








Biotransformation of Perrottetin F by *Aspergillus niger*: New Bioactive Secondary Metabolites

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Abstract: Biotransformation of bis-bibenzyl perrottetin F (1), isolated from the liverwort *Lunularia cruciata* by *Aspergillus niger*, has been investigated. New metabolites (2-4) have been isolated using reversed phase semipreparative HPLC and their structures were established to be 8-hydroxyperrottetin F, C-7-C-8 cleaved product, and perrottetin F 6'-sulfate using 1D and 2D NMR, HR-ESI-MS, IR and UV spectroscopy. The antimicrobial and cytotoxic properties of these compounds were also evaluated. Given the suggested cytotoxic properties of the parent compound, antiproliferative activity against healthy human lung fibroblasts (MRC5) and human lung carcinoma (A549) of three metabolites were evaluated revealing their lower cytotoxic properties in comparison to the starting compound - perrottetin F. The antimicrobial properties of these compounds were also evaluated, with the inhibitory activity against the *Pseudomonas aeruginosa* PAO1 and *Staphylococcus aureus* determined between 100 µM and 450 µM. The metabolites showed remarkable ability to inhibit synthesis of bacterial quorum-sensing signal molecules such as short chain acyl homoserine lactones (AHLs). Therefore, biotransformation method represents fast and effective tool for obtaining new bioactive structures.

Keywords: Biotransformation; perrottetin F; liverworts; *Aspergillus niger*; cytotoxic activity; 1D and 2D NMR.
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1. Introduction

The Marchantiophyta (liverworts) produce various aromatic compounds among them bis-bibenzyls, such marchantin, riccardin and perrottetin series which are very specific and rare natural products found only in this division. Up to date, 103 cyclic and acyclic bis-bibenzyls have been isolated from the liverworts [1-5]. Bis-bibenzyl skeleton comprises four aromatic rings (A-D),

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biosynthesized from dimerization of lunularic acid through dihydrocoumaric acid and prelunularin [5]. Perrottetin F (**1**) has been found in the thalloid liverworts *Radula perrottetii*, *R. kojana* and *Lunularia cruciata* [4,6]. This acyclic bis-bibenzyl was obtained as the major component from the methanol extract of *L. cruciata* [7]. Bis-bibenzyls possess wide range of biological activities such as cytotoxicity, 5-lipoxygenase and NO production inhibitory activity, antitumor, farnesoid X receptor (FXR)-activating activity, antimicrobial and many others [8,9]. Anti-influenza PA endonuclease potential of perrottetin F is described previously [10].

Known as 'green technology', biotransformation is used for converting various compounds into derivatives with possible biological activities [11]. Microbiological transformation of natural compounds can be useful for pharmaceutical drug development, in order to obtain more active compounds or to increase the yields of the pharmaceutical active components. It is an important strategy in the structural modification of natural compounds by using microorganisms, animal tissues or enzymes with great potential to produce novel, more efficient compounds [12]. Compounds obtained by biotransformation may improve properties in comparison to the starting component. It may also be good alternative to chemical synthesis of new drugs [13]. Literature data mainly show biotransformation of phytosterols, steroids, terpenes, alkaloids, and flavonoids [14]. It was noted that bacteria made complete steroid structure degradation while, micromycetes were active in multiple steroid hydroxylations. Some diterpenes (stemodin, stemodinone and stemarin) were biotransformed by *A. niger* to give three hydroxylated compounds [15]. Sesquiterpene cinnamodial was biotransformed to lactonized products [16].

In this study, an easy and cheap method for obtaining bioactive bis-bibenzyl derivatives using micromycete *A. niger* is presented. As per authors best knowledge there is only one report on biotransformation of bis-bibenzyl [9] and this is the first investigation on perrottetin F (**1**) biotransformation, from which three new derivatives were obtained, and their biological activities were investigated.

2. Materials and Methods

2.1. General Experimental Procedures

Optical rotations were measured on a Rudolph Research Analytical Autopol IV automatic polarimeter with methanol as solvent and concentration of compounds 1.00 mg/mL. UV spectra were recorded on a GBC Cintra UV/VIS spectrometer with methanol as solvent in concentration range $3\text{--}9 \times 10^{-5}$ M. IR spectra were obtained on a Thermoscientific Nicolet 6700 FT-IR spectrometer. NMR spectra were measured on a Bruker Avance III 500 spectrometer at 500.26 MHz for ^1H and 125.80 MHz for ^{13}C , with CD_3OD as a solvent. HR-ESI-MS data were obtained on Agilent 6210 Time-of-Flight LC/MS system equipped with an ESI interface (ESITOFMS). Solvent was methanol, mobile phase was: 0.2% $\text{HCOOH}/\text{CH}_3\text{CN}$ 1:1, 0.2 mL/min. The ESI was operated in a negative mode and nitrogen was used as the drying gas (12 L/min) and nebulizing gas at 350 °C (45 psi). The OCT RF voltage was set to 250 V and the capillary voltage was set to 4.0 kV. The voltages applied to the fragmentor and skimmer were 140 V and 60 V, respectively. Scanning was performed from 100 to 1500 m/z (mass-to-charge ratio). For column chromatography (CC) silica gel 60 (SiO_2 ; 0.063 – 0.200 mm, Merck) was used. Analytical TLC was carried out on silica gel 60 GF254 20 × 20 cm plates, with a layer thickness of 0.25 mm (Merck). Preparative TLC was carried out on silica gel 60 GF254 (SiO_2 ; <0.063 mm, Merck) packed 20 × 20 cm plates, thickness 0.75 mm. Semipreparative HPLC separation was performed on an Agilent Instrument 1100 series equipped with a DAD (Agilent Technologies, G1315C). The column used was a Zorbax Eclipse XDB C18 (9.4 mm × 250 mm, 5 μm). Solvents for HPLC separation were of chromatographic grade. Solvents for CC were freshly distilled.

2.2. Plant material, Extraction and Isolation

A voucher specimen (No. 17290) has been deposited in the Herbarium at the Institute of Botany and Botanical Garden "Jevremovac", University of Belgrade (BEOU). Extraction and

fractionation of crude methanol extract of liverwort *Lunularia cruciata* (L.) Dumort. ex Lindb. (Lunulariaceae) by silica gel CC and isolation of dominant compound perrottetin F have been done according to previously described procedure [7].

2.3. *Aspergillus niger* TBUYN-2 Growth and Biotransformation Procedure

Micromycete *Aspergillus niger* TBUYN-2 obtained from the soil collected in Osaka prefecture [9] was grown in Czapek-peptone medium (pH 7) in incubator shaker at 120 rpm, 25 °C for 3 days. When micromycete achieved full growth, perrottetin F (**1**) (50 mg was dissolved in 0.5 mL of ethanol) was added into three flasks (150 mg in total) and left in thermoshaker at 120 rpm, 25 °C for 4 days. EtOH (0.5 mL) was added in the flask with *A. niger* and medium, as control.

After 4 days, medium and mycelia were separated in a Buchner separation funnel. For extraction of compounds from media EtOAc was used, while mycelia were left in acetone for next 7 days. pH of media was 3.5. To get basic-neutral fraction (BNF) from media and mycelia (pH 9) 50% Na₂CO₃ was added. Acidic fraction (AF) (pH 2-3) was obtained by adding 5% H₂SO₄ solution in media. After extraction analytical TLC was carried out on aluminium silica sheet gel 60F₂₅₄ (MERCK). Chromatograms were developed using CH₂Cl₂: EtOH (7:1 v/v) as the developing solvent. The presence of biotransformed compounds were detected under UV light (254 nm) and spraying the 30% H₂SO₄ solution on TLC sheet. The biotransformation products were separated using semi-preparative HPLC.

2.4. Cytotoxicity Assay

Cytotoxicity of compounds (**1-4**) was measured using the methods described previously [17]. A549 (human lung carcinoma, obtained from ATCC) and MRC5 cells (human lung fibroblast, obtained from ATCC) were plated in a 96-well flat-bottom plate at a concentration of 1×10^4 cells per well, grown in humidified atmosphere of 95% air and 5% CO₂ at 37 °C, and maintained as monolayer cultures in RPMI-1640 medium supplemented with 100 µg/mL streptomycin, 100 U/mL penicillin, and 10% (v/v) fetal bovine serum (FBS). After 24 h of A549 and MRC5 cells incubation, the media containing increasing concentrations of each tested compound (with the highest concentration tested being 500 µM) were added to the cells. Stock solutions of bis-benzyls were prepared in DMSO (50 mM). Control cultures received the solvent DMSO and blank wells contained 200 µL of growth medium. After 48 h of incubation, cells proliferation was determined using 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) reduction assay. MTT solution (0.5 mg/mL) was added after treatment and incubated for 1 h, followed by dimethyl sulfoxide incubation. Cell proliferation was determined from the absorbance (A) at 540 nm on Tecan Infinite 200 Pro multiplate reader (Tecan Group, Männedorf, Switzerland). The MTT assay was performed two times in quadruplicate and the results were presented as percentage of the control (untreated cells) that was arbitrarily set to 100%. The cell viability rate (%) was calculated as follows: (A of the treated group/A control group) × 100.

2.5. Antibacterial Activity Assays

Pseudomonas aeruginosa PAO1 ATCC 27853, *Staphylococcus aureus* ATCC 25923 and *S. marcescens* ATCC 27117 used in this study were from the American Type Culture Collection (ATCC). *P. aeruginosa* PAOJP2, *P. aeruginosa* PA14-R3, and PAO1ΔpqsAmini-CTXlux-PpqsA were kindly provided by Dr. Livia Leoni, Department of Biology, Roma Tre University, Italy. *Chromobacterium violaceum* CV026 [18] was provided by Prof. Vittorio Venturi (ICGB, Trieste, Italy). Bacteria were routinely grown in Luria-Bertani (LB) medium (1% (w/v) NaCl, 1% (w/v) Tryptone, 0.5% (w/v) yeast extract) with or without agar (1.5% w/v) with shaking (180 rpm) at 37 °C. Tryptone soy broth (TSB) (Oxoid) was used for growth of *P. aeruginosa* PAO1 in the assay of biofilm formation. Antibiotics kanamycin (Kanamycin sulfate from *Streptomyces kanamyceticus*, BioReagent, Sigma-Aldrich, purity >95%), streptomycin (Streptomycin sulfate salt, BioReagent, Sigma-Aldrich, purity >96%) and tetracycline (BioReagent, Sigma-Aldrich, purity >96%) were incorporated into growth medium at the concentration of 30 µg/mL or 200 µg/mL in the case of kanamycin, 20 µg/mL for streptomycin and 100 µg/mL for tetracycline.

2.5.1. Minimum Inhibitory Concentration (MIC) Determination

The minimum inhibitory concentrations of bis-bibenzyls were determined according to standard broth microdilution assays recommended by the Clinical and Laboratory Standards Institute (M07-A9). Stock solutions of bis-bibenzyls were prepared in DMSO (50 mM). The highest tested concentration of each compound was 500 μ M. The inoculums were 10^5 colony forming units (CFU)/mL. The MIC value corresponds to the lowest concentration that inhibited the growth after 24 h at 37 °C.

2.5.2. Inhibition of *P. aeruginosa* PAO1 Biofilm Formation

P. aeruginosa biofilms were grown in TSB medium without shaking conditions. The effect of perrottetin F (**1**) and biotransformants on biofilm forming ability was tested at concentrations of 50% of determined MIC (0.5MIC) for each compound, on polystyrene flat-bottomed microtitre 96 well plates as previously described with modifications [19]. Briefly, overnight culture of *P. aeruginosa* was diluted in a fresh LB medium to optical density of OD_{600 nm}=0.2 and 100 μ L of the diluted culture with appropriate compound at the concentration of 0.5 MIC was added to the plate. After incubation for 24 h at 37 °C, the plate was washed twice with sterile water, dried, stained for 30 min with 0.1 % crystal violet in order to determine biofilm formed. Stained cells were washed with sterile water to remove unbound crystal violet. After drying, 150 μ L of 30% (v/v) acetic acid was added to solubilize the dye. After 10 min, the content of the wells was homogenized, and the OD of samples were measured at 550 nm using microplate reader Infinite 200PRO (Tecan). The experiment was done in quintuplicate and repeated two times.

2.5.3. Activity on Bacterial Pigment Production

Effect of bis-bibenzyls on violacein synthesis was evaluated with *Chromobacterium violaceum* CV026 indicator strain, as reported previously [18, 20]. Briefly, semisolid LB agar (0.3%, w/v; 5mL) was seeded with 50 μ L of an overnight culture of *C. violaceum* CV026 supplemented by *N*-hexanoyl- L-homoserine lactone (Sigma) to a final concentration of 5 μ M and poured over the surface of LB agar plates. When the overlaid agar had solidified, sterilized discs containing 250 μ g of each compound were placed on the plates. Petri dishes were incubated in the upright position overnight (30 °C) and examined for violacein synthesis. Inhibition of violacein synthesis was defined by the presence of white haloes in a purple background.

Overnight culture of *S. marcescens* was diluted 100-fold in molten semi-solid LB agar (0.3 % w/v) and poured over solid LB medium. Cellulose discs containing compounds (250 μ g/disc) were placed on solidified agar and incubated for 24 h at 30 °C. Inhibition of prodigiosin synthesis was identified by the absence of red colour around the disc.

2.5.4. Activity on Acyl Homoserine Lactones (AHLs) and Alkyl Quinolones (AHQs) Production using Biosensors

For the detection and quantification of long-chain (3OC12-HSL) and short-chain AHLs (C4-HSL) in AHLs production assays with two *P. aeruginosa* biosensors based on luminescent bacteria: *P. aeruginosa* PA14-R3 Δ lasIP_{rsaI}::lux and *P. aeruginosa* PAO1 Δ rhlIpKD-rhlA sensors for monitoring C12-AHL and C4-AHL are used, respectively [21, 22].

P. aeruginosa PA14-R3 Δ lasIP_{rsaI}::lux and *P. aeruginosa* PAO1 Δ rhlIpKD-rhlA biosensors reporter strains were grown overnight at 37 °C on LA agar plate. Plate for *P. aeruginosa* PAO1 Δ rhlIpKD-rhlA was supplemented with kanamycin (Sigma), 200 μ g/mL. Bacterial colonies were resuspended in LB medium with or without appropriate antibiotic and overnight cultures of the biosensors were diluted to OD₆₀₀=0.045 in LB. Bacterial suspension consists of 100 μ L of *P. aeruginosa* PA14-R3 Δ lasIP_{rsaI}::lux sensor supplemented by 3OC12-HSL lactone (Sigma) to a final concentration of 5 μ M or the suspension consists of 100 μ L of *P. aeruginosa* PAO1 Δ rhlIpKD-rhlA sensor supplemented by C4-HSL (Sigma), to a final concentration of 5 μ M, were dispensed in black,

clear-bottom 96-well microtiter plates. Microtiter plates were incubated at 37 °C with shaking, after 4 h of incubation OD600 and bioluminescence were simultaneously measured on Tecan Infinite 200 Pro multiplate reader (Tecan Group Ltd., Männedorf, Switzerland). Luminescence values were normalized by the cell density and expressed as percentage of the corresponding untreated control values. All experiments were performed in triplicate at least three times.

For detection and quantification of quorum sensing molecules 2-alkyl-4-quinolones (AHQs), such as 2-heptyl -3-hydroxy-4 quinolone (PQS) and 2-heptyl-4-quinolone (HHQ), a lux-based *P. aeruginosa* AHQ sensor was employed in liquid microtiter plate assay. 100 µl of *P. aeruginosa* PAO1 ΔpqsAmini-CTXluxPpqsA biosensor supplemented by HHQ lactone (Sigma), to a final concentration of 5 µM, was used for detection and quantification of AHQs molecules as described previously [23].

Compound 2: brown, amorphous solid; $[\alpha]_{\text{D}}^{22} +16.0$ (*c* 1.0, MeOH); UV (MeOH) λ_{max} (log ϵ) 212 (4.54), 275 (3.76); IR (KBr) ν_{max} 3413, 2923, 2855, 1600, 1506, 1456, 1352, 1342, 1322, 1257, 1215, 1168, 1033, 964, 853, 785 cm^{-1} ; ^1H NMR and ^{13}C NMR, see Table 1; HR-ESI-MS m/z 457.1668 $[\text{M} - \text{H}]^-$ (calcd for $\text{C}_{28}\text{H}_{25}\text{O}_6 - \text{H}$, 457.1651).

Compound 3: brown, amorphous solid; UV (MeOH) λ_{max} (log ϵ) 224 (3.56), 275 sh (4.32); IR (KBr) ν_{max} 3452, 2924, 2854, 1632, 1507, 1454, 1384, 1213, 1166, 1123, 1083, 1034, 792, 686 cm^{-1} ; ^1H NMR and ^{13}C NMR, see Table 1; HR-ESI-MS m/z 351.1244 $[\text{M} - \text{H}]^-$ (calcd for $\text{C}_{21}\text{H}_{20}\text{O}_5 - \text{H}$, 351.1233).

Compound 4: white, amorphous solid; UV (MeOH) λ_{max} (log ϵ) 215 (4.85), 272 (4.13), 282 (4.11); IR (KBr) ν_{max} 3413, 2925, 2857, 1595, 1590, 1505, 1455, 1281, 1242, 1215, 1166, 1060, 1021, 999, 856, 780, 729, 695 cm^{-1} ; ^1H NMR and ^{13}C NMR, see Table 1; HR-ESI-MS m/z 521.1283 $[\text{M} - \text{H}]^-$ (calcd for $\text{C}_{28}\text{H}_{26}\text{O}_8\text{S} - \text{H}$, 521.1270).

3. Results and Discussion

3.1. Isolation and Identification of *A. niger* Biotransformed Products of Perrottetin F

From the EtOAc extract of basic-neutral fraction of the *A. niger* medium, after separation on reversed-phased semipreparative HPLC compounds **2** and **3** have been isolated. From the acetone extract of the mycelium, compounds **2** and **4** have been isolated.

Compound **2** was obtained as a brown amorphous substance. Molecular ion $[\text{M} - \text{H}]^-$ at m/z 457.1668 of its HR-ESI-MS spectrum and the ^{13}C NMR spectrum (Table 1, Figures S7, S10 and S11, Supporting Information) were in accordance with molecular formula $\text{C}_{28}\text{H}_{26}\text{O}_6$. The UV absorption bands at 212 and 275 nm and the IR bands at 3413 and 1600 cm^{-1} indicated the presence of phenol moieties in **2**. In the ^1H NMR spectrum signals of the A-, C-, and D-rings were very similar to those of perrottetin F (**1**) (Figures S1 and S8, Supporting Information). However, the signals of the B-ring protons (connected by the COSY spectrum, Figure S12, Supplementary Material) were downfield shifted. One proton signal appeared at δ_{H} , 4.60 (t, 7.0 Hz) and there were six “bridge” protons (δ_{H} 2.5-2.9 region, Table 1) when compared with **1**. These data and the HSQC spectrum (Figure S15, Supporting Information), indicated the presence of a hydroxyl group at one of the four benzyl positions. Strong NOE correlations of the proton at δ_{H} 4.60 with H-10 and H-14, as well as weaker NOE correlations with H-3 and H-5 (Figure S14, Supporting Information) revealed that OH was at C-8. The COSY correlations H-8/H-7a, H-7b and the HMBC correlations H-10, H-14/C-8 gave additional confirmation (Figure S13 and S18, Supporting Information). Thus, the structure of the first new bis-bibenzyl **2** was established as 8-hydroxyperrottetin F. Its absolute configuration remained to be clarified.

Compound **3** was isolated as a brown amorphous substance. Its molecular $[\text{M} - \text{H}]^-$ ion at m/z 351.1244 from the HR-ESI-MS and ^{13}C NMR spectrum (Figures S19 and S22, Supporting Information) revealed the molecular formula $\text{C}_{21}\text{H}_{20}\text{O}_5$, i.e. seven C atoms less compared with compound **1**. The UV absorption bands at 224 and 275 nm and the IR bands at 3452 and 1632 cm^{-1} indicated the presence of phenol moieties in **3**. The 1D and 2D NMR data revealed the presence of

three benzene rings with the hydroxyl group at one benzyl position. The singlet at δ_H 4.45 (δ_C 65.0) originated from the methylene group, suggested terminal hydroxyl substituent. The HMBC correlations of this singlet with the overlapped signals of C-3 and C-5 and NOE correlations H-3, H-5/H-7 (Figures S27 and S24, Supporting Information) referred to the 7-hydroxyl derivative. The rest of 1H and ^{13}C NMR signals, very similar to those of the compounds **1** and **2**, were ascribed to the C and D rings atoms (Figures S1, S8, S23-S25, Supporting Information). Thus, the structure of compound **3** was established as shown in Figure 1.

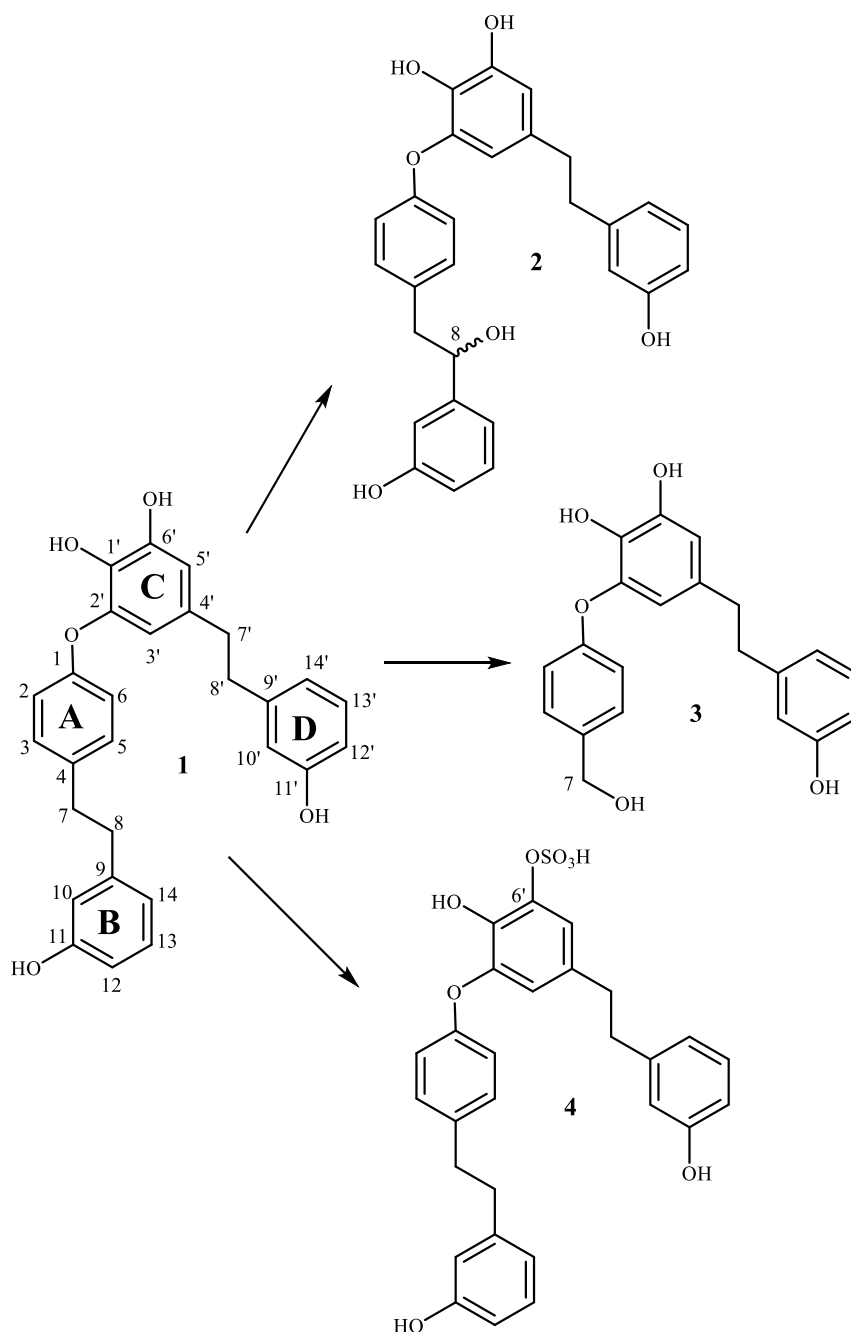


Figure 1. Biotransformed products of perrottetin F (**1**) obtained by *A. niger*.

Table 1. ¹H and ¹³C NMR Data of Compounds **2-4** recorded in CD₃OD (500 MHz for ¹H and 125 MHz for ¹³C)

Position	2		3		4	
	δ C	δ H	δ C	δ H	δ C	δ H
1	157.8	-	158.9	-	157.5	-
2	118.2	6.61 d (8.5)	118.3	6.74 d (8.5)	118.3	6.75 d (8.5)
3	131.7	6.90 d (8.5)	129.7	7.17 d (8.5)	130.7	7.07 d (8.5)
4	134.0	-	136.7	-	137.3	-
5	131.7	6.90 d (8.5)	129.7	7.17 d (8.5)	130.7	7.07 d (8.5)
6	118.2	6.61 d (8.5)	118.3	6.74 d (8.5)	118.3	6.75 d (8.5)
7	46.0	2.80 m	65.0	4.45 s	38.3	2.83 m
8	76.7	4.60 t (7.0)	-	-	39.3	2.80 m
9	147.5	-	-	-	144.8	-
10	114.2	6.63 t (2.0)	-	-	116.5	6.62 t (2.0)
11	158.4	-	-	-	158.5	-
12	115.2	6.52 dd (7.5; 2.0)	-	-	113.9	6.59 m
13	130.3	6.96 t (7.5)	-	-	130.4	7.05 t (8.0)
14	118.7	6.61 m ^b	-	-	121.0	6.64 m
1'	136.1	-	136.4	-	140.9	-
2'	145.6	-	145.3	-	142.6	-
3'	112.7	5.97 d (2.0)	113.0	6.03 d (2.0)	118.9	6.43 d (2.0)
4'	134.3	-	134.4	-	134.1	-
5'	112.4	6.33 d (2.0)	112.8	6.38 d (2.0)	119.5	7.05 d (2.0)
6'	147.8	-	148.0	-	146.2	-
7'	38.4	2.56 m	38.5	2.60 m	38.3	2.75 m
8'	39.1	2.56 m	39.2	2.60 m	39.0	2.75 m
9'	144.6	-	144.6	-	144.4	-
10'	116.5	6.40 t (2.0)	116.6	6.47 dd (8.0; 2.0) ^c	116.6	6.55 t (2.0)
11'	158.3	-	158.4	-	158.4	-
12'	113.8	6.43 d (8.0)	113.9	6.43 t (2.0)	113.9	6.57 dd (7.5; 1.0)
<u>13'</u>	<u>130.2</u>	<u>6.89 t (8.0)</u>	<u>130.3</u>	<u>6.92 t (7.5)</u>	<u>130.4</u>	<u>7.02 t (8.0)</u>
<u>14'</u>	<u>121.0</u>	<u>6.43 dd (8.0; 2.0)</u>	<u>121.0</u>	<u>6.47 dd (8.0; 2.0)^c</u>	<u>121.0</u>	<u>6.60 m</u>

^a *J* values are given in parenthesis^b signals overlaid with signals for H-2 and H-6^c signals overlaid

For compound **4**, obtained as a white amorphous substance, two molecular formulas C₂₈H₂₆O₈S and C₂₈H₂₆O₁₀ were possible regarding the molecular ion [M – H][–] at *m/z* 521.1283 from the HR-ESI-MS spectrum and the ¹³C NMR spectrum (Figures S28 and S30, Supporting Information). Calculating isotopic distribution and percentage of the [M+2-H][–] ion from the HR-ESI-MS spectrum, our experimental value of 9.1% was close to the theoretical value of 10.9% for the [M+2-H][–] ion for the molecular formula C₂₈H₂₆O₈S. In the case of C₂₈H₂₆O₁₀, [M+2-H][–] ion prediction was 6.7%. NMR analyses revealed great similarity of ¹H NMR spectrum of **4** to those of its biotransformation parent compound perrottetin F (**1**). The ¹H NMR spectrum of **4** was almost identical to that of perrottetin F (**1**) (Figures S29 and S1, Supporting Information), except the H-3' and H-5' signals of **4** that were downfield shifted (for 0.3 and 0.6 ppm, respectively), in comparison to compound **1**. This indicated another substituent at C-6' position, since multiplicity of both, H-3' and H-5', remained the same (two doublets with *J*=2.0 Hz). The difference of chemical shifts of C-ring carbons between **4** and perrottetin F (especially C-3', C-5', but also C-1', C-2' and C-6') additionally confirmed significant change in C-ring (Figures S30 and S3, Supporting Information). Having in mind the presence of sulphur in **4**, ie. the molecular formula C₂₈H₂₆O₈S, it was concluded that sulfate group was at C-6' position instead of OH group present in perrottetin F (**1**). All 2D NMR correlations confirmed the structure of perrottetin F-6'-sulfate (**4**) (Figures S31-S36, Supporting Information).

Although this is the first time that biotransformation of perrottetin F (**1**) was investigated, the obtained compounds are in agreement with literature data for the similar compounds obtained by the

microbial biotransformation using *A. niger*. The “target” positions in molecules during *A. niger* action were mostly allylic or benzyl C-atoms, or double bonds [24]. Successful biotransformation of thymol at the benzyl position by *A. niger* was reported [25]. In this work, compound **2** was obtained by hydroxylation at benzyl position C-8, and **3** by oxidative cleavage at the benzyl position C-7. Additionally, sulfate derivatives after *Aspergillus* spp. biotransformation were previously reported [26] and are in accordance with our findings (compound **4**).

3.2. Biological Activity Evaluations of Three Biotransformation Products in Comparison to Perrottetin F (**1**)

The cytotoxic activity (antiproliferative effect) of biotransformants was evaluated against two cell lines and compared to perrottetin F (**1**) (Table 2). Compounds (**2-4**) obtained by biotransformation of perrottetin F were less effective in inhibiting the growth of human lung carcinoma (A549) as well as human healthy lung fibroblasts (MRC5) in comparison to the parent compound. The reduction was between 1.6- to 16.6-fold for A549 and 2.3- to 8.3-fold for MRC5 cells, with the **4** being the least and **2** being the most active amongst biotransformants (Table 2). Some literature data proved that presence of sulfated groups may reduce cytotoxic potential of molecule [27]. For example, flavonoids luteolin and hispidulin showed higher cytotoxicity on melanoma cells line (B16F10) compared with luteolin-7-sulphate and hispidulin 7-sulphate [27]. This can be reason of slightly lower activity of compound **4** in regard to perrottetin F.

Biotransformed product **2** also showed the slight increase in the selectivity toward cancer in comparison to healthy cell line compared to perrottetin F.

Table 2. The cytotoxicity (IC₅₀, μM) and antibacterial properties (MIC, μM) of three biotransformation products in comparison to perrottetin F

	IC ₅₀ (μM)		MIC (μM)	
	A549	MRC5	<i>S. aureus</i>	<i>P. aeruginosa</i> PAO1
Perrottetin F (1)	15 ^a	30	100	150
2	25	70	450	200
3	50	80	400	250
4	250	250	400	250

^aThe results are expressed as means calculated from three independent experiments with standard deviation between 1-3%.

Next, we focused on antibacterial potential of perrottetin F (**1**) and its three biotransformed products against two common nosocomial opportunistic pathogens: *S. aureus* and *P. aeruginosa* PAO1 (Table 2). Very weak antibacterial activities against *S. aureus* and *P. aeruginosa* PAO1 with minimal inhibitory concentrations (MICs) ranging from 100 μM for perrottetin F to 450 μM for **2** were recorded. Again, the activity of biotransformants was 1.3- to 4.5-fold reduced in comparison to the parent compound (Table 2).

The effect of compounds (**2-4**) on bacterial pigment production was assessed using *C. violaceum* CV026 and *S. marcescens* (Figure S37, Supporting Information). *C. violaceum* CV026 is white, violacein negative, and has no constitutive ability of violacein pigment production, but CV026 can sense exogenously added AHLs and responds by producing purple violacein [18]. In this assay, the tested compounds demonstrated only weak antagonistic activity on violacein production in *C. violaceum* CV026 when 250 μg of test compound was applied per disc. Inhibition of violacein production with **2** and **3** was visible as a slightly brighter area around the discs (Figure S37A, Supporting Information). The interference of perrottetin F and biotransformants with pigment production in *S. marcescens* was examined by assessing modulation of prodigiosin production in a disc assay (Figure S37B, Supporting Information). It was obvious from the obtained results that compounds did not trigger inhibition of prodigiosin synthesis, however, added compounds could change the color of the halo around the discs, from red to pink. The pink zones were larger in the presence of biotransformants than in the presence of parent compound perrottetin F (**1**) (Figure S37B, Supporting Information). Taken together, obtained results with *C. violaceum* CV026 and *S. marcescens*

suggested that perrottetin F as well as three metabolites (**2-4**) had moderate effect on violacein and prodigiosin production under tested conditions.

S. marcescens has N-acylhomoserine lactones (AHLs)- dependent QS system and produces at least four AHLs (3-oxo-C6-HSL, C6-HSL, C7-HSL, and C8-HSL), which regulate production of prodigiosin and phenotypic characteristics such as motility and biofilm formation that have been connected to virulence of this strain [28]. On the other side, *P. aeruginosa* also possesses four QS systems: two acyl-homoserine lactone (AHL)-mediated QS systems, one 2-alkyl-4-quinolones (AHQs)-mediated QS system and the IQS-dependent QS system. Three autoinducer synthases LasI, RhlI, and PqsABCDH produce autoinducers 3-oxo-C12-homoserine lactone (HSL), C4-HSL, and 2-heptyl-3-hydroxy-4-quinolone, respectively. To evaluate the potential of the perrottetin F (**1**) and biotransformed products **2-4** modulators of specific QS pathway in *P. aeruginosa*, three mutant biosensors strains dependent on exogenous AHLs were used: *P. aeruginosa* PA14-R3ΔlasIPrsal::lux, used to determine 3OC12-HSL production (LasI activity), *P. aeruginosa* PAO1 ΔrhlIpKD-rhlA, used for C4-HSL production (RhlI activity), and *P. aeruginosa* PAO1 ΔpqsA mini-CTX luxPpqsA, used for assessment of PQS production (PqsABCDH activity).

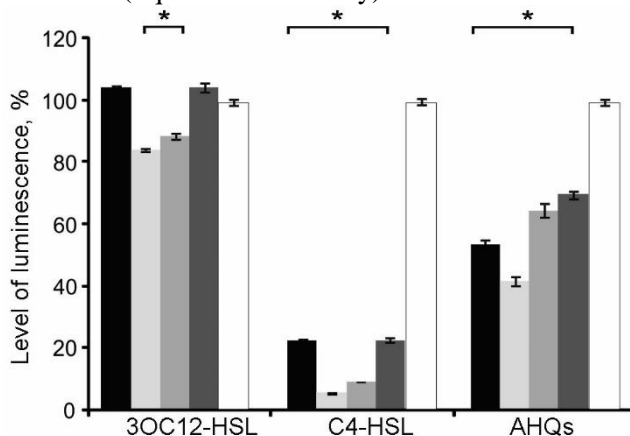


Figure 2. Effect of perrottetin F and compounds **2-4** (100 μ M) on acyl homoserine lactone (AHL) and AHQs production in *P. aeruginosa*. The *P. aeruginosa* PA14- R3ΔlasIPrsal : lux strain was used to directly measure the levels of 3OC12- HSL and *P. aeruginosa* PAO1 ΔrhlIpKD-rhlA reporter strain was used to directly measure the levels of C4-HSL. Detection of AHQs (PQS and HHQ) was performed by the biosensor *P. aeruginosa* PAO1 ΔpqsA mini-CTX luxPpqsA. Biosensors grown in the presence of DMSO served as the 100% luminescence control (□); ■- perrottetin F (**1**), □- **2**, ■- **3**, ■- **4**

Obtained data (Figure 2) suggest that perrottetin F and biotransformants reduced emitted bioluminescence in biosensor strains, in a range of 10% to 90% compared to negative control (DMSO). All these compounds exerted the most potent effects on C4-HSL (Rhl activity), moderate effects on AHQs production (PqsABCDH activity) and weak effects on 3OC12-HSL production (Las activity). Biotransformed products, especially **2** and **3**, had more pronounced effect in comparison to parent perrottetin F. Compounds **2** and **3** significantly reduced the bioluminescence in *P. aeruginosa* PAO1 ΔrhlIpKD-rhlA negatively for 95% and 90 %, respectively, in comparison to DMSO control (Figure 2). Also, these compounds significantly reduced the bioluminescence in *P. aeruginosa* PAO1 ΔpqsAmini-CTX luxPpqsA for 60% and 35%, respectively, and in addition, reduced bioluminescence in *P. aeruginosa* PA14-R3ΔlasIPrsal::lux, for 16% and 12% respectively. Perrottetin F (**1**) and compound **4** inhibited the bioluminescence in *P. aeruginosa* PAO1 ΔrhlIpKD-rhlA for 80%, and negatively affected the bioluminescence in *P. aeruginosa* PAO1 ΔpqsAmini-CTX luxPpqsA for 50% and 30%, respectively, while these compounds had no effect on the bioluminescence in *P. aeruginosa* PA14-R3ΔlasIPrsal::lux.

Taken together, these results suggested that the tested compounds, that showed no significant effect on the growth of *P. aeruginosa* and *S. aureus* (Table 2), were able to successfully modulate the production of signaling molecules that are part of the bacterial QS system (Figure 2).

According to literature data, acyclic bis-bibenzyl, perrottetin F (**1**) and cyclic bis-bibenzyls, marchantins A, B, and D showed cytotoxicity against KB cells (3.7–20 μ M), DNA polymerase β

inhibitory (ID_{50} 14.4–97.5 μ M), and anti-HIV-1 (5.30–23.7 μ g/mL) activity [29]. Perrottetin F exhibited very strong nitric oxide (NO) production inhibitory activity, in culture media, on RWA 264.7 cells [4]. Perrottetin F also inhibited influenza PA (subunit of RNA polymerase) endonuclease activity at 10 μ M [4, 8]. Perrottetin E, bis-bibenzyl isolated from *Radula perrottetii* inhibited KB cells at a concentration of 12.5 μ g/mL [4].

A series of bis-bibenzyls, including riccardin-, marchantin- and isoplagiochin-class structures were evaluated for anti-MRSA activity against two strains of *S. aureus* MRSA [30]. The structure–activity relationships and the results of molecular dynamics simulations showed that bis-bibenzyls with potent anti-MRSA activity commonly have a 4-hydroxyl group at the D-benzene ring and a 2-hydroxyl group at the C-benzene ring in the hydrophilic part of the molecule, and an unsubstituted phenoxyphenyl group in the hydrophobic part of the molecule containing the A–B-benzene rings. It was found previously that the number of hydroxyl groups on the riccardin framework is important for anti-MRSA activity, thus, bis-bibenzyls derivative with hydroxyl group did not show apparent anti-MRSA activity, whereas bis-bibenzyls with two or three phenolic hydroxyl groups showed anti-MRSA activity. Also, it was shown that riccardin-class bis(bibenzyl)-type cyclic phenolic compounds exhibit potent antibacterial activity towards MRSA *S. aureus*, comparable to that of vancomycin and linezolid [30]. Cyclic bis-bibenzyl, marchantin A from many *Marchantia* species, showed antibacterial activity against *Bacillus cereus* (MIC 12.5 μ g/mL), *B. megaterium* (25 μ g/mL), *B. subtilis* (25 μ g/mL), *Cryptococcus neoformans* (12.5 μ g/mL), *Escherichia coli*, *Pseudomonas aeruginosa* and *Staphylococcus aureus* (3.13–25 μ g/mL) [4].

AHL-mediated systems regulate the expression of many virulence factors, including LasA protease, phospholipase, exotoxin A, pyocyanin, rhamnolipids, elastase, and also regulate biofilms development. *Pseudomonas aeruginosa* PAO1 are known to produce a diverse 2-alkyl-4-quinolones (AHQs) which act as QS signal molecules and are involved in the regulation of many virulence factors. In the quinolone-dependent QS system, PqsABCDE and PqsH are involved in the synthesis of 2-heptyl-4-quinolone (HHQ) and 2-heptyl-3-hydroxy-4-quinolone (PQS) [23]. Both HHQ and PQS can bind to the transcriptional regulator MvfR and form a complex to regulate gene expression. *P. aeruginosa* PQS and HHQ regulate the production of pyocyanin, rhamnolipids, elastase and influence a biofilm formation [28]. The production of diverse AHQ, which act as QS signal molecules and are also involved in the regulation of many virulence factors in *P. aeruginosa* PAO1 (i.e. elastase, pyocyanin, rhamnolipids, biofilm development).

Biotransformation of bis-bibenzyls has not yet been investigated. Biotransformed products of these molecules found in the most thalloid liverworts could be useful in the study of antimicrobial, anticancer and other biologically active compounds. The microbial transformation is safe, cheap and constitutes an environmentally very friendly method which could be a starting platform for the synthesis of new pharmacological drugs. In the present research, three new bis-bibenzyl derivatives are obtained through biotransformation of perrottetin F by *A. niger* and their cytotoxic and antimicrobial properties were tested. Moreover, biotransformants showed an extraordinary ability to inhibit synthesis of bacterial quorum-sensing signal molecules such as short chain acyl homoserine lactones (AHLs).

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Supporting Information

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