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### EVALUATION OF THE ANTIBACTERIAL POTENTIAL OF DESERT TRUFFLES (*Terfezia spp*) EXTRACTS AGAINST METHICILLIN RESISTANT *Staphylococcus aureus* (MRSA)

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#### KEYWORDS

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Crude extracts

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*Terfezia claveryi*

*Terfezia nivea*

#### ABSTRACT

Desert truffles are well known for its medicinal and functional food values. These mushrooms are widely used for its antibacterial properties and as a source of protein in Saudi Arabia and many other countries around the world. Methicillin Resistant *Staphylococcus aureus* (MRSA) has been known for its virulence and multidrug resistant that became threat for hospital employees and community members. Present study was designed to test the common occurrence of MRSA isolates in hospital employees and patients by nasal swab samplings and then evaluate the antibacterial potential of desert truffles against these MRSA. PCR assays revealed that 17 (34%) bacterial isolates of nasal carriage found MRSA positive while in case of gender no significant association was reported in MRSA positive males and females. Further it was also reported that highest rate of MRSA positive isolates was found for the age group of 20-29 years, among the youngest participants. For testing antibacterial potential of Desert truffles, two wells of 6 mm diameter were punched on Muller-Hinton Agar that already had MRSA culture, one of them filled cautiously with 100µl of *Terfezia claveryi* extract and the other one with 100µl of *Terfezia nivea*, stored at 4°C for 2 hours. After incubation at 37°C for 24 h, the fresh crude extracts of *T. claveryi* showed a significant zone of inhibition (27±7.75mm) while this inhibition was reported (22.5±11.26mm) in case of *T. nivea*. These findings clearly indicated that extracts has remarkable

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antibacterial activity against MRSA. This investigation supports the traditional use of truffles for the treatment of eye infection and proved its effectiveness against MRSA which could be considered as promising antibiotic drug in near future.

## 1 Introduction

Now in these days use of antibiotics has frequently increased. The excessive and improper use of antibiotics not only increasing the resistance in microorganisms but also adversely affect the immune system. The severe side effects of the currently used antibiotics and its high production cost pushed researchers to find out new alternative source of antibiotics which should be safe and effective. Ancient literatures suggest the use of desert truffles as an alternative of commercial antibiotics (Wang & Marcone, 2011; Patel et al., 2017; Schillaci et al., 2017).

Desert truffles are hypogeous fungi that have been used as medicine and food for centuries in Saudi Arabia and many other countries around the world (Lo & Kam, 2006; Al-Qarawi & Mridha, 2012; Charoensiddhi et al., 2017). According to the local experts, these truffles grew in very rare environment when limited rain occurred between February to April. When there is more rain falls, the chances to have high yield of truffles increased. After 40-50 days of the first rain fall people start searching truffles in cracks formed on the ground (Sawaya et al., 2006; Hamza et al., 2016a). The interactions between growing truffles are very vital in ecological systems as the mycorrhizal species have the potential to avoid erosion and desertification by sand stabilization. Mycorrhizal fungi also modify water relations in host plants (Benzeggouta, 2014). Among various reported species of desert truffles two types "*T. nivea* and *T. claveryi*" are very common in Saudi Arabian climatic conditions. Among these two, *T. nivea* is locally known as Zubaidi and had a meat like texture with white colored skin while *T. claveryi* is similar to the potatoes in texture and have brown colored skin and locally known as Khalasi. Both of these belong to the family of *Terfeziaceae* and were used in the current research.

The importance and chemical composition of desert truffles has been well documented particularly in their proteins and amino acids contents which is higher than many other edible mushrooms. It proved highly nutritious and digestible for humans and comprises approximately 85% amino acids, 27% protein, 60% carbohydrates (glycerol, glucose, fructose, mannitol, inositol, and trehalose in varying quantities), 2 to 5% ascorbic acid, 3 to 7.5 % fat (unsaturated and saturated fatty acids), and 7 to 13% crude fiber (Dundar et al., 2012; Garcia-Vaquero et al., 2017).

The extracts of desert truffles have anti-bacterial properties against a wide range of bacteria and widely used in the treatment

of trachoma (Alhussaini et al., 2016). It is also successfully used against Gram-positive human pathogenic reference strain *Staphylococcus aureus* ATCC 29213 and Gram-negative strain *Pseudomonas aeruginosa* ATCC 15442 (Janakat et al., 2005; Al-Qarawi & Mridha, 2012; Pádua et al., 2015; Neggaz et al., 2015; Schillaci et al., 2017). Gouzi et al. (2011) tried aqueous extract of the truffles against *Staphylococcus aureus* and *Pseudomonas aeruginosa* and exhibits remarkable zone of inhibition. Desert truffles comprise a vast unexploited therapeutic compounds such as phenolics, tocopherols, ascorbic acid and carotenoids which have remarkable medicinal properties along with antioxidant potential (Al-Laith, 2010; Balboa et al., 2013; Özyürek et al., 2014; Sanjeeva et al., 2016; Gargano et al., 2017; Pinteus et al., 2017).

Desert truffles have nutritional value due to their proteins, carbohydrates, fats, fibers, and low energy. They also have immune-modulating, hepatoprotective, antidepressant, antibacterial, antifungal, antiviral, antioxidant, and antiradical properties due to their content of phenol, carotenoid, anthocyanin, ascorbic acid, flavonoid, tannin, glycoside, ergosterol, etc. Hence, the introduction of desert truffles in the pharmacological field is important, especially in the treating of eye infections and cancer (Janakat & Nassar, 2010; Hamza et al., 2016b; Khadri et al., 2017; Owaid, 2018). Moreover, studies on higher mushrooms showed antibacterial activity against *Escherichia coli*, *Pseudomonas aeruginosa* and *Staphylococcus aureus* (Imtiaj & Lee, 2007).

*Staphylococcus* is common nasal bacteria can cause serious infectious diseases in skin and other mucous membranes and at severe infections it might cause lung infection and pneumonia. Methicillin resistant *S. aureus* (MRSA) usually developed and showed resistance against beta-lactam antibiotics. One of the most effective methods for preventing the spread of MRSA requires detection of colonized HCWs and measuring the associated risk factors of colonization (Kaur & Chate, 2015). MRSA is strain of *Staphylococcus aureus* that developed by multidrug resistance to  $\beta$ -lactam antibiotics which include Penicillin, Methicillin, dicloxacillin, nafcillin, Oxacillin (Appelbaum, 2007; Kaur & Chate, 2015). The first report of methicillin resistant *S. aureus* (MRSA) was from London in 1961 (Cookson, 2011). This bacterium is capable in causing severe serious infections in humans forming skin infections such as boils and abscesses (Hiramatsu et al., 2014). They can also burrowed deep into the body, causing potentially life-threatening infections in bones, joints, surgical wounds, the bloodstream, heart valves and lungs (MayoClinic,

2015). The virulence of this *S. aureus* comes from its ability to adapt to a variety of changing environmental conditions and to modulate its pathogenicity. It can establish asymptomatic carriage, which permits widespread dissemination among human hosts. It also has a remarkable proclivity to acquire resistance to multiple antimicrobial agents, which causes therapeutic challenges for physicians (Moellering, 2010). It has been suggested that the *mecA* gene is responsible for *S. aureus* resistance to methicillin. *MecA* encodes an altered penicillin-binding protein (PBP2a) with a low affinity for  $\beta$ -lactam antibiotics (Grundmann et al., 2006). MRSA was previously considered as a nosocomial pathogen, but in the past two decades, reports suggest an increasing trend for community-associated MRSA (CAMRSA). These clones may replace current health care-associated MRSA (HA-MRSA) clones in the future. This hypothesis is supported not only by mathematical models but also by reports that have shown invasion of CA-MRSA clones to hospitals (Moellering, 2010). First described in Minnesota, CA-MRSA has now attracted global attention (DeLeo et al., 2010). MRSA related to livestock infections has also been reported (Bortolami et al., 2017). However, this type of MRSA seems to be limited to some countries, especially the ones where pig farms are common (Angen et al., 2017). Some European countries have maintained low rates of MRSA (Köck et al., 2010; Stefani et al., 2012). However, an increase in the worldwide prevalence of MRSA highlights the urgent need of finding alternative solutions to this emerging problem (Graveland et al., 2011; Stefani et al., 2012). Keeping in view all these facts and findings, the present study aimed to determine the antibacterial activity of aqueous extract obtained from two varieties of desert truffles against MRSA bacteria.

## 2 Materials and Methods

### 2.1 Preparation of the Desert Truffles Extracts

#### 2.1.1 Desert Truffles Collection

Two types of desert truffles *T. clavaryi* (Khallasi) and *T. nivea* (Zubaidi) (1Kg each) were collected from guard fence, Al-Ahasa Street, Riyadh (East), Saudi Arabia.

#### 2.1.2 Truffle Crude Extracts

The crude truffles extract was prepared according to method described by Janaket et al. (2004). Fifty grams of truffles were cut into small pieces and soaked in distilled water (1:3) for 24 h. The mixture was homogenized for 1 min at full speed and homogenate was filtered through double layer of sterile cheese cloth (sterile gauze was used instead of it) and centrifuged at  $3000 \times g$  for 10 min at  $4^{\circ}C$  in order to remove its soluble material. The

supernatant constitute the crude aqueous extract was recovered and filtered under sterile condition through a  $0.22\mu m$  filter. The obtained sterile filtrate was than kept at  $-15^{\circ}C$  until use.

### 2.2 Subjects

A retrospective observational chart review was conducted of surveillance MRSA nasal-swab screening and corresponding culture results for clinical isolates. MRSA screening nasal specimens were collected using sterile cotton swabs (ACI Saudi Plast labs, Jeddah, Saudi Arabia) as per laboratory standard operating procedures. Nasal samples were collected from Fifty Health Care Workers (HCWs) and Community Volunteers (CV) by ENT specialist from King Abdulaziz University Hospital (KAUH), Jeddah, Saudi Arabia. Eligible subjects for that trial also included those patients admitted to the Hospital with a history of MRSA infection or colonization  $\geq 90$  days from the day of admission. The subjects were approached and attempts were made to screen them on days 1, 2, and 3 of hospitalization. The swabs were simultaneously inoculated and assessed on Mannitol Salt agar for the presence of *S. aureus* colonies and further subculture on Muller-Hinton agar for the isolation and identification of MRSA.

A total of 50 MRSA isolates from nasal specimens were randomly selected for further analysis. These isolates were also tested for routine antibiotics by Kirby-Bauer disk diffusion method on Mueller-Hinton agar as per Clinical and Laboratory Standards Institute (CLSI).

The study was approved by Research Ethics Committee of Faculty of Medicine, King Abdulaziz University, Jeddah, Saudi Arabia. All volunteers hospital staff signed informed consent and the questioners before being included in this study because of the retrospective nature of the study.

### 2.3 Antibiotics Susceptibility Testing (AST)

#### 2.3.1 Disc Diffusion Test (DD test)

Antibiotic susceptibility tests of all MRSA isolates were performed using a disk diffusion method on Muller Hinton agar with antibiotic discs of Oxacillin (1  $\mu g$ ), Ampicillin (10  $\mu g$ ), Ciprofloxacin (5  $\mu g$ ) and Azithromycin (15  $\mu g$ ), then interpreted according to the Clinical & Laboratory Standards Institute (CLSI, 2007) guidelines. Each bacterial isolate were initially considered MRSA, if the test results of inhibition zones for Oxacillin was  $\leq 10$  mm.

### 2.3.2 Agar Well Diffusion Method

Bacteria (1 ml) were spread on the surface of Muller Hinton agar evenly and the excess was withdrawn by micro-pipette. Wells of six mm diameter were punched into the agar and filled with 100  $\mu$ l of the aqueous extracts of truffles *T. clavaryi* (Khallasi) and *T. nivea* (Zubaidi). Plates were kept first at 4°C for at least 2 hrs to allow the diffusion of any antibacterial metabolites and then there were incubated for 24 hrs at 37°C. The antibacterial effects of aqueous extracts of the two varieties of truffles were compared with those of Oxacillin, Ampicillin, Ciprofloxacin and Azithromycin antibiotics which usually employed against pathogenic strains. All experiments were carried out in Triplicate. The antimicrobial activity was determined by observing the zone of inhibition (ZOI).

### 2.4 DNA Extraction

DNA was extracted using Gene JET Genomic DNA Purification Kit #K0722 (Thermo Scientific, USA) according to the manufacturer's protocol.

### 2.5 Agarose Gel Preparation

Gels with a concentration of 2% (w/v) agarose were used for gel electrophoresis. Each gel consisted 2 g agarose and 100ml 1X TAE buffer. The solution was melted in a microwave (about 2-3 min) until the agarose was completely dissolved and the solution appeared clear. After melting the solution was cooled down to approximately 65°C and 2 $\mu$ l of ethidium bromide solution were added, mixed well and poured into a prepared gel casting tray for cooling, until the gel looked milky and became solid.

### 2.6 Polymerase chain reaction (PCR)

DNA (8  $\mu$ l) was amplified in 25  $\mu$ l reaction by using the master mix from Promega (GoTaq® Green Master Mix (2X), M7122, USA). The primers were designed based on the highly conserved of *mecA* gene from Eurofins Genomics (Germany).

The *mecA* gene was amplified by PCR using previously reported primers (Table 1). DNA amplification was carried out in thermal cycler (multigene™ optimax thermal cycler, model no. 66-TC9610) supplied from Lab net International, USA. The reaction cycles for each gene included, initial denaturation at 94°C for 4 min followed by 40 cycles of amplification (denaturation at 94°C for 30 secs, annealing at 55°C for 30 secs, and extension at 72°C

Table 1 Primer Sequence of *mecA* gene

Gene name	Primer sequence
mecA	FP : 5'ATAGAGATGCTGGTACAGG3'
	RP : 5'GCTTCCGATTGTCGATGC3'

for 1min) and a final extension at 72°C for 5 min. PCR products were run through a 2% agarose gel (in 1X TAE buffer) containing Ethidium bromide (2  $\mu$ l/100ml agarose solution) at 80 V for 45 min. In parallel, a 1Kb DNA ladder (CSL-MDNA-1Kb, Cleaver Scientific Ltd, UK) ranging from 250-10 Kb was applied to determine the size of the DNA. DNA fragments were visualized through UV light.

### 2.7 Statistical Analysis

The statistics were performed by using Windows Microsoft Excel 2013 software, using t-test to detect the significance at p- value (p<0.05).

## 3 Results and Discussion

In this study, the antibacterial properties of both truffles *T. clavaryi* and *T. nivea* aqueous extracts on the growth of MRSA were evaluated and promising results were obtained. In agar well diffusion method, *T. clavaryi* was reported more effective against MRSA (27 $\pm$ 7.75mm zone of inhibition) as compared to *T. nivea* (22.5 $\pm$ 11.26 mm zone of inhibition) (Table 2; Figure 1). These findings are in agreement with the study of Gouzi et al. (2011) and Casarica et al. (2016). Similarly, Neggaz & Fortas (2013) suggested that ethyl acetate extract of *Tirmania pinoyi*

Table 2 Comparison ZOI of antimicrobial activity between *T.clavaryi* and *T.nivea* aqueous extracts and antibiotics discs.

Test bacteria	<i>T.clavaryi</i> 100 $\mu$ l	<i>T.nivea</i> 100 $\mu$ l	OX 1 $\mu$ g	AMP 10 $\mu$ g	AZM 15 $\mu$ g	CIP 5 $\mu$ g
MRSA	27 $\pm$ 7.75	22.5 $\pm$ 11.26	9 $\pm$ 6.65	11 $\pm$ 6.35	21 $\pm$ 9.07	23 $\pm$ 2.5

Diameter of inhibition zones were measured in mm  $\pm$  SEM; Oxacillin (OX), Ampicillin (AMP), Azithromycin (AZM), Ciprofloxacin (CIP)



Figure 1 Desert Truffle extracts effect against MRSA, Z refers to Zubidi (*Terfiza nivea*) K refers to ikhallasi (*Terfiza calvayn*).

have significant inhibitory activity against Gram negative and Gram positive bacteria and have antifungal activity against *Candida albicans*. Results of these studies conclude that desert truffles had antibacterial and antifungal activity against many bacteria and fungi species that support the traditional use of truffles. The results obtained from antibiotic susceptibility/resistant test showed that MRSA exhibited high resistant against Ampicillin and Oxacillin and sensitive for Azithromycin and Ciprofloxacin, as compared to the desert truffles extract values. It was clearly reported that the aqueous extracts of both truffles possessed higher antibacterial activity against MRSA than the 1µg Oxacillin, 10 µg Ampicillin, 15 µg Azithromycin and 5 µg Ciprofloxacin (Table 2). These results are in agreement with the findings of Gouzi et al. (2011).

At confidence level 95%, the p value = 0.2765 considered non significant association between the truffles extracts and MRSA sensitive antibiotics (Ciprofloxacin and Azithromycin) but there were significance association between the truffles extracts and MRSA resistant antibiotics (Oxacillin and Ampicillin) with p value = 0.0353 (Table 3 & 4).

The *in vitro* antibacterial activity of 3 edible desert truffle mushrooms species aqueous extracts (*Tirmania pinoyi*, *Terfezia claveryi* and *Picoa juniper*) against Gram-positive human pathogenic reference strain *Staphylococcus aureus* ATCC 29213 and the Gram-negative strain *Pseudomonas aeruginosa* ATCC 15442 are recently reported by Schillaci et al. (2017). Aqueous extract of the truffle *Terfezia claveryi* contains a potent antimicrobial agent that is protein in nature and may be used in the treatment of eye infections caused by *P. aeruginosa*. Relative antimicrobial activities of these truffles fractions were found to be superior to most of reference antibiotics used (Janakat et al., 2005).

Disc diffusion test (DD test) was used to detect antibiotic sensitivity of MRSA isolates. Results of study revealed that out of 50 *S. aureus* isolates, 23 (46%) isolates were found resistant to Ampicillin and Oxacillin and sensitive for Azithromycin and Ciprofloxacin (Table 3 & 4). Staphylococcal resistance to Oxacillin occurs when the organism including an altered Penicillin-Binding Protein (PBP2A) that is coded by the *mecA* gene (Fishovitz et al., 2014). Oxacillin showed stability under storage conditions. Oxacillin resistant isolates were initially interpreted as MRSA based on DD test which were not proved after further investigations. It was also observed that phenotypic methods to detect methicillin resistance in *S. aureus* (MRSA) are inadequate and need more developments. These results also supported by Alipour et al. (2014) who described that DD assay from a single specimen is not optimal for the rapid detection of MRSA.

Table 3 MRSA isolates occurrence percentage to the antibiotics via susceptible / resistant test (DD test).

Antibiotics	Groups				Susceptible (Sensitive)	<i>S.aureus</i> (Resistant)		MRSA (Resistant)	
	SM	SF	VM	VF		No.	%	No.	%
Oxacillin	1	8	6	8	27	23	46%	-	-
Ampicillin	4	13	11	20	2	48	96%	22	(45.83%)
Azithromycin	2	3	2	6	37	13	26%	12	(92.31%)
Ciprofloxacin	0	4	1	0	45	5	10%	3	(60%)

Here SM. Staff male, SF. Staff female, VM.Volunteer male, VF. Volunteer female

Table 4 Relationship of biophysical parameters and the prevalence of MRSA

Parameter	Df	t-value	SD	P-value
Desert truffles extract	MRSA S.AB	1	1.41	0.2765**
	MRSA R.AB	1	6.36	0.0353
Age	3	2.45	19.37	0.0018
Gender	1	4.3	14.14	0.2597**
Group	3	2.45	6.56	0.0670**

\*\* No significant difference, S; Sensitive, R; Resistant AB; Antibiotics

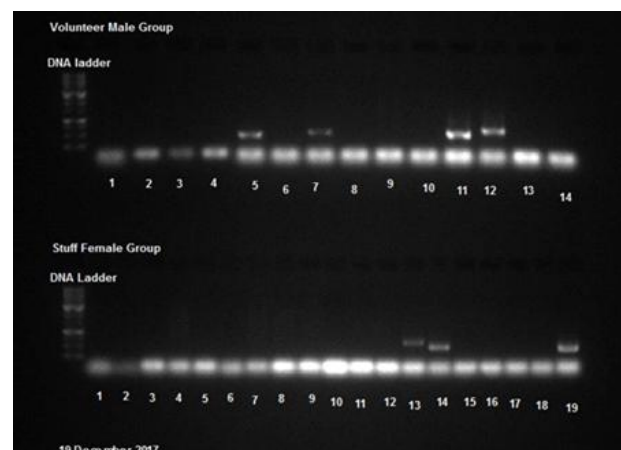


Figure 2: 2% agarose gel electrophoresis of *mecA* gene PCR products where lane 1:1 Kb DNA ladder and lane 1-14(VM), 1-19(SF); PCR product.

PCR assay results showed that among randomly selected 50 MRSA isolates, 17 MRSA isolates showed the presence of two types of plasmid bands at 500 bp and 750 bp after gel electrophoresis. The 500bp band was the most common (76.47%) while 750 bp band observed only in 23.52% gel electrophoresis samples which indicating the presence of *mecA* gene. It also showed that 17 (34%) MRSA strains isolated from nasal carriage found positive as MRSA in PCR analysis (Figure 2-4). Alipour et al. (2014) observed 37% of positive isolates by PCR assays and also recorded the presence of

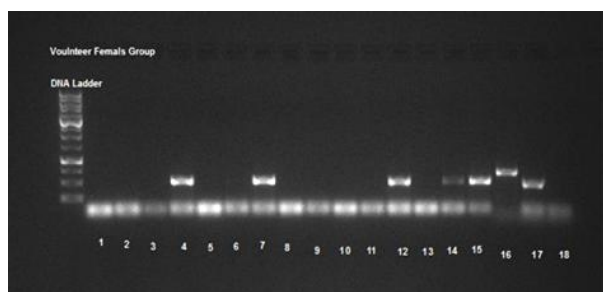


Figure 3: 2% gel electrophoresis of *mecA* gene PCR products where lane 1:1 Kb DNA ladder and lane 1-18(VF): PCR product



Figure 4: 2% gel electrophoresis of *mecA* gene PCR products where lane 1:1 Kb DNA ladder, a) lane 19-24 (VF) and b) lane 1-4(SM): PCR product.

*mecA*. Further results of study showing that 8% hospital staff comprised of 1 (2%) male (SM) and 3 (6%) female (SF) have MRSA occurrence. In volunteer female (VF), 9 out of 19 samples (18%) were positive; while in volunteer male (VM) 4 out of 11 were found positive (Table 4). These findings are supported by the study of Jaffe et al. (2000). MRSA significance was tested for age, gender and groups. It showed that there was significant difference between MRSA and age ( $p$  value = 0.0018). Among these 20-29 age group respondents had 10 positive samples while 30-39 age group had 5 positive samples and 40-49 have 2 positive samples. The negative MRSA samples were observed at age range 50-59 (Table 3). The highest rate of MRSA was reported for the age group of 20-29 years, among the youngest participants. Positive throat carriage MRSA, who even not exposed to health care system, was more common in 30 years and younger participants (Mertz et al., 2009). The gender and MRSA positive results showed no significant association with ( $p$  value = 0.2597) even though 24% of female and 10% of Male had MRSA carriage. These findings were in contrary with the findings of Kupfer et al. (2010) who confirmed that male gender were at higher risk of MRSA carriage. Further, no significant association was reported between Hospital Staff Male (SM), Female (SF) and community Volunteer Male (VM) and Female (VF) groups ( $p$  value = 0.0670).

This result was also in contrast with the findings of Cesur & Çokça, (2004) study who reported that hospital staff are more likely to be MRSA carrier ( $p$  value = 0.013) than the outpatient but it was in agreement with the study of Alsulami et al. (2017) who reported non significant difference between groups of units.

Our findings showed the Community-Associated MRSA had highest prevalence rate than the hospital-associated MRSA. Previous studies also reported that increasing in the occurrence of community-acquired MRSA lineages plus emergence of pandemic and rare MRSA strains is occurring now in these days (Senok et al., 2016). Due to this reason, the Methicillin-Resistant *S. aureus* USA300 strain has become the dominant strain in hospitals as well as in the community because of its unique characteristics. Type IV SCCmec cassette in USA300 is smaller than types I-III in hospital-acquired MRSA (HA-MRSA). Community acquired CA-MRSA including USA300 carry fewer antibiotic resistance genes than HA-MRSA. Doubling time of USA300 is shorter by ~1.25-fold. Linkage of arginine catabolic element with SCCmec IV in USA 300 confers increased fitness and pathogenicity of this strain. These characteristics made this strain as a type causes most community-associated MRSA infections and is an increasingly common cause of health care-associated MRSA infections (Moellering, 2010). In this study, after combining the observations about MRSA prevalence and its solution from AST/DD test and PCR assays have accurately revealed that MRSA rate had raised among community associated MRSA (CA-MRSA). Analyzing genotyping studies enabled us to track the emergence of a new, successful MRSA type in space and time across the countries. The prevalence of desert truffles species with multiple antibacterial traits emphasizes its potential for use and development of effective antimicrobial drug that could be used to improve the human health against MRSA. Further studies must be carried out in large and identical group numbers to have more precise outcomes.

## Conclusion

This study has shown that the aqueous extract of *T. nivea* and *T. clavaryi* desert truffles exhibited antibacterial activity against MRSA. The antibacterial potential of *T. clavaryi* was stronger and effective than the *T. nivea* extract. Molecular detection by PCR assays had identified the frequent prevalence of MRSA among community associated MRSA (CA-MRSA). This investigation supports the traditional use of truffles for the treatment of eye infection and proved its effectiveness against MRSA which could be considered as promising antibiotic drug of future use.

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### Conflict of Interest

Authors would hereby like to declare that there is no conflict of interests that could possibly arise.

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