Antibacterial activity of multidrug resistance *Escherichia coli* by some natural products

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ABSTRACT

Multidrug resistance *Escherichia coli* have one of most common resistance in the world. A total of 50 blood samples from Mansoura University Hospitals were collected, isolated, and identified E. coli, by molecular and chemical characterization using the Vitek AST automated technique. The results demonstrated antibiotic sensitivity results showed resistance against ciprofloxacin and moxifloxacin 100%. Ampicillin 100% Ampicillin/Sulbactam (91.93%), Aztreonam (96.93), Cefazolin (72.45%), Ceftazidime (100%), Ceftriaxone (94.68%), Cefepime (96%), Ciprofloxacin (100%).Gentamicn (73%),Moxifloxacin (100%),Nitrofurantoin 83.33%. Piperacillin/Tazobactam (100%), Tobramycin (97.53%), Trimethoprim/Sulfamethoxazoleol (70%), Ticarcillin (100%), Ticarcillin/Cavulanic acid (100%). Laboratory cultures of some fluoroquinolone groups, such as ciprofloxacin, levofloxacin, ofloxacin and norfloxacin, were also found to be resistant to E. coli bacteria. Two strains of Escherichia coli were used, and the antibacterial activity of varying concentrations (25, 50, 75, and 100) of aqueous garlic extract and peppermint was tested against the two strains. The findings revealed that although mint had no effect at all, garlic had a complete effect in all concentrations.

Keywords: MDR, Mint, Resistant E. Coli, Anti-bacterial, Garlic aqueous extract.

1. INTRODUCTION

Over the past decade, the intensive use of quinolones in human and veterinary medicines as well as in food animals has led to the rapid emergence of quinolones resistance 2016). (Haque al., et Fluoroquinolones (FQ) are effective antibacterial medications used to treat a wide range of nosocomial and community diseases. FO resistance rates vary substantially by sex, age, type of urinary infection, and geographic region, although it has recently been revealed that *E. coli* isolated from community-acquired UTIs are becoming more resistant to FQ, with up to 29% of women carrying this strain (Hossain *et al.*, 2016). Since the 1980s, when fluoroquinolones were first used in clinical treatments, this class of antibiotics has been frequently chosen to treat infections caused by *Escherichia coli*, *Salmonella*, and other Enterobacteriaceae in both humans and animals. This is due to the fluoroquinolones'

oral administration benefit and bactericidal activity against Gram-negative bacteria (Kim et al., 2009). The rate of quinolone resistance differs widely from one country to another. According to the Center for Disease Dynamics, Economics and Policy the percentage of fluoroquinolone resistant E. coli among invasive isolates was 29% in United States of America, ranged from 7% to 46% European countries. and in reached>70% in some countries as India (78%) and Pakistan (72%) (Jeong et al., 2005). One of the most prevalent causes of various typical bacterial infections in both humans and animals continues to be Escherichia coli. Enteritis, uTIs, septicemia, and other clinical infections such infant meningitis are frequently brought on by E. coli (Briales et al., 2012). Chromosome mutations that change DNA gyrase, the medicines' biological target, or decrease intracellular drug concentration are two examples of fluoroquinolone resistance mechanisms in bacteria. Both chromosomal and plasmid-mediated mechanisms are involved in bacterial resistance to quinolones and fluoroquinolones (van der Zee et al., 2016). Escherichia coli are one of the microorganisms most frequently studied worldwide. they are Gram-negative bacilli, facultative anaerobic, rod-shaped bacterium, which are most often found in the gastrointestinal tract as a normal colonizer of warm blood organisms (mainly in mammals, but are also present in birds, reptiles and fish) (Ehwarieme et al., 2021). Escherichia coli can emerge from primary infection sites like percutaneous vascular systems asymptomatically. Bacteremia can develop into sepsis when bacteria in the blood result in a significant systemic response that leads to sepsis syndrome, severe sepsis (sepsisinduced malfunction of at least one organ or or septic shock. Septicemia system). accounted for 1.3% of all fatalities in the United States in 2000, making it the tenth

most common cause of death (Aslam *et al.*, 2020). Many societies have employed garlic (Allium sativum L.) to combat infectious diseases because it is effective against grampositive bacteria. (Adler *et al.*, 2002). It is advised to use garlic to cure bacteria that are resistant to many drugs. (Dini *et al.*, 2011). **The goals of this research** are to evaluate the antibiotic resistance phenotype of *E. coli* that tested the effect of garlic and mint extract against multidrug-resistant *E. coli*.

2. MATERIALS AND METHODS 2.1. Population: Collection of samples

About 50 blood samples from patients (Approval No. H18RE34) who received care while hospitalised and whose data was thoroughly evaluated and subsequently studied were used for the testing. Samples were taken between October 2019 and February 2021 from Mansoura University Hospitals in Egypt. Blood was used to gather the samples. Samples were taken before the administration of antibiotics. Under sterile conditions, blood samples (5–10 ml) were drawn and inoculated into a blood culture container.

2.2. Automated Broth Dilution AST Systems

Vitek AST Automated System VITEK® 2 COMPACT (bioMérieux) Marcy l'Etoile, France Vitek 2 measures the turbidity changes over time (growth curve) and compare a growth control well with wells of drug concentrations. other MicroScan WalkAway, Vitek, and Vitek 2) is a broth microdilution method that utilizes a standard 96 microwell panel. These microwells contain serial dilutions of dehydrated antimicrobial agents. Results are obtained after about 18 hours by turbidimetric readings of overnight conventional panels or after about 7 hours by fluorometric readings

of rapid panels. Fluorometric analysis depends on the degradation of fluorogenic substrates by viable bacteria. The phoenix system detects the susceptibility results by gravity-based inoculation process. Monitoring of the growth depends on a redox indicator system which provide results in 8 to 12 hours.

2.3. Bacterial DNA extraction

Rapid DNA extraction was performed using methods of (**Salah** *et al.*, **2019**). Where 10 samples were isolated as in Figure (2).

2.4. Sequencing of the 16SrRNA gene of the isolates

The thermal cycling conditions were 96 °C for 10 seconds, 50 °C for 5 seconds, and 60 °C for 4 seconds for the sequencing reactions in the 9700 thermal cycler with a volume of 20 L. (25 cycles). Afterwards, the cycle sequencing reaction was isolated from the extra dye terminators and primers using the DyeExTM 2.0 Spin Kit (Qiagen PN 63204). The resulting sequences were examined using the Finch TV (version 1.4.0) programme, and the phylogenetic tree was constructed using the closest published type strain sequences created using the Seaview programme.

2.5. Prepare aqueous garlic and mint extract

Fresh garlic obtained from the local market were used to make aqueous garlic extract. Garlic weighing 100 grams have been cleaned. Then, a 60-sec immersion in 70% (volume/volume) ethanol was performed as a method to sterilize the surface (Kalyan, 2000). Garlic was homogenized using a sterile mortar and pestle after the ethanol had been vaporized in a sterile laminar flow chamber. The extract was filtered with 0.45-µm membrane, after centrifugation at 6000 rpm for 10 m. This extract contains 100% concentration of the original substance. Finally, 25, 50, 75 and 100% of the concentrated garlic extract was produced by diluting 100% of it with sterile distilled water (Durairaj *et al.*, 2009). The mint from the local market was extracted in the same way as before.

2.6. Evaluation of the antibacterial activity of garlic and mint extract

Using a well-agar diffusion method two tested bacterial strains of *E. coli* were cultured on MacKonky Agar, then incubated at 37 °C for 8 h. The bacterial culture was spread on the surface of the nutrient agar plate, and a cork borer was used to puncture wells with a diameter of 6 mm. 100 μ L of the following concentrations were added into each well: standard ciprofloxacin 5 mg/mL (a reference antibiotic) in sterile distilled water 25, 50, 75, and sterile 100% garlic and mint extract.

2.7.MIC to garli

Two strains of *E. coli* bacteria were tested for their response to various garlic doses. 100, 75, 50, 25, 12.5, 6.25 and 3.13.

3. RESULTS AND DISCUSSION

3.1. Identification of *Escherichia coli* by Vitek

In the current investigation, 50 blood samples in total were gathered. Results of a Vitek sample test are displayed. All samples (n = 50) contained *Escherichia coli*. Furthermore, all samples (n = 50) had a probability of 93% to 99% for *Escherichia coli* identification by Vitek.

3.2. Antibiotic susceptibility profile:

According to the results shown in Table (1), the isolated E. Coli is resistant to the following drugs: ciprofloxacin (100%), gentamicin (73%), moxifloxacin (100%), tobramycin (97.53%), piperacillin/tazobactam (100%),trimethoprim/sulfame.70%, ticarcillin (100%), and ticarcillin/cavulanic acid (100%). It is also resistant to ampicillin (10 The resistance ratios for amikacin, imipenem, meropenem, ertapenem. nitrofurantoin, and tigecycline are (10.52, 14.45, 15.3, 17, 17, 27.05, and 1.07, respectively), despite E. coli being less resistant to these drugs. A full allergy to colistin in the interim. The findings in Table

(2) show that every isolate is a multidrugresistant bacterium. 72% of the 50 blood samples had ESBL-*E. coli* in them.

3.3. Sensitivity test

Sensitivity test was performed in the lab on a few fluoroquinolone drugs and the resistance of bacteria to the antibiotic class was seen as in Figure (1).

Table 1. Bacterial identification	of the	two isolates.

Isolates	Identification	Identified Accession number	Highest homology	Indenty %
1	<i>Escherichia coli</i> Strain A1	MW719074	<i>Escherichia coli</i> strain GenBank: MT448673.1	100
2	<i>Escherichia coli</i> Strain A2	MW719253	<i>Escherichia coli</i> strain GenBank: MT735392.1	99.9

Table 2. Antibacterial activity of Garlic extracts against E. coli strains.

TREATMENTS	CONC.	ZONE OF INHIBITION MM(MEANS ± S.EROR.)	
		E. coli MW719074	E. coli MW719253
CIPROFLOXACIN	5mg	0	0
	25%	10.3 ± 0.1552	9.1 ± 0.2445
GARLIC EXTRACTS	50%	15 ± 0.8221	14 ± 0.365
	75%	17 ± 0.4226	16 ± 0.287
	100%	20 ± 0.3225	19 ± 0.726



Fig 1. Testing of a few fluoroquinolone drugs' sensitivities in lab.

3.4. DNA extraction

In this study, 10 samples were isolated and DNA was extracted as in Figure (2).

3.5. Molecular identification of the selected isolates.

The selected isolates can be identified thanks to the study of the 16S rRNA gene sequence (Figure, 3). Two samples were isolated and there results were in Figure (4).

3.6. Antimicrobial activity of hydrolyzed garlic and mint

The results in Tables (2 and 3) and Figures (5 and 6) demonstrated the potential

antibacterial activity of 25, 50, 75, and 100% of garlic aqueous extract against *Escherichia coli* MW719074 and MW719253, as well as ciprofloxacin 5 mg (used as a reference). According to the results. While mint has no effect at all, garlic extracts do have an antibacterial effect against two types of bacteria, and the activity increases as the concentration of aqueous garlic extracts increases.

The results demonstrate in figure 7 represent 25% garlic extract was minimam inhibitory concentration (MIC) of two *E. coli*

TREATMENTS	CONC.	ZONE OF INHIBITION MM(MEANS ± S.EROR.)		
		E. coli MW719074	E. coli MW719253	
CIPROFLOXACIN	5mg	0	0	
	25%	0	0	
	50%	0	0	
MINT EXTRACTS	75%	0	0	
	100%	0	0	

Table 3. Antibacterial activity of Mint against E. coli strain

TREATMENTS	CONC.	ZONE OF INHIBITION MM(MEANS ± S.EROR.)	
		E. coli MW719074	E. coli MW719253
CIPROFLOXACIN	5mg	0	0
	3.13%	0	0
GARLIC EXTRACTS	6.25%	0	0
	12.5%	0	0
	25%	10.3 ± 0.1552	9.1 ± 0.2445
	50%	15 ± 0.8221	14 ± 0.365
	75%	17 ± 0.4226	16 ± 0.287
	100%	20 ± 0.3225	19 ± 0.726

Table 4. Antibacterial activity of different concentrations of garlic against *E. coli* strain(MIC)

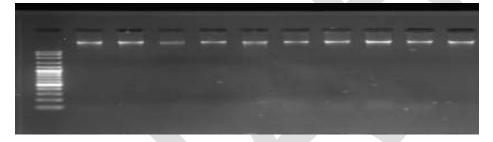


Fig 2. 10 DNA extraction samples were isolated (1,2,3,4,5,6,7,8,9,10)

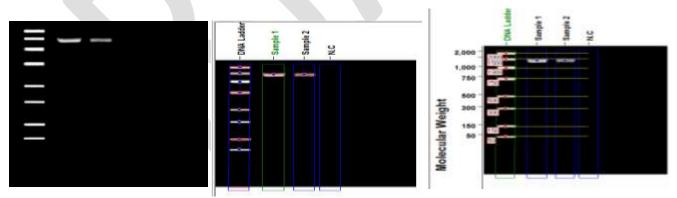


Fig 3. Specific genomic products for 16SrRNA bacterial strains with \approx 1500bp. and Computerized and molecular weight detection for Specific genomic products for 16SrRNA with \approx 1500 bp.

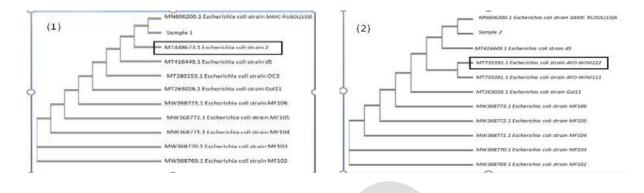


Fig 4. Phyllogenetic tree for the two isolates.

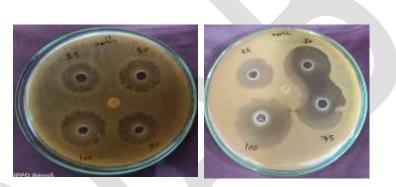


Fig 5. Activity of garlic against two strains of Escherichia coli.

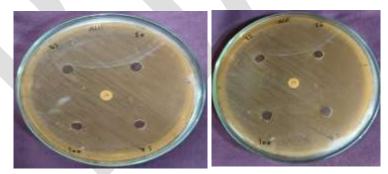


Fig 6. Activity of mint against two strains of *Escherichia coli*.

3.7. Discussion

All isolated samples identified MDR E. *coli*. Antibiotics are all over the place, which could explain the MDR. The causes that are most likely to contribute to the spread of antibiotic resistance include globalization, excessive antibiotic use in aquaculture and animal husbandry, the use of numerous broad-spectrum medications, and a lack of effective antimicrobial management (Vivas et al., 2019). The results showed that E. coli are resistant to a number of antibiotics, including ampicillin, ampicillin/sulbactam, aztreonam, cefazolin, ceftazidime, ceftriaxone. cefepime, ciprofloxacin, gentamicin, moxifloxacin, nitrofurantoin, piperacillin/tazobactam, tobramycin, trimethoprim (Aabed et al., 2021). E. coli isolated from healthy children in China resistance to has extreme Ampicillin (54.0%), Arimethoprim/Sulphurmethoxazole (47.5%) and Tetracycline (58.9%) (Zhao et al., 2021). This research results showed that E. coli bacteria are resistant to a number of antibiotics. including ampicillin, ampicillin/sulbactam, aztreonam, cefazolin, ceftazidime. ceftriaxone. cefepime. gentamicin, moxifloxacin, ciprofloxacin, nitrofurantoin, beprazobactamide. 100% of the evaluated ESBL-E. coli bacteria were ampicillin resistant. (Abalkhail et al., 2022). In this research 100% of the ESBL-E rating. *coli* resistant to ampicillin I agree with this. Major antibiotic-resistant E. coli strains are becoming more prevalent in almost all of Europe. (Chang et al., 2015). This research agreement with (Abalkhail et al., 2022), all E. coli isolates are MDR and that 78% of E. *coli* samples are ESBL E. coli. 15 antimicrobial medications. including Ceftriaxone, Cefotaxime, Ceftazidime, Cefpime, Cefuroxime. Cephalothin, Cefoxitin, Amoxicillin-Ampicillin, Trimethoprimclavulanate. Aztreonam, sulfamethoxazole. Gentamicin. and

Tigecycline, showed high resistance in ESBL-E. coli. E. coli with ESBLs present in Iran, Pakistan, and 40% (Hashemizadeh et al., 2018; Ali et al., 2016). This study proved aqueous garlic extract has antibacterial properties in different doses against E. coli MW719074 and MW719253. Garlic and ginger are natural herbs that have powerful antibacterial properties against pathogens that are resistant to several drugs and can be utilised to prevent the development of drugresistant microbial illnesses (Karuppiah and Rajaram, et al., 2012). It was agreed that garlic affected all its concentrations, and the bacteria were affected by mint extract, and the results showed that mint did not affect all its concentrations at all. Many researches on suggested that allicin in garlic is the most vital compounds that makes antibacterial properties and restrictions the speed of RNA synthesis; the preliminary task of allicin is tricking RNA (Deresse, 2010). Allicin containing garlic dominate extracts inhibitory and bactericidal activities against Burkholderia cepacia (Wallockthe Richards et al., 2014). Ethyl acetate, Ethanol, Chloroform, Hexane, Petroleum ether and Methanol. extracts of the Garlic leaves and bulbs were possessing highest antibacterial activity against pathogenic bacteria (Prasad et al., 2018). Ultrasonicated garlic extract (garlic nanoparticles) displayed highest effective against Escherichia coli, Streptococcus mutans, Staphylococcus aureus, and Poryphyromonas gingivalis (Gabriel et al., 2022).Garlic aqueous extract was to damage bacteria and inhibit from working of the protein generating RNA synthesis (Mozaffari et al., 2014). The high antimicrobial activities of many bacterial types by garlic are related to the garlic contents such as allicin, ajoenes, and allyl sulfides (Bhatwalkar et al., 2021).

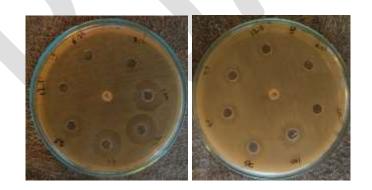
The possibility of interbreeding different species of the genus Mentha results in the fact that their taxonomy is complicated, and

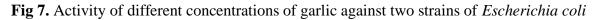
chemical composition thus their is heterogeneous. Therefore, the chemical composition and biological properties of individual hybrids can significantly vary from one another, depending on many factors, both environmental and genetic. The heterogeneous botanical systematics of the constantly emerging new mint cultivars problems with creates their proper identification, and thus with the appropriate

4. Conclusion

This investigation demonstrated that all *E. coli* isolates were resistant to many drugs, including ampicillin, ampicillin/sulbactam, azitreonam, cefazolin, ceftazidime, moxifloxacin, gentamicin, moxifloxacin, nitrofurantoin, piperacillin/tazobactam, tobramycin, trimethoprim/sulfamethoxazole, ticarcillin, ciprofloxacin. Aqueous garlic extract has antibacterial activity o against *E. coli* MW719074, and *E. coli* MW719253. It determination of their chemical profile and possible use for therapeutic purposes. The active constituents of *Mentha* species leaves include flavonoids (eriocitrin, hesperidin, diosmin, luteolin and their glycosides), phenolic acids (derivatives of caffeic acid, e.g. rosmarinic acid), terpenoids, and volatile compounds (Fecka and Turek, 2007; Mahendran and Rahman, 2020; Nilo et al., 2017).

must be Egyptian food and drug authority (EFDA) launch an electronic awareness campaign, which aimed to raise the awareness in community towards the necessity to avoid taking the antibiotic without a prescription, and warn of complications of excessive use of antibiotics. In addition, stressing the need to adhere to prescribed doses of antibiotics. While mint did not affect this strain of bacteria.





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