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Chemical composition of fennel essential oil and its impact on *Staphylococcus aureus* exotoxin production

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Abstract In this study, fennel oil was isolated by hydrodistillation, and the chemical composition was determined by gas chromatography/mass spectral analysis. The antimicrobial activity of fennel oil against Staphylococcus aureus was evaluated by broth microdilution. A haemolysis assay, tumour necrosis factor (TNF) release assay, western blot, and real-time reverse transcription (RT)-PCR were applied to investigate the influence of fennel oil on the production of S. aureus virulence-related exoproteins. The data show that fennel oil, which contains a high level of trans-anethole, was active against S. aureus, with MICs ranging from 64 to 256 µg/ml. Furthermore, fennel oil, when used at subinhibitory concentrations, could dosedependently decrease the expression of S. aureus exotoxins, including α -toxin, Staphylococcal enterotoxins (SEs) and toxic shock syndrome toxin 1 (TSST-1).

Keywords Staphylococcus aureus · Fennel essential oil · Antimicrobial activity · Subinhibitory concentrations · Virulence factors

Jiazhang Qiu and Hongen Li contributed equally to this work.

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Introduction

Staphylococcus aureus is a major medical pathogen that causes a wide variety of infections, from simple abscesses to fatal sepsis, as well as toxinoses, such as food poisoning and toxic shock syndrome (Lowy 1998). Its pathogenic versatility is largely attributed to its ability to produce and secrete a number of virulence factors, including superantigen toxins, haemolytic cytotoxins, and surface proteins.

 α -toxin is a pore-forming haemolytic toxin that causes membrane damage to many types of mammalian cells. The α -toxin monomers bind to cell membranes and then associate into a heptameric complex to form a pore (Song et al. 1996). Staphylococcal enterotoxins (SEs) and toxic shock syndrome toxin 1 (TSST-1), which are known as pyrogenic toxin superantigens (PTSAgs), share certain structural and biological properties and stimulate the proliferation of T lymphocytes and the release of T-cell-derived cytokines (e.g., TNF- α) (Dinges et al. 2000). These toxins were initially implicated in staphylococcal food poisoning (SEs) and TSS (TSST-1).

Because *S. aureus* develops resistance to new antibiotics almost as fast as they are introduced, there is a continuous demand to find new and improved antimicrobial agents or to develop alternative therapeutic strategies in both the food and pharmaceutical industries. One of the areas that is currently the subject of considerable interest is plant extracts, and particularly their essential oils, for their potent antimicrobial properties against a broad spectrum of microorganisms (Smith-Palmer et al. 1998). Fennel (*Foeniculum vulgare* Miller) is a major aromatic plant belonging to the *Apiaceae* family, which has long been considered as a medicinal and spice herb. Fennel oil is commonly used as a flavouring agent in food products and as a constituent of cosmetic and pharmaceutical products; furthermore, its antimicrobial properties against a wide range of microorganisms have been well established (Elgayyar et al. 2001; Aprotosoaie et al. 2008). It has long been known that some plant essential oils could affect the expression of staphylococcal exotoxins (Smith-Palmer et al. 2004). Consequently, the present study aimed to evaluate the influence of subinhibitory fennel oil concentrations on the expression of α -toxin, the two major enterotoxins (SEA and SEB), and TSST-1 in *S. aureus*.

Materials and methods

Bacterial strains

The methicillin-susceptible *S. aureus* strain ATCC 29213 was obtained from ATCC. Nineteen *S. aureus* clinical isolates (6 MSSA and 13 MRSA) were acquired from the First Hospital of Jilin University; the clinical MRSA strains 2985 and 3701, which have the ability to produce α -toxin, SEA and SEB, and TSST-1, were selected for further tests. Bacteria were stored as a 30% glycerol stock at -80° C until testing.

Plant material and essential oil extraction

Fennel (*Foeniculum vulgare* Miller) was collected from the Guangxi Province of China in July 2009. The plant material was air-dried in a shady and aerated room until the weight was stable. The fennel was crushed, and the fennel oil was extracted by hydrodistillation for 4 h using a Clevenger-type apparatus. The essential oil was dried over anhydrous sodium sulphate.

Gas chromatography/mass spectral analysis

The chemical composition of the fennel oil was detected by GC/MS performed on a GCMS-QP2010 Plus Gas Chromatograph/Mass Spectrometer (Shimazu Co., Ltd., Kyoto, Japan) equipped with a fused silica capillary column (Rix-5 ms; 30 m × 0.25 mm × 0.25 μ m; Shimadzu, Kyoto, Japan). The carrier gas was helium (1 ml/min). The oven temperature was maintained at 60°C for 10 min and programmed to reach 250°C at a rate of 10°C/min. The split ratio was adjusted to 100:1, and the injector temperature was set at 280°C. Mass spectra were obtained by electronic impact at 70 V, and the mass range was from *m*/*z* 50 to 500. Identification of the constituents was performed by a computer-based library search.

MIC determination

Stock solutions of fennel oil at different concentrations were prepared in dimethyl sulphoxide (DMSO) (Sigma-

Aldrich). The MICs (minimum inhibitory concentrations) of fennel oil against 20 *S. aureus* strains were evaluated using a broth microdilution method as described by Carson et al. (1995), with minor modifications. All tests were performed in Mueller–Hinton broth (MHB) (BD Biosciences, Inc., MD, USA) supplemented with Tween-80 (Sigma-Aldrich) at a final concentration of 0.5%. Serial doubling dilutions of fennel oil were prepared in a 96-well plate over the range of $8-1,024 \mu g/ml$. Following the inoculation of 5×10^5 cfu/ml of overnight broth cultures into each well, the plates were incubated aerobically at 37° C for 24 h. The MIC was defined as the lowest concentration of fennel oil at which the bacteria do not demonstrate visible growth.

Growth curve assay

An overnight culture of *S. aureus* was diluted into 1000 ml fresh MHB supplemented with 0.5% Tween-80 and grown at 37°C with shaking at 200 rpm to obtain a starting OD ₆₀₀ value of 0.3, and 100-ml volumes of the culture were aliquoted into six 250-ml Erlenmeyer flasks. Five of the flasks were supplemented with fennel oil (dissolved in DMSO) at concentrations of 6.25, 12.5, 25, 50 and 100% MIC. Bacteria were further cultured at 37°C with agitation at 200 rpm under aerobic conditions, and the growth of the cells was monitored by measuring the OD ₆₀₀ values at the indicated time points.

Haemolysis assay

The haemolysis assay was performed as previously described (Qiu et al. 2010b) with rabbit erythrocytes. In brief, bacteria were grown in MHB supplemented with 0.5% Tween-80 at 37°C with graded subinhibitory concentrations of fennel oil until reaching the post-exponential growth phase (OD_{600 nm} of 2.5). Culture supernatants were collected and were filter sterilised with a 0.22-µm (pore size) acetate syringe filter. A 0.1-ml volume of culture supernatant was mixed with 2.5% defibrinated rabbit blood in PBS buffer. After 15 min at 37°C, the unlysed blood cells were pelleted by centrifugation $(5,500 \times g, room$ temperature, 1 min). The haemolytic activity was determined by measuring the optical density (at 543 nm) of the cell-free supernatant. The control culture supernatant served as the 100% hemolysis control, and the % hemolysis was calculated by comparison to the control culture.

Tumour necrosis factor (TNF) release assay

The TNF release assay was performed by an established method described by Qiu et al. (2010a). In short, overnight bacterial cultures grown in RPMI-1640 (Invitrogen, CA,

USA) were incubated into fresh, prewarmed RPMI-1640 medium (500 ml) supplemented with 0.5% Tween-80. Following incubation at 37°C for 30 min with aeration, cultures were divided into aliquots of 100 ml. Increased concentrations of fennel oil were added to bacterial suspensions, and the cultures were further incubated for 4 h at 37°C with constant shaking. S. aureus supernatants were collected by centrifugation, filtered through a 0.2-µm filter and immediately analysed as described below.

The animal studies were conducted in accordance with the experimental practices and standards approved by the Animal Welfare and Research Ethics Committee at Jilin University. Specific-pathogen-free BALB/c mice (male, 6-8 weeks old, weighing 18-22 g) were supplied by the Experimental Animal Center of Jilin University (Changchun, China). Mice were euthanised by cervical dislocation. Single spleen cell suspensions were prepared in RPMI-1640, washed and resuspended in complete RPMI-1640 medium. A total of 10^6 (150 µl) cells were added to wells of 96-well plat tissue culture plates and then supplemented with 50 µl of S. aureus culture supernatants. After incubation for 16 h at 37°C, the supernatants were harvested by centrifugation $(1,000 \times g \text{ for 5 min})$. TNF in the supernatants was measured using the Mouse TNF-a ELISA MAXTM Standard Set (Biolegend, Inc., San Diego, USA).

Western blot assay

Bacteria were cultured, and supernatants were collected in the same fashion as for the haemolysis assay. Proteins in equal volumes of culture supernatants (20 µl) were subjected to sodium dodecyl sulphate (SDS)-polyacrylamide (12%) gel electrophoresis at 120 V. Proteins were transferred to polyvinylidene fluoride membranes (Wako Pure Chemical Industries, Ltd, Osaka, Japan) using a semidry transfer cell (Bio-Rad, Munich, Germany). Membranes were incubated overnight at 4°C in 10% milk powder as a blocking reagent. The production of α -toxin, SEA, SEB, and TSST-1 in S. aureus was detected by incubation with the indicated antibodies. Antibodies to SEA, SEB, and α hemolysin were purchased from Sigma-Aldrich and diluted to 1:10000, 1:5000 and 1:8000, respectively; then, a horseradish peroxidase-conjugated anti-rabbit antiserum (Sigma-Aldrich) diluted to 1:4000 was used as the secondary antibody. The antibody to TSST-1 (Santa Cruz Biotechnology, California, USA) was diluted to 1:200 according to the manufacturer's recommendations; then, a horseradish peroxidase-conjugated anti-mouse antiserum (Sigma-Aldrich) diluted to 1:5000 was used as the secondary antibody. The blots were developed using ECL substrate (GE Healthcare, Buckinghamshire, UK).

Proteolytic activity assay

Samples of culture supernatant (100 µl) were added to 1 ml of azocasein (Sigma-Aldrich; 1 mg/ml in 100 mmol Tris-HCl, PH 7.2) and incubated at 37°C for 1 h. The reaction was terminated by addition of 1 ml of trichloroacetic acid (5%, w/v) and mixing; undigested azocasein was allowed to precipitate for 30 min. The mixture was then centrifuged at $10,000 \times g$ for 10 min and the absorbance of the supernate read at 328 nm. One unit of protease activity was defined as giving an absorbance of 0.001 after incubation for 1 h at 37°C.

RNA isolation and real-time RT-PCR

The S. aureus strain ATCC 29213 was incubated with or without the addition of 50% MIC of fennel oil to the postexponential growth phase (OD_{600 nm} of 2.5), as described in the haemolysis assay. Total RNA was isolated using the RNeasy Mini kit (Qiagen, Hilden, Germany) in accordance

imers used for real- R	Primer	Sequence	Genomic location
	16S rRNA-fw	5'-GCTGCCCTTTGTATTGTC-3'	287-305
	16S rRNA-rv	5'-AGATGTTGGGTTAAGTCCC-3'	446-465
	hla-fw	5'-TTGGTGCAAATGTTTC-3'	485-501
	hla-rv	5'-TCACTTTCCAGCCTACT-3'	569-586
	sea-fw	5'-ATGGTGCTTATTATGGTTATC-3'	335-356
	sea-rv	5'-CGTTTCCAAAGGTACTGTATT-3'	477-498
	seb-fw	5'-TGTTCGGGTATTTGAAGATGG-3'	480-501
	seb-rv	5'-CGTTTCATAAGGCGAGTTGTT-3'	612-633
	<i>tst</i> -fw	5'-ACCCCTGTTCCCTTATCATC-3'	73-93
	tst-rv	5'-AAAAGCGTCAGACCCACTAC-3'	159-180
	agrA-fw	5'-TGATAATCCTTATGAGGTGCTT-3'	111-133
	agrA-rv	5'-CACTGTGACTCGTAACGAAAA-3'	253-274

Table 1 Pr time RT-PC

with the manufacturer's directions. The contaminating DNA was removed using the optional on-column RNasefree DNase I step (Qiagen, Hilden, Germany). RNA was reverse transcribed into cDNA using the Takara RNA PCR kit (AMV) Ver. 3.0 (Takara, Kyoto, Japan), according to the manufacturer's protocol. The genes and primer sequences employed for the real-time RT-PCR analysis are listed in Table 1. The real-time PCR was performed using the 7000 Sequence Detection System (Applied Biosystems, Courtaboeuf, France) and SYBR Premix Ex Taq (Takara). Reaction mixtures were initially incubated for 30 s at 95°C, followed by 35 cycles of 5 s at 95°C, 30 s at 55°C, and 20 s at 72°C. A melt-curve analysis was also carried out to assess PCR specificity and resulted in single primerspecific melting temperatures. All samples were analysed in triplicate, and the housekeeping gene 16S rRNA was used as an endogenous control. In this study, relative quantification based on the relative expression of a target gene versus the 16S rRNA gene was utilised to determine the transcript level changes between samples.

Statistical analysis

Statistical analysis was performed with SPSS 12.0 statistical software. The data are presented as the mean values \pm SD (n = 3). An independent Student's *t*-test was used to analyse the data. Differences were considered statistically significant when the *p* value was less than 0.05.

Results

The chemical composition of fennel oil is shown in Table 2. Fennel oil contains a high level of *trans*-anethole (88.91%).

Table 2 The main components and their relative contents (%) of fennel oil

Compound	Retention time (min)	Relative contents (%)	
<i>m</i> -cymene	5.667	0.37	
D-limonene	5.750	1.64	
Eucalyptol	5.833	0.13	
γ-terpinen	6.183	0.34	
L-fenchone	6.750	0.7	
Anisole	8.992	2.89	
Camphor	9.092	0.35	
Cis-anethole	9.175	0.25	
Anisaldehyde	9.592	2.54	
Trans-anethole	10.825	88.91	
p-acetonylanisole	11.283	0.31	
Apiole	14.300	0.15	



Fig. 1 Growth curves for *S. aureus* ATCC 29213 grown in MHB with increasing concentrations of fennel oil

The MIC of fennel oil for each of the 20 S. aureus strains was evaluated and ranged from 64 to 256 µg/ml. The MIC values of fennel oil against S. aureus ATCC 29213, MRSA 2985 and MRSA 3701 were 128 µg/ml. As shown in Fig. 1, fennel oil at concentrations from 6.25 to 50% MIC had no significant influence on the growth of S. aureus strain ATCC 29213. However, when supplemented with 100% MIC of fennel oil, the growth rate was significantly decreased; after 30, 180 and 360 min of fennel oil treatment, the OD₆₀₀ values were 54.5, 53.1 and 58.9% of the fennel oil-free culture, respectively. Although the growth kinetics can vary greatly between strains, the growth of MRSA 2985 and MRSA 3701 were affected by these concentrations of fennel oil in a similar manner. In other words, the addition of 6.25, 12.5, 25, and 50% MIC of fennel oil had no significant effects on MRSA 2985 and MRSA 3701 growth (data not shown).

As shown in Table 3, when supplemented with 6.25% MIC of fennel oil, the haemolysis of *S. aureus* strains ATCC 29213, MRSA 2985 and MRSA 3701 culture supernatants were 66.7, 55.8 and 42.5% of their drug-free cultures, respectively. Remarkably, no haemolytic activities were observed when cultured with 50% MIC of fennel oil. This dose-dependent inhibition of haemolysis was observed in all of the investigated strains. As shown in Fig. 2, the culture supernatants of *S. aureus* grown in the presence of graded subinhibitory concentrations of fennel oil stimulated significantly lower levels of TNF- α release. In addition, fennel oil itself did not stimulate or inhibit TNF- α production at 100% MIC (data not shown). Apparently, fennel oil repressed the TNF-inducing activity of *S. aureus* culture supernatants in a dose-dependent fashion.

Figure 3 shows the α -toxin, SEA and SEB, and TSST-1 levels in the *S. aureus* culture supernatants after exposure

Strains	Haemolysis (%) of rabbit erythrocytes by culture supernatant ^a					
	0	6.25 MIC	12.5 MIC	25 MIC	50 MIC	
ATCC 29213	100	66.7 ± 3.8	25.9 ± 5.2**	$6.4 \pm 2.3^{**}$	None ^b	
MRSA 2985	100	55.8 ± 5.7	$45.5 \pm 4.6^{*}$	$13.7 \pm 5.1^{**}$	None	
MRSA 3701	100	$42.5 \pm 5.4*$	$15.8 \pm 7.2^{**}$	None	None	

Table 3 Haemolytic activities in the culture supernatants of S. aureus in the presence of various concentrations of fennel oil

^a Haemolytic activity in fennel oil-free S. aureus culture supernatants was set to 100%

^b No haemolytic activity was detected

Values are means \pm SD (n = 3). *indicates P < 0.05 and **indicates P < 0.01 compared to the corresponding control



Fig. 2 TNF- α release from spleen cells stimulated with *S. aureus* culture supernatants grown in RPMI 1640 with graded subinhibitory concentrations of fennel oil. *Error bars* (n = 3) indicate standard deviation, *indicates *P* < 0.05 and **indicates *P* < 0.01

to fennel oil. Treatment with increasing concentrations of fennel oil resulted in a dose-dependent decrease in the production of α -toxin, SEA and SEB, and TSST-1. Growth with 6.25% MIC of fennel oil resulted in a recognisable reduction in the secretion of these toxins, whereas at 50% MIC, no or little immunoreactive proteins could be detected in all of the strains tested.

There was no significant influence on protease secretion by ATCC 29213, MRSA 2985 or MRSA 3701 cultured with graded subinhibitory concentrations of fennel oil (Fig. 4).

Table 4 shows that treatment with 50% MIC of fennel oil significantly decreases the transcription levels of *hla*, *sea*, *seb*, *tst*, and *agrA* of the *S. aureus* strain ATCC 29213.

Discussion

The increasing emergence of multi-drug-resistant pathogens has intensified the need to find novel antimicrobial agents for the prevention and treatment of bacterial infections. In recent decades, plant essential oils have been gaining great attention, and their antibacterial activities against a broad spectrum of micro-organisms have been well established (Smith-Palmer et al. 1998, Smith-Palmer 1999; Valero and Salmeron 2003). Due to their multicomponent nature, plant essential oils are more difficult for bacteria to develop resistance to than many common used antibiotics, which have a single target site (Smith-Palmer et al. 2004). In the last few years, many studies have been conducted in different countries to demonstrate the significance of these oils in therapeutic treatments (Benoit-Vical et al. 2006; Senatore et al. 2007). In the present study, fennel oil was active against both MSSA and MRSA, with MICs ranging from 64-256 µg/ml. The results were consistent with previous studies (Mohsenzadeh 2007: Aprotosoaie et al. 2008), indicating that fennel oil is a potentially effective antimicrobial agent against S. aureus, and it may deserve further investigation for its potential therapeutic efficacy in S. aureus infections.

Previous studies have indicated that the expression of S. aureus exotoxins could be influenced by subinhibitory concentrations of antimicrobial agents. Therefore, the antibiotic-induced modulation of virulence factors may result in either aggravation or attenuation of the infection. For example, some β -lactam antibiotics strongly induce the production of virulence-related exoproteins (Ohlsen et al. 1998), suggesting that the symptoms of S. aureus infections may be intensified when patients are treated with these antibiotics. In contrast, some protein-synthesis-suppressing antibiotics, such as clindamycin, linezolid, and quinupristin/dalfopristin, have been shown to disrupt the expression of S. aureus virulence factors (Herbert et al. 2001; Bernardo et al. 2004; Koszczol et al. 2006). As a consequence, these antibiotics are recommended for the management of S. aureus-produced toxic syndromes. Furthermore, it has also been demonstrated that some plant essential oils (e.g., oils of cinnamon, bay and clove) can influence production of exotoxins when used at sub inhibitory concentrations (Smith-Palmer et al. 2004). The data presented here show the ability of subinhibitory concentrations of fennel oil to **Fig. 3** Western blot analysis of α -toxin, TSST-1, SEA and SEB production by the *S. aureus* strains ATCC 29213 (**a**), MRSA 2985 (**b**) and MRSA 3701 (**c**) after growth in the presence of increasing concentrations of fennel oil



(a)

Fig. 4 Protease units of *S. aureus* culture supernatants. Values represent the mean \pm SD for three independent experiments

 Table 4
 Relative expression levels of *hla, sea, seb, tst* and *agrA* in *S. aureus* ATCC 29213 after treatment with 50% MIC of fennel oil

Gene	Protein	Fold changes \pm SD ^a
hla	α-toxin	-9.2 ± 2.9
sea	Staphylococcal enterotoxin A	-7.7 ± 2.1
seb	Staphylococcal enterotoxin B	-10.5 ± 3.4
tst	Toxic shock syndrome toxin-1	-6.9 ± 2.5
agrA	Accessory gene regulator A	-6.2 ± 1.9

 $^{\rm a}$ Means reduction; values were presented as mean \pm SD of three independent experiments

cause a significant decrease in the production of major exotoxins by *S. aureus*, indicating that fennel oil may be useful for the treatment of *S. aureus* infections when used in combination with β -lactam antibiotics. Furthermore, as fennel oil contains extraordinarily high amount of *trans*-anethole, it might be possible that the effects seen are due mainly to *trans*-anethole.

Considering the tendency of consumers to avoid foods containing chemical and artificial preservatives,



investigators are pursuing natural antimicrobial substances from plant sources, especially their essential oils (Lee et al. 2002). The potential application of essential oils to foods has been well investigated in recent years (Smith-Palmer et al. 2001; Valero and Salmeron 2003). In addition to the inhibition of growth of bacterial cells, researchers are also interested in the inhibition of toxin production. For example, food-borne staphylococcal poisoning is caused by the ingestion of one or more enterotoxins that were pre-formed in foods contaminated with *S. aureus*. More importantly, SEs are resistant to treatment with heat, strong acid and alkali. Therefore, the ability of fennel oil to inhibit the production of staphylococcal enterotoxins may increase the likelihood of fennel oil being applied as a novel natural food preservative.

Exoprotein expression by S. aureus is co-ordinately controlled by numerous global regulators that act at the transcriptional level (Booth et al. 1997). Previous studies have indicated that sublethal concentrations of antibiotics could affect the translation of certain regulatory gene products in S. aureus, which, in turn, influence the transcription of toxin-encoding genes (Herbert et al. 2001; Qiu et al. 2010b). Therefore, it would be reasonable to infer that the fennel oil-induced inhibition of global regulators might lead to the decreased production of exotoxins. The agr locus is one of the well-characterised staphylococcal global regulators and enhances the post-exponential-phase expression of secreted proteins (e.g., a-toxin, SEB and TSST-1) (Arvidson and Tegmark 2001). The real-time RT-PCR data indicated that the transcriptional level of agrA in S. aureus strain ATCC 29213 was significantly decreased after treatment with 50% MIC of fennel oil. The regulatory mechanism by which S. aureus controls the expression of virulence factors is extremely complicated, involving an interactive, hierarchical regulatory cascade among the gene products of agr and sar as well as other regulators (Chan and Foster 1998). Therefore, we presume that the reduced production of *a*-toxin, SEB and TSST-1 may, in part, depend on the inhibition of the agr locus induced by fennel oil. The expression of SEA is not controlled by the agr regulatory system, and the action of regulatory genes on SEA production in S. aureus is still unclear (Arvidson and Tegmark 2001). Therefore, it is definite that the impact of fennel oil on SEA production cannot be mediated through influence on *agr*, and the regulatory mechanism that governs fennel-induced reduction of SEA production remains to be determined.

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Conflict of interest The authors declare that there have no conflict of interest.

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