724
Rye bran	1547	3540	1503	462	729	1017	ND
Wheat bran	138	882	672	868	410	2787	ND
Oat bran	567	297	766	90	440	712	ND
Barley bran	71	140	133	42	42	541	ND

Altogether, wheat grains may contain up to 1 mg lignans/100 g and the large majority of these reside in the bran layer containing up to 20 mg lignans /100 g bran. 7-Hydroxymatairesinol and syringaresinol are the dominant lignans in wheat although secoisolariciresinol, lariciresinol, matairesinol and other less characterized lignans, such as 7-oxomatairesinol are also present in variable concentrations depending on the variety [33, 34] (Table 2).

Table 2: Concentration of total lignans in wheat bran and rye bran.

Wheat bran	Amounts of total lignan 5-20 mg/100 g (<i>depending on the variety</i>)
Total lignan	7-hydroxymatairesinol, syringaresinol, secoisolariciresinol, lariciresinol, matairesinol
New lignan	Iso-hydroxymatairesinol, 7-oxomatairesinol
Rye bran	Amounts of total lignan 10-20 mg/100 g (<i>depending on the variety</i>)
Total lignan	syringaresinol, pinoresinol, lariciresinol, isolariciresinol, 7-hydroxymatairesinol, matairesinol, secoisolariciresinol.

Different varieties of wheat have been profiled for their content of lignans. In a work by Dinelli *et al.* [35], old and modern Italian wheat varieties cropped in the same location and growing season have been analyzed. The study shows that both free and bound phenols are very variable in content and, on the average, the bound fraction contributed 72.0% to the total phenolic content. The obtained metabolomic fingerprints showed the presence of several classes of compounds which included flavonoids, coumarins, stilbenes, proanthocyanidins and lignans, some of which may have health promoting effects. The HPLC-ESI-TOF-MS analysis identified 104 compounds and also highlighted remarkable differences in the phytochemical fingerprints of old and modern wheat varieties. Six ancient wheat genotypes (Bianco Nostrale, Frassineto, Gentil Rosso, Gentil Rosso Mutico, Marzuolo d'Aqui, Verna) showed phenolic profiles with a number of total compounds and isomer forms including the lignan syringaresinol, with much higher contents than in the modern cultivars, indicating that they might be better sources of phenolic compounds with added value in terms of potential health-promoting components [35].

Rye: Among cereal grains, rye has the highest concentration of lignans, mostly as glycosides which, like in wheat, are concentrated in the outer layers of the kernel [36]. The most abundant are pinoresinol, syringaresinol, 7-hydroxymatairesinol, lariciresinol and isolariciresinol. When considering the whole rye grain, 7-hydroxymatairesinol is the dominant lignin, followed by syringaresinol (Table 2).

In studies aimed at identifying novel urinary biomarkers of wholegrain rye bread intake versus refined wheat bread, the profiling of several metabolites, including END and ENL, allowed the identification of wholegrain rye as a food source providing a higher concentration of urinary phenols [37]. In their study, Bondia-Pons *et al.* experimentally found higher levels of five lignans after intake of wholegrain rye bread than after intake of refined white bread. This was associated with a lower fasting plasma insulin level and lower 24-h urinary C-peptide excretion [37]. The highest concentration of plasma and urinary levels of entrolignans have

been measured in subjects traditionally consuming wholegrain rye at the highest amounts in North and East Europe, Russia and China, where the production of rye is mostly concentrated (1 to 4 million tons/year).

Cereal consumption and bioprinciples of accessibility:

Regarding the nutritional aspect of cereal grains and bran, studies suggest that whole grains have superior nutritional values compared with foods enriched with fibers. This is because the bioactive components are more abundant in the whole grain and particularly concentrated in the aleurone layer. Moreover, beyond lignans, the bran fraction of cereals contains most of the micronutrients and complex carbohydrates of the grains, in particular beta-glucans and arabinoylans, which are promising nutraceuticals. Nonetheless, as recently evidenced by an EU study consortia on European cohorts, the consumption of whole seeds or grains supposed to provide health benefits, often does not fulfil the expectations (www.healthgrain.eu). This might be due to the fact that although cereals are good sources of lignans, their bioabsorption from the consumption of the whole grains is limited. This is likely due to their macro- and microstructure and consequently texture of the raw flour which influences their functionality in the gastrointestinal tract, specifically component absorption, bioavailability, and metabolism to enterolignans [38]. The studies carried out by two groups in the Healthgrain consortium working on wholegrain products, including wheat, showed in fact that subjects consuming whole grains did not achieve significant changes in traditional biomarkers of MetS (www.healthgrain.eu). It is now known that this is partially due to the inaccessibility of the bioactive principles contained in the natural matrices because of their insufficient breakdown and digestion but also absorption. The major problem identified by bioavailability studies was the poor bioaccessibility of the active principles. Two studies from Anson *et al.* [39, 40], evidenced that release and conversion of phenolic acids to their microbial metabolites was enhanced by bioprocessing of wheat bran to more bioavailable formulations. In the bioprocessed bread, the bran was processed by yeast fermentation combined with enzyme treatment, consisting of cell-wall-degrading enzymes, mainly xylanase, cellulase, glucanase and feruloyl-esterase [40].

Nutritional impact on health of lignans. The metabolic syndrome (MetS): The high prevalence of MetS and excessive adipose tissue accumulation (in particular, abdominal obesity) is a major threat to public health, being associated with a substantial decrease in health-related quality of life and an increase in economic costs due to disease treatment, as well as lower social productivity.

MetS is a multiplex and life-based risk factor that consists of several correlated risks of metabolic origin. In addition to dyslipidemia, hypertension and hyperglycemia, the syndrome carries a proinflammatory state [41]. Subjects affected by MetS are at increased risk of cardiovascular diseases (2-fold), type-2 diabetes (5-fold), gallbladder disease, certain types of cancer and psychosocial problems (www.lipgene.tcd.ie,

www.nutrition.org.uk/lipgene, www.oecd.org) [42]. More than 130 million Europeans are estimated to have MetS and 56 million type-2 diabetes [43] and the numbers are estimated to increase for the next decades. Thus, MetS is ‘the’ major health challenge in Europe and the western countries today but further, the same tendency is also seen in many less developed countries, e.g. India. New strategies and very extensive research and initiatives related to the long-term prevention and reduction of incidence and severity of cardiovascular diseases, type-2 diabetes and obesity have been undertaken by the EU Commission and the WHO for both prevention and cure, to halt the problem [44-46]. Amongst the identification of fields of intervention, the better characterization of the bioactive compounds within many foods seems to be a promising area. In fact, although poor diets and physical inactivity are the driving forces behind MetS, appropriate dietary strategies may positively impact on disease predisposition and evolution.

For the assessment of the nutritional effects of lignan consumption on human health, different European networks such as the EPIC-Potsdam Study have been particularly suited to serve as epidemiological reference cohorts for results comparison. They have been recently used to investigate the role of lignans on disease occurrence using medically verified data. The analyzed endpoints of chronic diseases following lignan intake were: a) incidence of the metabolic syndrome (NHANES-National Health and Nutrition Examination Survey [47]; b) type-2 diabetes (EPIC-InterAct Study) [9]; c) cardiometabolic risk factors (NHANES) [11]; d) specific types of cancers (DietCompLyf Study) [12]; (EPIC Study) [10, 48-51], (EPIC-Norfolk) [52-54]; and e) overall mortality (EPIC-Spain) [10]. From these studies it is apparent that European subpopulations in which the major source of lignans is cereals [9], display considerable lower disease frequency and beneficial effects in the control of metabolic diseases like dyslipidemia and type-2 diabetes. Further, other recent observations in different populations report that lignan intake is associated with lower levels of triglycerides, lower fasting glucose, and fasting insulin serum levels in males aged 20-60 years. These data are associated with higher HDL-cholesterol levels [2, 11, 29, 55-58]. Some studies have also been conducted in animal models in which partially purified lignan fractions have been shown to improve blood lipids and glycemic control [59, 60]. The results from all these studies underline the need to understand better the mechanisms of action of these compounds in maintaining a healthy metabolism.

Mechanisms of lignans on health: Although impressive advances have been seen in the last years as regards the mechanisms of action for the most thoroughly investigated dietary polyphenols (lignans, stilbenes, isoflavones, etc.), most of their health-related effects have been attributed to their activity as antioxidants [61, 62]. However, a certain amount of data is now becoming available on the activity of these compounds at doses lower than those required to exert biological antioxidant activities, but sufficient to activate fundamental cell pathways, several of which are regulated by the nuclear receptors (NRs). The availability of data regarding the regulation of different NRs at relatively low doses, stimulated the characterization of the effects of lignans in tissues and cells at doses that are consistent with the doses reached in the blood through the diet (nutritional doses) [7, 18]. Among others, it has been shown that, at nutritional doses, some lignans can regulate the activity of specific NRs such as the estrogen receptors (ERs) with an efficiency that in some cases approximate to that of the endogenous ligand 17 β -estradiol [18]. This highlighted the importance of their actions as hormone mimics suggesting that they are effective on several physiological functions. Based on these observations, molecular and cellular studies are now starting to support the data provided by

epidemiological studies and dietary interventions. Significant effects have in fact been shown for regulatory mechanisms related to lipid metabolism, glucose homeostasis, cholesterol biosynthesis and insulin biosynthesis and secretion, [18, 60, 61, 63-65].

Several data on lignans as estrogenic compounds regulating ER-dependent pathways have been published [19, 66]. However, recent progress in the area indicate that the term “phytoestrogen” may be poorly representative of the complex activities of lignans when considering as functional targets metabolic functions regulated through the ERs in the liver or adipose tissues, and not representative at all when considering lignan activity through metabolic sensors like the peroxisome proliferator-activated receptors. It is in fact becoming frequent to observe that the same molecules that regulate the ERs, also regulate other NRs that play fundamental actions in metabolic tissues. The engagement and regulation of PPAR α , PPAR γ , and LXR α that are central regulators of glucose metabolism, fatty acid metabolism and insulin action, evidence the lignans as real SNRMs (Selective Nuclear Receptors Modulators). Known examples are the flaxseed lignan secoisolariciresinol diglucoside (SDG), which can regulate the ERs [6], PPARs [67] and LXR α [17], and unpublished observations. A lignan isolated from medicinal plants, honokiol [68], is a non-adipogenic PPAR γ agonist [69] and a ligand of the adipogenic RXR β receptor [70]. Meso-dihydroguaiaretic acid (MDGA), a major component of *Myristica fragrans*, a traditionally used food spice in several countries [71], competes with the LXR α agonist T0901317 for binding to the LXR α domain [17]. The sesame lignans sesamin and episesamin, can stimulate fatty acid synthesis and oxidation in the liver via activation of PPAR α , can attenuate nutritional fibrosing steatohepatitis through up-regulation of PPAR γ [72], down-regulate PPAR γ antiinflammatory effects in 3T3-L1 preadipocytes [16], and can suppress macrophage-derived chemokine expression in human monocytes via epigenetic ER- and PPAR α -mediated regulation [73]. The mammalian enterolignans, enterodiol (ENL) and enterolactone (END) generated through the metabolism of dietary lignans by the intestinal microbiota bind both the ERs [18] and likely the LXR α , as evidenced by unpublished *in silico* studies (personal communication from Ingemar Pongratz). Activity has also been reported for (+)-pinoresinol through the negative regulation of the human thyroid hormone receptor β [74], likely interfering with adipose cell differentiation [75]. Negative regulation has also been shown by ENL through the androgen receptor (AR) and by sesamin through the pregnane X receptor (PXR), while secoisolariciresinol and ENL are able to activate PXR in reporter assays suggesting an effect in the metabolism of CYP3A substrates [76]. A few cereal lignans have also been tested *in vitro* and found to be effective inhibitors of adipocyte differentiation. Pinoresinol, cycloisolariciresinol, lariciresinol, secoisolariciresinol in the concentration range of 1-1000 nM, are effective in inhibiting 3T3-L1 cell differentiation [77] and although their fine mechanism of action has still to be clarified, it is likely that it involves the ability to regulate nuclear receptors such as PPARs, ERs and LXR α s [65, 78, 79].

Microbial metabolism of lignans: Lignans occur in food in the form of esters, glycoconjugates, and polymers. These compounds are not directly bioavailable and require metabolic steps which involve both mammalian (glucuronidation and to a lesser degree, sulfation) and gut microbial processes [80]. When lignan glycosides are consumed with the diet, they are released from the matrix after mastication, are deglycosylated and are converted in the liver to glucuronidated [81], methylated and sulfated derivatives [82] that can be identified in human urine [83], but also to enterodiol and enterolactone by the intestinal microbiota [20] (Figure 4).

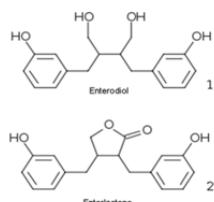


Figure 4: Mammalian lignans. The enterolignans (or mammalian lignan) enterodiol (END) and enterolactone (ENL) are lignan metabolites that are produced in the intestine from plant lignans by the gut microbiota.

Probiotic lactobacilli and bifidobacteria, which have evolved within the colonic ecosystem where indigestible oligo- and polysaccharides are their sole carbon sources, bear several glycosyl-hydrolases and can contribute to the release of the aglycones from glycoconjugated phytochemicals [80]. β -Glucosidase-positive probiotic bacteria were proved to release the aglycones of lignans *in vitro* [84]. Despite the structural diversity of food lignans and their very different plant origins (i.e. oily matrices versus cereal bran), lignans are converted to enterolignans by the gut microbiota [20]. Following metabolism, enterodiol and enterolactone may reach a level high enough for signal pathway regulation in the liver, lung, kidney, heart and brain tissues [83] and pharmacokinetic studies show that these microbial metabolites have high systemic clearance and short half-lives in the blood circulation, before being excreted as hepatic conjugates with the urine [85]. Dietary consumption of probiotics and limited digestible or indigestible food constituents such as oligosaccharides (prebiotics) and polyphenols or both (synbiotics), regulate the numbers and types of microbes which can influence energy utilization from the diet and regulate energy expenditure and storage. However, the genera of the microbiota that contribute to the metabolism of lignans are not yet sufficiently characterized, although investigations are underway and some gut microbiota strands have been identified as major lignan metabolizers in humans [86] and in animals [4, 87] and may thus represent possible beneficial factors for health. Associations are starting to be reported between the composition of the intestinal microbiota, lignan metabolites and the risk of diseases. Recent results suggest that urinary levels of ENL and/or END are associated with a lower risk

of type 2-diabetes in US women [5], inflammation [88] and obesity and metabolic alterations in US men [55], and to the inhibition of inflammation and pro-carcinogenic microenvironments in animal tissues [66].

Conclusions: Today, more than 20 different lignans have been identified in seeds and cereals consumed as food worldwide and many others are found in cosmetic and medicinal plants [89]. Many of these have not been studied before and may possess potent biological activity, but also, amongst the most studied ones, data are not sufficient to draw definitive considerations for the present. However, the health effects of the most abundant food lignans in ameliorating different chronic diseases and related co-morbidities are being extensively characterized regarding absorption, bioavailability and activity following intake of whole foods, enriched foods, and fractions and extracts. Although so far most of these studies deal mainly with oilseeds lignans, an increasing number of observations indicate that regular intakes of wholegrain cereals and their products may exert a level of protection not observed with the ingestion of refined cereals, and recent research suggests that part of these effects may well be contributed by the lignan components [90].

To conclude, the reviewed body of research suggests that given additional efforts in clinical trials, it may be possible to make 'function' claims for lignans related to maintaining a healthy metabolism and the prevention/amelioration of the metabolic syndrome and related pathologies. Importantly, this effect may be elicited through mechanisms that involve their direct agonistic/antagonistic action with different types of nuclear receptors.

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The Rare Amino Acid Building Block 3-(3-furyl)-Alanine in the Formation of Non-ribosomal Peptides

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Microorganisms have made considerable contributions to the production of peptide secondary metabolites, many of them with therapeutic potential, e.g., the fungus-derived immunosuppressant cyclosporine A and the antibiotic daptomycin originating from *Streptomyces*. Most of the medically used peptides are the product of non-ribosomal peptide synthetases (NRPS), incorporating apart from proteinogenic also unique, non-proteinogenic amino acids into the peptides. An extremely rare such amino acid is 3-(3-furyl)-alanine. So far, only few peptides have been found that contain this residue, including the rhizonins, bingchamide B and endolides. The producer of the rhizonins was proven to be the bacterial endosymbiont *Burkholderia endofungorum* inside the fungus *Rhizopus microsporus*. The microbial origin, chemistry and bioactivity of the 3-(3-furyl)-alanine containing peptides are the focus of this review.

Keywords: Peptides, NRPS, Furyl-alanine, Endolide.

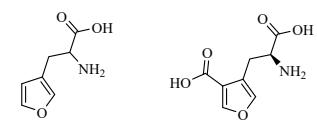
Among the huge class of secondary metabolites, peptides are a restricted group of structurally diverse compounds produced by prokaryotes like bacteria, and eukaryotes, e.g. fungi [1]. They are often essential for the survival and growth of the microbial producers in the rivalry of their natural habitat, acting as siderophores, toxins or signaling molecules [2]. Many of these peptides have important medical applications as antibiotics, e.g., daptomycin, and immunosuppressants, e.g., cyclosporine A [3]. In terms of biosynthesis, some of these peptides originate from ribosome-derived precursor molecules and gain their final structure by post-translational modifications, e.g., microcins, microviridins and polytheonamides [4]. A second group of these peptides, including most of the medically used ones, are produced by non-ribosomal peptide synthetases (NRPS), which function as multi-modular enzymes connecting small building blocks, predominantly amino acids, but in some cases also carboxy acids [5]. The latter group often contains unique, non-proteinogenic amino acids, such as hydroxyphenylglycines in vancomycin and (4R)-4-[(E)-2-but enyl]-4-methyl-L-threonine in cyclosporine A [2]. Over 500 different amino acids are known to date, either found as partial structures in peptides, as free low molecular weight natural products, or are metabolic intermediates [6]. 3-(3-Furyl)-alanine (**1**) is such a non-proteinogenic, heterocyclic amino acid incorporated in non-ribosomal peptides, but to date not encountered free in nature (Figure 1). In contrast, the 3-(3-furyl)-alanine carboxy derivative, L-3-(3-carboxyfuran-4-yl)-alanine (**2**) was isolated from the fruiting bodies of the gilled non-edible mushrooms *Phyllotopsis nidulans* and *Tricholomopsis rutilans* (Figure 1) [7,8]. For this component no biological activity has been described to date. Chemically synthesized L-3-(3-furyl)- and L-3-(2-furyl)-alanine, however, exhibit growth inhibitory activity towards both bacteria and fungi [9], making them potentially useful agrochemical fungicides, industrial microbicides and wood preservatives [10].

3-(3-Furyl)-alanine as a building block in bioactive natural products: Even though 3-(3-furyl)-alanine is so far not known as an unbound amino acid, fungi and bacteria incorporate this residue into peptide structures with diverse bioactivities (Figure 2, Table 1).

Table 1: Peptides incorporating 3-(3-furyl)-alanine.

Source	Peptide	Activity
<i>Burkholderia endofungorum</i> (bacterial)	Rhizonin A, B (3,4)	Hepatotoxic effects in rats [15] and plant growth promoting activity [17]
<i>Streptomyces bingchengensis</i> (bacterial)	Bingchamide B (5)	<i>In vitro</i> cytotoxic activity towards human colon cancer cell line HCT-116 ($IC_{50}= 18 \mu\text{M}$) [18]
<i>Stachyliodium</i> sp. (fungal)	Endolide A (6)	Affinity to the vasopressin receptor 1A ($K_i = 7.04 \mu\text{M}$) [19]
	Endolide B (7)	Affinity to the serotonin receptor subtype 5HT _{2b} ($K_i = 0.77 \mu\text{M}$) [19]

The first described naturally occurring peptides containing 3-(3-furyl)-alanine were rhizonin A (**3**) and B (**4**), at that time isolated from *Rhizopus microsporus*, a fungus used to prepare fermented food [11]. Interestingly, the two 3-(3-furyl)-alanine units in rhizonin A possess opposite absolute configurations as determined from X-ray crystallography. The rhizonins were regarded as the first mycotoxins isolated from fungi (Zygomycota), with most known mycotoxin producing strains belong to the fungal phyla Ascomycota or Basidiomycota [12]. It was later found that the true producer of rhizonins is in fact the bacterial endosymbiont *Burkholderia endofungorum*, which resides inside the fungal cytosol [13]. This was also true for the polyketide-derived antimitotic macrolide rhizoxin, initially believed to be a product of *R. microsporus* but later discovered to be produced by *Burkholderia rhizoxinica* [14].



3-(3-furyl)-alanine (**1**) L-3-(3-carboxyfuran-4-yl)-alanine (**2**)

Figure 1: 3-(3-Furyl)-alanine and its carboxy-derivative.

Rhizonin A exhibited potent hepatotoxic effects in rats [15], and cytotoxicity, which was mainly ascribed to its 3-(3-furyl)-alanine partial structure. Synthetic replacement of this moiety with phenylalanine gave analogues **8** and **9** with reduced cytotoxicity and

improved *in vitro* fat-accumulation inhibitory activity in 3T3-L1 murine adipocytes [16]. Controversially, rhizonin A showed plant growth promoting activity in lettuce seed germination, and thus, the rhizobacterial strain *Burkholderia* sp. KCTC11096 is to be explored for supporting sustainable agricultural production [17].

The second report of a 3-(3-furyl)-alanine building block is related to the cyclic pentapeptide, bingchamide B (**5**), isolated from a soil-dwelling strain of *Streptomyces bingchengensis* [18]. The absolute configuration of the amino acid building blocks of bingchamide B, including 3-(3-furyl)-alanine, was not determined. Bingchamides exhibited *in vitro* cytotoxic activity towards human colon cancer cell line HCT-116 with an IC_{50} value of $18 \mu\text{g mL}^{-1}$ for bingchamide B and an IC_{50} value of $14.1 \mu\text{g mL}^{-1}$ for its natural analogue bingchamide A (**10**), featuring a phenylalanine residue in place of the 3-(3-furyl)-alanine [18].

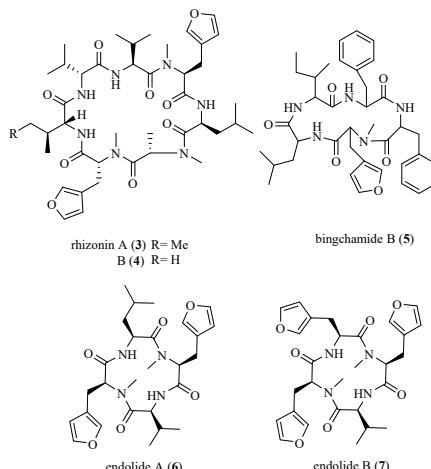


Figure 2: Peptides featuring a 3-(3-furyl)-alanine amino acid residue.

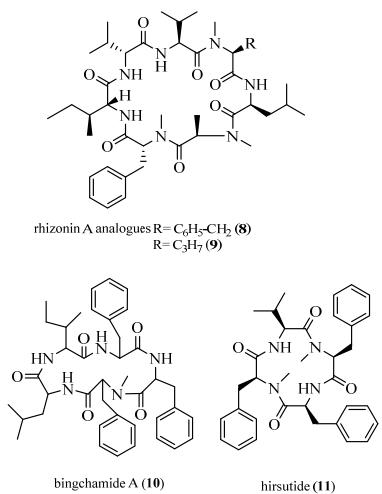


Figure 3: Bioactive peptide analogues featuring phenylalanine amino acid residues.

Recently, the novel tetrapeptides endolide A (**6**) and B (**7**) were isolated from a solid culture of the marine sponge-derived fungus *Stachyliidium* sp. [19]. Here, the two 3-(3-furyl)-alanine units in endolide A were assigned the L-configuration as determined from X-ray crystallography and Marfey's analysis. Hitherto the endolides were regarded as the first fungal peptides containing 3-(3-furyl)-alanine. Whether the endolides are presumably produced by an endofungal bacterium, as in the case of rhizonins, could not be unequivocally proven, as it was not yet possible to isolate and culture axenic bacteria from the *Stachyliidium* sp. fungus [19].

Unlike the other 3-(3-furyl)-alanine containing peptides rhizonin A and bingchamide B, the endolides did not exhibit any *in vitro* cytotoxicity. Interestingly, they have affinity to G protein-coupled receptors (GPCRs) in radioligand binding assays. Endolide A showed affinity to the vasopressin receptor 1A with a K_i of $7.04 \mu\text{M}$, while endolide B had selective affinity to the serotonin receptor subtype 5HT_{2b} with a K_i of $0.77 \mu\text{M}$ [19].

A structural analogue of the endolides, hirsutide (**11**), was isolated from the entomopathogenic fungus *Hirsutella* sp. Hirsutide features phenylalanine residues instead of the 3-(3-furyl)-alanine residues in endolide B [20]. Initially no bioactivities were reported for hirsutide. However, its recent chemical synthesis and detailed pharmacological screening revealed potent antibacterial activity against the Gram-negative bacteria *Pseudomonas aeruginosa* and *Klebsiella pneumoniae*, with additional anthelmintic activity and minor cytotoxic effects against Dalton's lymphoma ascites and Ehrlich's ascites carcinoma cell lines with IC_{50} values of 14 and $22 \mu\text{M}$, respectively [21].

Putative biosynthesis of peptides and 3-(3-furyl)-alanine: Peptide secondary metabolites are assembled either through the ribosomal machinery, *i.e.*, through direct gene translation [4], or by multi-modular non-ribosomal peptide synthetases (NRPS) [2]. In the translation system, decoding of the trinucleotide sequence (codon) information stored in the messenger RNA determines the peptide amino acid sequence. Here each codon assigns a single amino acid from the known pool of 20 L-configured proteinogenic amino acids [22].

On the other hand, NRPS assembly lines are divided into modules termed initiation and elongation modules, where each module is responsible for integrating a dedicated substrate into the growing peptide chain. This is done with the help of catalytic domains, a sub-division of each module. Fundamental domains are indispensable for the selection and activation of the building blocks (adenylation, A-domain), the tethering of the product to the enzyme (peptidyl carrier protein, PCP-domain) and peptide bond formation (condensation, C-domain). Structural diversity arises due to the presence of auxiliary domains imparting unique structural features to the non-ribosomal peptides. These include epimerization (E) domains for incorporating D-amino acids, methylation (Mt) domains for introducing methyl groups and oxidation (Ox) and reduction (R) domains for the formation of heterocyclic rings. Finally, a terminal thioesterase (TE) domain catalyzes peptide release and at times cyclization [23].

A closer look into the structural architecture of the rhizonins, bingchamides and endolides proposes a non-ribosomal biosynthetic origin. This is implied due to the incorporated amino acid residues, *i.e.*, N-methylated, sometimes D-configured, and structurally unusual non-proteinogenic in nature, *i.e.*, 3-(3-furyl)-alanine. For the rhizonins, total DNA isolated from the cultured *B. endofungorum* afforded NRPS gene fragments. However, the exact NRPS genes involved in the biosynthesis of rhizonins were not yet revealed and further studies are underway [13]. Analysis of the *S. bingchengensis* genome yielded gene clusters postulated to encode NRPSs. These include modules with adenylation domain substrate specificities, correlating with the structures of bingchamides [24]. However, no proof for a responsible biosynthetic gene cluster is currently available.

In NRP biosynthesis, not only the multi-modular NRPS assembly line is of importance, but also the formation of the building blocks. Often the respective enzymes are encoded as part of or adjacent to

the biosynthetic gene cluster [25,26]. So far, no biosynthetic studies were conducted to reveal the biosynthetic origin of 3-(3-furyl)-alanine, apart from it being considered a potential substrate for the NRPS biosynthetic machinery. Taking known biosynthetic pathways for similar furans into consideration, several pathways for 3-(3-furyl)-alanine may be proposed here.

A furan ring as such is encountered in many secondary metabolites. In terpenoids, furan ring biosynthesis utilizes active isoprene units, *i.e.*, isopentenyl diphosphate (IPP) and dimethylallyl diphosphate (DMAPP), as seen in the diterpenoids divinatorins (**12**) and bibifuran (**13**) [27,28]. The signaling molecules methylenomycin furans (MMFs) (**14**) and the furanylcarbonyl containing acyl rhamnosides (**15**) are formed by a mixed acetate-glycerol pathway [29-32]. The latter involves the condensation of a dihydroxyacetone-derived three-carbon unit originating from glycerol and a β -ketothioester intermediate derived from a polyketide synthase or fatty acid metabolism (Figure 5). In contrast, the hydroxyfuranone (**16**) and methoxyfuranone (**17**), volatile strawberry aroma compounds, are obtained from carbohydrate metabolism, *i.e.*, from D-fructose [33].

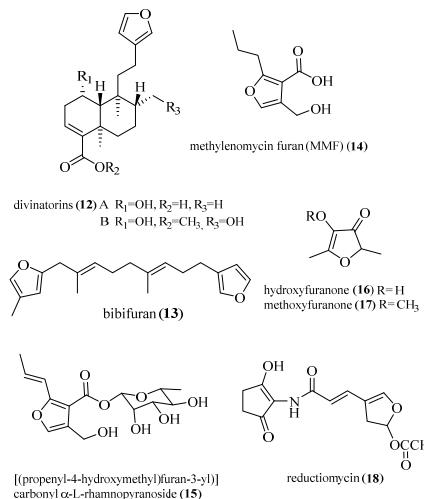


Figure 4: Selected metabolites featuring a furan moiety.

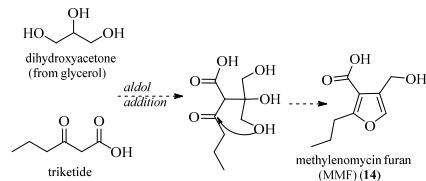


Figure 5: Biosynthesis of methylenomycin furan through a mixed acetate-glycerol pathway, adopted from literature [27,30].

However, the association of the furan skeleton with an amino acid, *i.e.*, alanine, implies other precursor molecules to be involved in its biosynthesis. 3-(3-Furyl)-alanine is an analog of the shikimate-derived aromatic amino acid phenylalanine [34]. Thus, the structural relationship between the 3-(3-furyl)-alanine and the

phenylalanine skeleton, may point to the possibility of furyl-alanine being formed by ring cleavage of a phenylalanine intermediate. Such a ring cleavage is known for several metabolites such as the mycotoxin patulin [35], the betalain plant pigments [36], and pyrrolobenzodiazepine anti-tumor antibiotics [37]. Feeding of isotopically labeled phenylalanine to the respective microbial cultures could clarify this presumption.

On the other side however, the unique dihydrofuranacrylic acid moiety of the antibiotic reductionmycin (**18**), a *Streptomyces* metabolite, also shares the basic carbon skeleton with 3-(3-furyl)-alanine [38]. In this case, labeling studies showed that the dihydrofuranacrylic acid moiety is derived from a symmetrical product of the shikimate pathway, *i.e.*, *p*-hydroxybenzoic acid, by aromatic ring cleavage (Figure 6) [39].

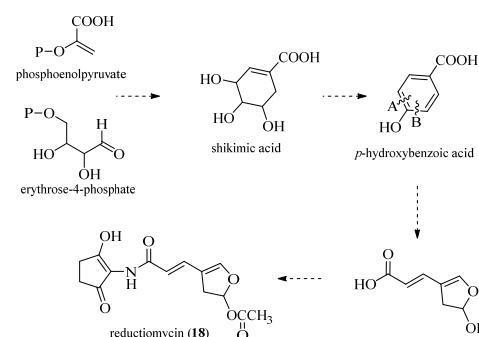


Figure 6: Biosynthesis of the dihydrofuranacrylic acid moiety of reductionmycin via the shikimate pathway, adopted from literature [38,39]. Cleavage occurs either at site A, or alternatively at B, resulting in different isotope labeling patterns (not shown here).

Which metabolic pathway underlies the biosynthesis of the unique 3-(3-furyl)-alanine moiety in the peptides, and whether all the peptides incorporating such a moiety have a similar biosynthetic origin is yet to be revealed.

Conclusion: Microorganisms have made considerable contributions to the production of peptide secondary metabolites, many of them with therapeutic potential, *e.g.*, the immunosuppressant cyclosporine A [40] and the antibiotic daptomycin [41]. Indeed, peptides that occur in nature as secondary metabolites share structural features like a cyclic architecture, *N*-methylation of amide bonds, and incorporation of atypical building blocks, such as D-amino acids, carboxy acids and non-proteinogenic amino acids. Such features bestow several attractive attributes in terms of receptor affinity and selectivity and improved proteolytic stability, making them interesting drug scaffolds [3]. With the ongoing interest in peptide-based drugs [42], 3-(3-furyl)-alanine containing peptides add an interesting structural variation to this group of natural products. However, a deeper understanding of their biosynthesis and bioactivities should be sought, in order to be implemented in the drug-design process.

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