



Antimicrobial Activity and Microbial Transformation of Ethyl *p*-Methoxycinnamate Extracted from *Kaempferia galanga*

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ABSTRACT

Ethyl *p*-methoxycinnamate (EPMC), a major constituent of the *Kaempferia galanga* rhizome, was transformed to ethyl *p*-hydroxycinnamate (EPHC) using *Aspergillus niger*. The EPHC metabolite was elucidated by NMR spectroscopic technique. Antimicrobial microbial study found that EPHC was active against all strains tested with a good minimum inhibitory concentration (MIC). It was active against both *Staphylococcus aureus* and *Bacillus cereus* at MIC 333 µg/ml while against *Pseudomonas aeruginosa*, *Escherichia coli* and *Candida albicans* at MIC 111 µg/ml. It was also shown that EPHC exhibited more growth inhibition potential than EPMC. Besides that, EPHC has shown the minimum bactericidal concentration (MBC) toward *B. cereus*, *P. aeruginosa* and *E. coli* at the concentration of 1000 µg/mL while EPMC did not show killing potential toward the tested microorganisms.

Key words: Biotransformation, *Kaempferia galanga* Linn, Ethyl *p*-methoxycinnamate, Ethyl *p*-hydroxycinnamate, *Aspergillus niger*, Antimicrobial.

INTRODUCTION

The need for a new antibiotic against pathogenic microorganism has increased due to the current problems of resistance associated with them¹. Plant based antimicrobial agents have enormous therapeutically potential as they can serve the purpose without any side effect, that often associated with synthetic agents.

Malaysian natural product plants have been studied extensively. These include medicinal plant species from *Andrographis*^{2,3,4,5}, *Musa*⁶, *Plumeria*⁷, *Citrus*^{8,9}, *Cymbopogon*¹⁰, *Artocarpus*¹¹, *Alpinia*¹² and *Centella*¹³. However, plant species of Zingiberaceae continue to attract much phytochemical interests due to their culinary uses, besides their biological and pharmaceutical activities. *Kaempferia galanga* Linn., belongs to the Zingiberaceae family, is one

of the plants that mainly distributed in tropics and subtropics of Asia. Besides Malaysia, it can be found in Southern China, Indochina and India¹⁴. It has been known from ancient times that the essential oil of *K. galanga* plays an important role in native medicine as it possesses interesting biological activities such as antimicrobial¹⁵, antihypertensive¹⁶, anticancer¹⁷ and antiproliferative activity¹⁸.

The emergence of biotransformation technique to produce new natural drug as an antimicrobial agent was successfully reported. According to Muller¹⁹, fungi are effective in increasing the chemical diversity by producing new derivatives with lower toxicity but improved biological activity. While Kim *et al*²⁰, described that the biotransformation afforded another metabolite that has higher antimicrobial activity compared to the parent metabolite. Thus, this approach successfully enhances the antimicrobial activity of the compound. For example, rhapontigenin produced from rhapontin by biotransformation showed 4-16 times higher antimicrobial activity than rhapontin. The activity was higher against Gram-positive strains than Gram-negative strains²⁰. Another study by Srivastava *et al*²¹, indicates that *A. flavus* can transform artemisinin to deoxyartemisinin (Figure 1) and shows antibacterial activity against *S. aureus*, *S. epidermidis* and *S. mutans* at a minimum inhibitory concentration (MIC) of 1 mg/mL compared to artemisinin whose MIC was >2 mg/mL. Siddhardha *et al.*²² showed biotransformation of (-)- α -santonin using *A. parasiticus* resulted in the production of 3,4-epoxy- α -santonin, 4,5-dihydro- α -santonin and 1,2-dihydro- α -santonin and evaluated for the biological activity of the transformed products. 3,4-epoxy- α -santonin exhibited higher degree of antibacterial and antifungal activity. 4,5-dihydro- α -santonin showed slight increase in the activity, compared to (-)- α -santonin. 1,2-dihydro- α -santonin is equipotent to the substrate (-)- α -santonin. This is the first report on the biotransformation of (-)- α -santonin by *A. parasiticus* and evaluation of the biological activities of the biotransformed products.

Therefore, this present study was initiated to screen the antimicrobial activity of ethyl *p*-methoxycinnamate (EPMC), extracted from *K. galanga* Linn., and compare the bioactivity with its biotransformation metabolite, ethyl *p*-

hydroxycinnamate (EPHC) produced by *A. niger*.

EXPERIMENTAL

Materials

The rhizomes of *K. galanga* L. were collected at different locations throughout Malaysia in December 2013 and stored at -20 °C prior to analysis. Five reference strains of human pathogens used for antimicrobial assay were *Staphylococcus aureus* (ATCC25923), *Bacillus cereus* (ATCC11778), *Pseudomonas aeruginosa* (ATCC27853) and *Escherichia coli* (ATCC8739) while one fungal strains was *Candida albicans* (IMRC533/11 A). The bacteria were maintained on Muller-Hinton Agar (MHA) and Muller-Hinton Broth (MHB) and stored in the refrigerator at 4°C. Meanwhile, *C. albicans* was maintained on Sabouraud's Dextrose Agar (SDA) and stored in the refrigerator at 4°C. Both bacteria and fungus *C. albicans* were subcultured onto fresh media at regular interval.

Extraction of *K. galanga* oil and purification of ethyl *p*-methoxycinnamate

The fresh rhizomes of *K. galanga* were washed and cut into smaller pieces. The rhizomes were then extracted using steam distillation and supercritical fluid extraction (SFE) according to the previously reported method²³. The essential oil was collected and stored at -4°C for future analyses. For purification of ethyl *p*-methoxycinnamate (EPMC, 1) the essential oils were stirred with boiling water and then recrystallized at cold temperature (-4°C). After crystallization, the mixture was filtered through Whatman No 1 and then the crystal was kept in the desiccator for 24 hours prior to further analysis.

Fungus culture preparation and biotransformation procedure.

Aspergillus niger fungal culture preparation Pure fungal culture of *A. niger* was obtained from Microbiology Laboratory Kulliyyah of Science, International Islamic University Malaysia. The pure fungus was streaked on Sabouraud Dextrose Agar slant (SDA) at 30 °C for a week and stored at 4 °C. After cultivation, the well grown mycelia on the agar slants were placed in the conical flask (250 ml) and inoculated with 10.0 ml of sterilized medium broth containing glucose, glycerol, peptone, yeast extract, KH₂PO₄, NaCl and distilled H₂O. The flask

was incubated using an incubator shaker for 48 hr at 120 rpm and 30°C. Then, mycelia suspensions were transferred into 17 flasks containing sterile broth medium (100 ml of each) and incubated for 24 hour using the similar condition as reported earlier²⁴.

Biotransformation of the bioactive compounds

Ethyl p-methoxycinnamate, **1** (480 mg,) was dissolved in dimethyl sulfoxide (DMSO) (24 ml) and distributed among 48 flasks containing 24 hour stage culture (10 mg/0.5 ml in each flask) and continuously shaken for 24 h using a rotary shaker (120 rpm) at 30 °C. At the same time, a control flask having substrate without fungus and a control flask containing fungus without substrate were analysed in order to check the substrate ability and to determine the endogenous metabolite, respectively.

Separation and purification of biotransformed product

After incubation, the culture media and mycelium were separated using cotton in a funnel. Then, mycelium was washed with ethyl acetate (1.5 liter) while the culture media were extracted 3 times with ethyl acetate (2 liter). The combined organic extract was dried with anhydrous sodium sulphate (Na₂SO₄) and evaporated using a rotary evaporator. The similar procedures were employed for the control group. The residues from both experimental and control group were analysed by TLC to confirm the presence of biotransformed product. The biotransformed products were isolated by column chromatography using silica gel column (200-300 mesh, Merck Ltd.) with hexane: ethyl acetate as solvent.

Structural elucidation of the biotransformed product

Gas Chromatography/Mass Spectrometry (GC/MS)

The crystal was further analyzed using an Agilent 5975C GC-MSD system (Agilent, Avondale, USA) equipped with a 30 m x 0.25 mm i.d. x 0.25 μm, HP-5 capillary column (SGE, Australia) and 70 eV of the electron impact technique. The carrier gas was helium at a flow rate of 1.3 mL/min. The crystal was dissolved in dichloromethane (DCM) at concentration of 1%. Then, 1 μL of sample was injected into the column. The injector and

detector temperatures were 250 °C and 230 °C, respectively. The other analytical conditions were as follows: Temperature programming: 60 °C, as initial temperature, for 5 min, 8 °C/min to 180 °C at 10 °C/min to 240 °C, holding for 5 min. Mass spectroscopy analysis was carried out using positive electron ionization method in order to obtain fragments of the compounds. The identification of compounds were based on a comparison of their retention times with those of authentic standards and by comparison of their mass spectra with those of data in the Wiley Registry of Mass Spectral Data and National Institute of Standards and Technology (NIST) libraries¹⁸. Low resolution mass spectrometry (m/z) was carried out on a VG Biotech Quattro II triple quadrupole instrument.

Nuclear Magnetic Resonance (NMR) analysis

The nuclear magnetic resonance (NMR) spectroscopy was used to elucidate the structure of compound (**1 & 2**) using 1D (1H- and ¹³C-NMR) and 2D (COSY 45°, HSQC, HMBC and NOESY) experiments according to the standard protocols. The NMR analysis was carried out at the Department of Chemistry, Universiti Kebangsaan Malaysia. ¹H and ¹³C NMR spectra were recorded on a Bruker 400 MHz FT-NMR spectrometer. The crystal was dissolved in CD₂Cl₂ and chemical shifts (α) are reported in ppm downfield of tetramethylsilane. Coupling constants (J) are quoted in Hz.

Antimicrobial Assays

Minimum Inhibitory Concentration (MIC)

The test was performed using 96-well plates and prepared in a three-fold dilutions manner, where the concentration in each well is three times less than the preceding well. The first well for each sample (single compounds) was prepared in a 1000 μg/mL concentration by mixing 30 μL of the single compounds (10 mg/mL concentration) with 270 μL broth media. After mixing the samples with the broth, the mixture (100 μL) was transferred to the next well, in the same column, containing 200 μL broth media. The similar procedure was repeated for the subsequent well until the end of each column. The final concentrations formed in each well after this process were 1000 (first well), 333, 111, 37, 12.3, 4.1, 1.37, and 0.45 μg/mL²⁵.

Minimum Bactericidal/Fungicidal Concentration (MBC)

The minimum bactericidal/fungicidal concentration (MBC/MFC) test was carried out according to Betts *et al.*²⁶, with slight modifications in order to determine whether the MIC values only inhibit the growth or even kill the microorganism. 100 μ L of each broth in the well that represents MIC values for all bacteria and fungus was spread in MHA and SDA, respectively. Plates were kept in incubator for 24 hr. The MBC/MFC could be determined when the microorganisms were killed and unable to regrow back.

RESULTS AND DISCUSSION

Microbial transformation of ethyl *p*-methoxycinnamate (**1**) by *A. niger* afforded a main compound, identified as ethyl *p*-hydroxycinnamate (**2**) (Figure 2) in good yield (24%). The molecular formula of **2** proved to be C₁₁H₁₂O₃ from ¹H-NMR

and ¹³C-NMR data. The NMR spectral data of compound **2** are summarized in the Table 1. These finding presented a clear correlation to support the proposed structure and identified by comparing their spectra with literature²⁷.

The antimicrobial assay in terms of the minimum inhibitory concentrations (MIC) of ethyl *p*-methoxycinnamate (EPMC) and ethyl *p*-hydroxycinnamate (EPHC) against different pathogenic bacteria and fungus are shown in Table 2.

MIC tests were done selectively against four types of bacteria which were *B. cereus*, *S. aureus*, *E. coli*, *P. aeruginosa* and one fungus which was *C. albicans*. The result showed that both compounds have good antimicrobial activities against all tested bacteria and fungus. The MIC values also indicated that both compounds (EPMC & EPHC) were able to inhibit Gram-negative bacteria at lower

Table 1: ¹H- NMR, ¹³C-NMR, COSY and HMBC Spectral Data of EPHC, 2

Position	¹³ C-NMR	¹ H- NMR	COSY	HMBC
1	127.07, C	-	-	
2, 6	129.94, CH	7.4 (2H, d, $J_5 = J_9 = 8.4$ Hz)	H3,5	C 1,4,7
3, 5	115.80, CH	6.8 (2H, d, $J_6 = J_8 = 9.0$ Hz)	H2,6	C 3,4, 5
4	157.98, C	-	-	
7	144.18, CH	7.6 (1H, d, $J = 15.6$ Hz)	H8	C 1, 8, 9
8	115.55, CH	6.3 (1H, d, $J = 16.2$ Hz)	H7	C 1,7,9
9	167.42, C	-	-	
10	60.46, CH ₂	4.2 (2H, q, $J = 7.2$ Hz)	H11	C 11, 9
11	14.12, CH ₃	1.3 (3H, t, $J = 7.2$ Hz)	H10	C 10

Table 2. Minimum Inhibitory Concentration (MIC) of ethyl *p*-methoxycinnamate (EPMC) and ethyl *p*-hydroxycinnamate (EPHC)

Microorganisms	Minimum Inhibitory Concentration (MIC) value (μ g/mL)	
	EPMC, 1	EPHC, 2
<i>Staphylococcus aureus</i>	1000	333
<i>Bacillus cereus</i>	1000	333
<i>Escherichia coli</i>	333	111
<i>Pseudomonas aeruginosa</i>	333	111
<i>Candida albicans</i>	333	111

Table 3. Minimum Bactericidal/Fungicidal Concentration (MBC/MFC) of ethyl *p*-methoxycinnamate (EPMC) and ethyl *p*-hydroxycinnamate (EPHC)

Microorganisms	MBF/MFC value (µg/mL)	
	EPMC, 1	EPHC, 2
<i>Staphylococcus aureus</i>	-	1000
<i>Bacillus cereus</i>	-	1000
<i>Escherichia coli</i>	-	1000
<i>Pseudomonas aeruginosa</i>	-	1000
<i>Candida albicans</i>	-	333

concentrations compared to Gram-positive bacteria. The MIC value of ethyl *p*-hydroxycinnamate (EPHC) against both *E. coli* and *P. aeruginosa* was 111 µg/mL, which were more active than the value exhibited by ethyl *p*-methoxycinnamate (EPMC, MIC = 333 µg/mL) (Table 2). Meanwhile, the MIC values of EPHC against both *B. cereus* and *S. aureus* was 333 µg/mL, which were also lower than the value exhibited by EPMC (MIC=1000 µg/mL). Furthermore, EPHC was able to inhibit the growth of *C. albicans* with the concentration that exhibited inhibition potential (MIC) was 111 µg/mL which were lower than the value exhibited by EPMC (MIC= 333 µg/mL).

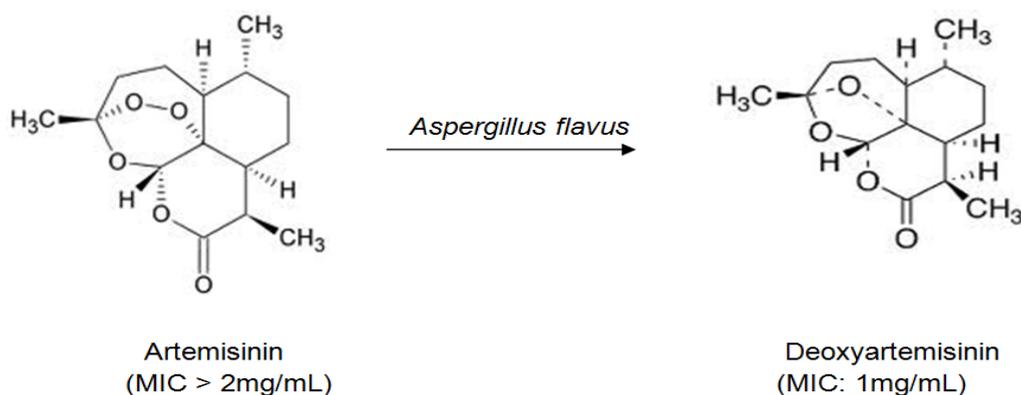


Fig. 1: Biotransformation of artemisinin to deoxyartemisinin by *Aspergillus flavus*

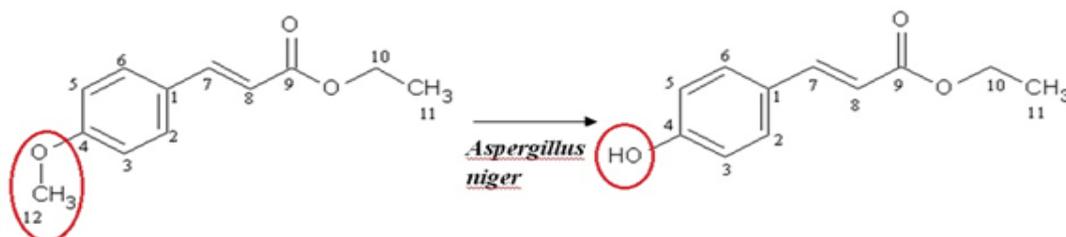


Fig. 2: Biotransformation pathway of ethyl *p*-methoxycinnamate into ethyl *p*-hydroxycinnamate acid by *Aspergillus niger*

In terms of minimum bactericidal concentration (MBC), there were bactericidal effect of EPHC against *S. aureus*, *B. cereus*, *E. coli* and *P. aeruginosa* at 1000 µg/mL and fungicidal effect (MFC) against *Candida albicans* at 333 µg/

mL (Table 3). However, there were no MBC was obtained for ethyl *p*-methoxycinnamate (EPMC) against all tested microorganisms. Hence, ethyl *p*-hydroxycinnamate (EPHC) has great potential to inhibit and killing the growth of Gram-positive

bacteria, Gram-negative bacteria and fungus compared to ethyl *p*-methoxycinnamate (EPMC). The present of hydroxyl group in structure may increase the antimicrobial activities of compound²⁸. The result has proved that the folkloric use of this plant in treating microbial infections and suggested that ethyl *p*-hydroxycinnamate (EPHC) could be exploited in future as one of the new antibiotics in line with the current drug development.

CONCLUSION

Biotransformation has been successfully utilized as a tool to generate pharmaceutical compounds from natural products. Through this process, ethyl *p*-methoxycinnamate (EPMC), extracted from *Kaempferia galanga* plant, was transformed using *Aspergillus niger* to ethyl

p-hydroxycinnamate (EPHC), which shown antimicrobial activities.

Looking at the antimicrobial activities, ethyl *p*-hydroxycinnamate (EPHC) has an inhibitory potential effect against *S. aureus*, *B. cereus*, *P. aeruginosa*, *E. coli* and *C. albican* better than the effect by ethyl *p*-methoxycinnamate (EPMC). EPHC has also shown bactericidal and fungicidal effects against all tested bacteria and fungus while EPMC did not show killing potential toward them.

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