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Chemical constituents and *in vitro* **antimicrobial activity of rhizome essential oils of** *Zingiber densissimum* **S.Q.Tong & Y.M.Xia and** *Kaempferia laotica* **Gagnep. growing wild in Vietnam**

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Abstract

Zingiber densissimum S.Q.Tong & Y.M.Xia and *Kaempferia laotica* Gagnep., belong to the Zingiberaceae family, which were found in Vietnam and Laos in recent years. To date, there has been very limited information on their phytochemical profiles and pharmacological activities. The present study was conducted to give insights into the chemical composition of the essential oils isolated from two plants and their antimicrobial activities. Gas Chromatographic–Mass Spectral (GC-MS) analysis was utilized to identify the chemical composition of the rhizome essential oils of *Z. densissimum* and *K. laotica*. The antimicrobial activities were evaluated on three Gram-positive bacterial strains, three Gram-negative bacterial strains, and a pathogenic yeast. The GC-MS analysis showed that β-pinene (38.36%), β-phellandrene (26.85%), and α-pinene (13.31%) were the most dominant components in *Z. densissimum* rhizome essential oil, while *K. laotica* rhizome essential oil was found to contain mainly camphene (23.44%), α-pinene (10.69%), 3-carene (8.60%) and β-pinene (6.85%). As for the antimicrobial activity test results, the rhizome essential oil of *Z. densissimum* displayed potent activities against *Candida albicans* (MIC value = 2 µg/mL), *Enterococcus faecalis*, *Staphylococcus aureus*, and *Escherichia coli* (MIC values = 8 µg/ mL). Meanwhile, the rhizome essential oil of *K. laotica* showed weaker activities against all investigated microbial strains (MIC values ranging from 32 to 256 µg/mL)*.* The study results indicated the chemical constituents of the essential oils prepared from these plants. The two essential oil samples were also shown to exhibit potential antimicrobial activities against several pathogenic bacterial and fungal strains.

Keywords

Zingiber densissimum, *Kaempferia laotica*, GC-MS analysis, Essential oil, Antimicrobial activities

Introduction

The genus *Zingiber* is the third largest of the Zingiberaceae family and contains over 140 species. Most of the species are edible and medicinal plants that possess aromatic odors. Many species from the genus, for example, *Z. zerumbet*, *Z. officinale* (commonly known as ginger), *Z. corallinum*, and *Z. mioga*, have long been used in folklore medicine for the treatment

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of nausea, vomiting, common cold, and for relieving symptoms from arthritis and abdominal pain¹. Among recorded species, Z. densissimum was first discovered by Tong and Xia in southern Yunnan, China, in 1987, and then by Triboun in Thailand in 2014. Recently, *Z. densissimum* has been found in Lam Dong Province, Vietnam.

The genus *Kaempferia* is a medium-sized genus in the Zingiberaceae family with around 60 species distributed in Asia, mainly in India and Southeast Asia. In Vietnam, only seven species were discovered in natural habitat, including some plants widely used in traditional medicine, such as *K. galanga*. Vietnamese people have long used some *Kaempferia* species for the treatment of indigestion, and the alcoholic extract can be used to relieve symptoms of arthritis. In 2015, Tuan Nguyen-Hoang et al.² discovered the natural occurrence of *K. laotica* in Ba Ria-Vung Tau Province, Vietnam. Literature research showed that *K. laotica* is only distributed in Xiangkhuang Province (Laos), Ubon Ratchalthani and Buri Ram Provinces (Northeast Thailand)².

To date, there have been no studies on these two species of Zingiberaceae in terms of phytochemistry and bioactivity. In the present study, *Z. densissimum* rhizomes and *K. laotica* rhizomes (Fig. S1) were examined about the chemical composition of the essential oils (EOs) and their antimicrobial activities. These findings are part of research about the understudied ginger species growing wild in Vietnam, contributing to our literature about the natural product profiles and antimicrobial efficacy of these species.

Materials and Methods Plant materials

In this study, *Z. densissimum* rhizomes were collected from Da Chais Commune, Lac Duong District, Lam Dong Province (GPS: 12°09'37.10"N 108°39'53.49"E), Vietnam in August 2022, while *K. laotica* rhizomes were gathered from Big Mountain, Vungtau City, Ba Ria Vung Tau Province (GPS: 10°22'47.72"N 107°03'29.81"E), Vietnam in July 2022. The identification and authentication of the two plants were done by Assoc. Prof. Dr. Nguyen

Hoang Tuan. Voucher specimens (NHTuan 057 and NHTuan 058, respectively) were deposited at the herbarium of Hanoi National University.

Essential oil isolation

The fresh rhizomes (500 g) were cleaned, cut into small pieces, mixed with 1.5 L of distilled water, and then hydrodistilled using a Clevenger apparatus. The isolation process lasted until no further EO could be extracted (3.5 h). The collected EOs were treated with anhydrous Na_2SO_4 (Merck) to remove the traces of water and stored in sealed vials in the dark at 4°C until analysis. The experiment was done in triplicate. The EOs were obtained in yields of 0.20% (v/w, *Z. densissimum* rhizome EO) and 1.09% (v/w, *K. laotica* rhizome EO), calculated on a fresh weight basis.

Gas chromatographic-mass spectral (GC-MS) analysis and identification of the constituents

The GC-MS analysis was employed to identify the chemical composition of the EOs from *Z. densissimum* and *K. laotica* rhizomes, which was performed on an Agilent Technologies 7890B GC System with an HP-5MS UI column (size: 30 m \times 0.25 mm, 0.25 µm film thickness), and coupled with an Agilent 5977B MSD model. Analysis conditions were set up as follows: Helium was the carrier gas (flow rate: 1.0 mL/min); injection volume: 1.0 μL with a split ratio of 25:1; oven temperature 0-2 min: 50°C; 2-22 min: 50-150°C (temperature rising rate: 5°C/min); 22-32 min: 150°C; 32- 45 min: 150-280°C (temperature rising rate: 10° C/min). The respective temperatures were also set up for the injector (300°C), MS Quad (150°C), transfer line temperatures (300°C), and MS source (230°C). The MS conditions were documented at 70 eV with a range of 50-550 amu at 2.0 scan/s. The individual chemical components were identified based on comparison of their retention indices (RI) and their mass spectral fragmentation patterns to those listed in the reference libraries (NIST17 and Adams book) $3-6$. The relative percentage of compounds was calculated from the peak area percentage in the GC chromatogram^{7,8}.

Antimicrobial activity test

The antimicrobial activities of the EO samples were evaluated on Gram-positive bacterial strains (including *Enterococcus faecalis* ATCC 299212, *Staphylococcus aureus* ATCC 25923, and *Bacillus cereus* ATCC 14579), Gramnegative bacterial strains (including *Escherichia coli* ATCC 25922, *Pseudomonas aeruginosa* ATCC 27853, *Salmonella enterica* ATCC 13076), as well as a pathogenic yeast (*Candida albicans* ATCC 10231), which were provided by National Institute for Food Control, Vietnam. The antimicrobial activities were assessed by microdilution broth susceptibility assay method, which was referenced from the standards issued by the Clinical and Laboratory Standards Institute (CLSI supplement M100, 2020)⁹. Luria Bertani Agar (LBA) and Sabouraud Dextrose Agar (SDA) were used in bacteria and yeast cultures on 96-well plates, respectively. The positive controls used in the study were streptomycin (for bacterial testing) and cycloheximide (for yeast testing), which were provided by the Institute of Drug Quality Control in Ho Chi Minh City, Vietnam.

The experiment was performed according to the following steps: (1) The EO samples were diluted in 10% DMSO to prepare eight different concentrations (2-256 μ g/mL); (2) The bacteria and fungi (yeast) were standardized with a concentration of 2×10^5 CFU/mL; (3) the mixtures of microbial and testing samples were incubated at 37°C (18-24 h) for bacteria and at 35-37°C (36-48 h) for yeast. The minimum inhibitory concentration (MIC) value was determined as the lowest concentration that completely prevented the visible *in vitro* microbial growth after 24 h of culture. This is accomplished by spreading cultures out on an agar plate. All experiments were done three times, in duplicates.

Results and Discussion Chemical composition analysis of the EOs

The GC-MS analysis detected 37 compounds in the rhizome EO of *Z. densissimum* (Fig. S2), of which monoterpene hydrocarbon was the largest chemical group, with 85.40% of the total content, followed by oxygenated sesquiterpenes,

sesquiterpene hydrocarbons, diterpenes, and oxygenated monoterpenes based on peak area in the GC chromatogram. β-Pinene was found to be the most dominant component, accounting for 38.36% of the total content, followed by β-phellandrene (26.85%) and α-pinene (13.31%). Noticeably, these three compounds belong to the monoterpene hydrocarbon group ([Table 1](#page-4-0)).

Similarly, the GC-MS analytical results revealed that a total of 52 components in the rhizome EO of *K. laotica* (Fig. S3), contained mainly monoterpene hydrocarbons (61.09%), followed by sesquiterpene hydrocarbons (15.18%), oxygenated monoterpenes (7.24%), oxygenated sesquiterpenes (4.62%) and diterpenes (0.47%). Camphene was found to be the most dominant volatile compound in the EO sample with a percentage of 23.44% based on peak area in the GC chromatogram. The EO from *K. laotica* rhizomes also contained a significant amount of α-pinene (10.69%) and β-pinene (6.85%) while a considerable amount of 3-carene was also detected (8.60%). Similar to the EO prepared from *Z. densissimum* rhizomes, all major components from the EO of *K. laotica* rhizomes belong to the monoterpene hydrocarbon group [\(Table 2](#page-5-0)).

The EOs of the most common ginger species, *Z. officinale*, contained mainly α-zingiberene, arcurcumene, and geranial, which indicated the vast difference between the chemical composition of the EOs of *Z. densissimum* and *Z. officinale*¹⁰. When compared with other *Zingiber* species, the EO of *Z. densissimum* was shown to be very different in terms of chemical composition. Specifically, the EOs of *Z. montanum* and *Z. cassumunar* contained sabinene (37- 56%), terpinen-4-ol (7-30%), and (*E*)-1-(3-4 dimethoxyphenyl) butadiene $(11-16\%)^{11,12}$ while *Z. nimmonii* EO was found to contain myrcene, α-caryophyllene, β-caryophyllene, α-humulene, and α -cadinol¹³.

As for the *Kaempferia* genus, the EO of *K. galanga* contained ethyl-p-methoxy-cinnamate, ethyl cinnamate, 1,8-cineole, borneol, camphene, linoleoyl, methyl-cinnamate, and pentadecane¹⁴. The EO of *K. angustifolia* collected in Thailand was found to contain camphene (27.98%), camphor (18.73%) and α-pinene (7.42%) while

Table 1 *cont.*

RT: Retention time (min); RI (obsd.): Retention indices calculated in the investigation; RI (lit.): Retention indices from literature

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K. marginata contained δ-3-carene (33.84%), β-pinene (12.21%) and camphene (11.51%) in its EO. Therefore, it was observed that the chemical composition of *K. laotica* EO, to some extent, was similar to those of *K. angustifolia* and *K. marginata* with camphene, α-pinene, β-pinene, and 3-carene as the most dominant compounds¹⁵. Compared to other species, camphor was found to be the main component in *Artemisia herba-alba* and *Lavandula dentata*, borneol in *Origanum vulgare* and *Thymus vulgaris*, and 1,8-cineole in *Rosmarinus officinalis*¹⁶.

Antimicrobial activity of the EOs

The EOs were evaluated for their antimicrobial activities against some pathogenic bacterial and fungal strains ([Table 3](#page-7-0)). Streptomycin and cycloheximide were used as positive controls, respectively. The testing results indicated that the rhizome EO of *Z. densissimum* strongly inhibited the growth of *C. albicans,* with a MIC value of 2 µg/mL, which was considerably lower than that of cycloheximide. As for antibacterial activities, the *Z. densissimum* rhizome EO only showed moderate activities,

with the MIC values ranging from 8 to $128 \mu g$ / mL. Specifically, *E. faecalis*, *S. aureus*, and *E. coli* were the most affected trains*,* with the MIC values of 8 µg/mL, while *B. cereus* and *S. enterica* were less sensitive to the EO, with MIC values of 16 and 32 μ g/mL, respectively. Noticeably, the EO only showed weak effects against the growth of *P. aeruginosa* with MIC value of 128 µg/mL.

In contrast, the EO of *K. laotica* rhizomes showed generally weaker antimicrobial activities in comparison with that of *Z. densissimum* rhizomes. Specifically, the most susceptible microbial strain with *K. laotica* rhizome EO was *B. cereus* (MIC = 32 μ g/mL), followed by *S. enterica* (MIC = 64 µg/mL), *E. coli* and *C. albicans* (MIC = 128 µg/mL), *S. aureus* and *P. aeruginosa* (MIC = $256 \mu g/mL$). As for *E. faecalis* strain, the MIC value could not be detected due to the weak activity.

In comparison with other species in the Zingiberaceae family, the rhizome EO from *Z. densissimum* showed more pronounced activity, while that from *K. laotica* was in similar ranges. The EO from *Hedychium yunnanense* rhizomes was reported to display antimicrobial activities on the same microbial strains, with MIC values of $32-64 \mu g/mL^{17}$. In another study, the EO from *K. attapeuensis* rhizomes showed the MIC values around 32 µg/mL against all tested microbial strains except for *P. aeruginosa*¹⁸.

The antimicrobial effects of the EOs are related to the major compounds and their synergy

with other minor ones present in the EOs. The major monoterpene hydrocarbons from the two plants' EOs were shown to be effective against a variety of microbial strains $19-21$, which explained the pharmacological results in the present study. β-Pinene was shown to display potent anti-fungal effects by interfering with the cell wall and reducing the *Candida* biofilm adhesion²². Camphene was effective against most dermatophytes, which are responsible for superficial skin infections, by changing the fatty acid composition of cell membrane, leading to inhibition of respiration and changes of cellular permeability²³. 3-Carene, a main component of pine and pepper volatile oils, was found to possess strong bactericidal effects against *Brochothrix thermosphacta* and *Pseudomonas fluorescens* through cell wall permeability changes, DNA structure disruption, and interfering with the metabolic processes 24 . α-Pinene, a monoterpene abundantly found in pine and some rosemary²⁵, was studied for its synergistic effects with antibiotics in the combat against antibiotic resistance. The monoterpene was shown to prevent the antimicrobial efflux of *Campylobacter jejuni* (the main cause of gastroenteritis), which was displayed by reducing the MIC value of concurrent antibiotics, such as ciprofloxacin, erythromycin and triclosan, up to 512 times²⁶. α-Pinene was also found to act via NorA efflux pump inhibition against *S. aureus* and significantly enhance the potential of norfloxacin²⁷.

Conclusion

The present study reveals the chemical composition of EOs from the two understudied ginger species, *Z. densissimum* and *K. laotica*, by the GC-MS analysis, in which β-pinene (38.36%), β-phellandrene (26.85%), and α-pinene (13.31%) were found the most abundant in *Z. densissimum* whereas the major compositions of *K. laotica* rhizome essential oil were α-pinene (10.69%), β-pinene (6.85%), and 3-carene (8.60%). The study results have shown that the EO of *Z. densissimum* rhizomes exhibited strong inhibitory activities against some microbial strains, specifically *E. faecalis*, *S. aureus*, *E. coli* (MIC value = 8 µg/mL), and *C. albican*s (MIC = 2 µg/mL) while *K. laotica*'s rhizome EO only showed weak antimicrobial effects against all investigated species (MIC values ranging from $32-256 \mu g/mL$). The study findings should encourage further scientific work on these two gingers, to clarify their phytochemical profiles and other pharmacological activities.

Conflict of Interest

No potential conflict of interest was reported by the authors.

Supplementary data

Figures S1 to S3 are given in supplementary file.

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