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Extracts of Turmeric (*Curcuma longa* L.) as a Potential Biocontrol of Multidrug-Resistant *Acinetobacter baumannii*



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Article Information

Received 25 May 2022, Revised 6 June 2022, Accepted 10 June 2022. Published online 30 June 2021 **Abstract:** Nosocomial infections with *Acinetobacter baumannii* are increasing every day, making treatment with traditional antibiotics more complicated. This work aimed to investigate the antibacterial activity of Curcuma longa extracts against carbapenemresistant isolates of A. baumannii. A total of 90 samples (Sputum, n=40 and Urine, n=50) were collected and biochemically identified via Vitek -2 system. Ten isolates were identified as A. baumannii (10/90; 11.1%). (10/90; 100 %) of isolates identified were multi drug resistance (MDR). Curcuma longa extracts aqueous and/or ethyl acetate of, exhibited a significant MBC value varying from 43.75 to 50.83 mg ml⁻¹ against A. baumannii isolates (A1, A2, A3, A5, A6, and A8), respectively. Within isolates A1 and A8, ethanol extracts revealed MBC with significant values (30 to 70 mg ml⁻¹) respectively. The main phytochemical compounds detected among 68 bioactive compounds in the extracts of ^C. longa over the GC- mass spectra analysis were Turmerone (59.42%), Beta-caryophyllene (16.66 %), Caryophyllene oxide (15.62 %), Isolongifol (12.14 %), Alpha-pinene (12.07 %), Citral (11.68 %), Curdione (11.0 %), B -Sesquiphellandrene (10.47 %), B-myrcene (8.12 %), D-Limonene (8.04 %). Bactericidal activity of C. longa against MDR A. baumannii were supported by phytochemical compounds that detected in GC-MS analysis.

Keywords: Acinetobacter baumannii, Curcuma longa, GC-Mass, Imipenem.

Introduction

Nosocomial infections with A.baumannii are a common cause of multiple respiratory, wound and urinary tract infections, and have become one of the most predominant illnesses and representing more than 54% of hospitalized patients (Ayoub & Hammoudi, 2020). Infection with A.baumannii is often difficult to treat because of resistance to multiple antibiotics and disinfectants (Williams et al., 2020). They represents second most common gram-negative bacteria after Pseudomonas aeruginosa, which identified in clinical specimens (He et al., 2011). In Egypt A. baumannii recovered from different location of hospitalized patients (Abdulzahra et al., 2018). A. baumannii induced respiratory pneumonia is widespread in clinical environments as their surviving as a commensal on the skin or hair of hospital staff and patients (Vrancianu et al., 2021). Urinary tract infections with A. baumannii were increasingly found in immune compromised patients, leading to a high significant morbidity and mortality (Ayenew et al., 2021). A. baumannii is generally antibiotic-resistant because of decreased

permeability, efflux pumping systems, inactivation of enzymes, and the formation of biofilm (Chakravarty *et al.*, 2020). Thus, they are often β -lactams, aminoglycosides, and quinolones resistant (Motbainor *et al.*, 2020).

β-lactams antibiotics such as cephalosporins, carbapenems, and penicillins represent approximately 60% of the used antibiotics (Magiorakos et al., 2012). A. baumannii resistance is mainly due to Extended-Spectrum Beta-Lactamases (ESBLs), which could disrupt various β-lactam antimicrobial agents as penicillins and their derivatives (Ghaima, 2018). C. longa commonly known as (Turmeric), is widely used as a spice and colouring agent, and well known for its antibacterial properties (Kumar & Sakhya, 2013). There is also urgently requirement to identify novel pharmacological agent and to understand the function of possible therapies in the treatment of A. baumannii infection. In the present study, the prevalence, patient clinical characteristics, antibiogram, of A. baumannii isolated from different clinical samples were investigated. Subsequently, phytochemical investigations of C. longa different extracts were carried chromatography-mass out using gas

spectrometer technique (GC-MS). The antibacterial activity of the extracts or imipenem was tested in vitro against *A. baumannii* isolates.

Materials and Methods

I. Sampling collection, isolation and processing

Ninety specimens; urine (n = 50), sputum (n = 40), and were randomly collected under complete aseptic conditions from different hospitals and clinics in Qena province, Egypt (From January to November 2019). Using sterile container, 5 ml of clean-catch mid-stream urine was collected from patients presumptive for urinary tract infection (UTI). Urine samples were inoculated on MacConkey agar and/or Blood agar (Merck, Germany) at 37°C for 24 hrs and observed for bacterial growth. Blood agar colonies were counted using colony counter and checked for significant bacteriuria. Culture that grew $\geq 10^5$ CFU/ ml, were measured as a significant bacteriuria.

For heterogeneous colonies, sub-culturing of individual distinct colonies was performed to ensure pure cultures. Sputum samples were taken from patients over a self-induced cough into sterile cups and sent for culture. Samples initially identified using standard laboratory methods including growth on MacConkey agar and/or Blood agar (Merck, Germany) and Gram staining. Plates were incubated at 37°C and examined for visible bacterial growth after 48 hrs of incubation. The isolated bacteria were further identified using the automated system VITEK-2 (bioMérieux, France).

II. Phenotypic identification of the isolates by ViteK-2 systems

All bacterial isolates were identified through morphological, conventional biochemical tests with Automated Identification (Biomerieux) Vitek-2 System. Forty-one tests, including 18 tests for sugar assimilation, 18 tests for sugar fermentation and 2 decarboxylase tests, and 3 miscellaneous tests (for urease, utilization of malonate, and tryptophane deaminase). With a vacuum device, the cards are inoculated with a 0.5 McFarland suspension of the organism prepared from 18- 20 hours-old blood agar plate (bioMe'rieux) and are then automatically sealed and manually inserted inside the Vitek-2 reader. Inoculators fluorescence is measured every 15 min, and the results of identification are determined after 3 h (Joyanes et al., 2001).

III. 2.3. Antimicrobial susceptibility testing

The antibiograms for all the recovered isolates were

determined by disk diffusion method (Bauer *et al.*, 1966). Each bacterial isolates was inoculated in tryptic soy broth and incubated at 37°C overnight for activation. Bacterial inoculum was adjusted to 0.5 McFarland turbidity, which equal 1.0×10^8 CFU/ml.

The antimicrobial agents used belonged to five different classes of antimicrobials; the aminoglycosides (Amikacin 30 ug/disc, Tobramycin 10 ug/disc and Gentamicin 10 ug/disc), β-lactams (Ticarcillin 100 ug/disc, Ticarcillin/Clavulanic Acid 75/10 ug/disc, Piperacillin 100 ug/disc, Piperacillin/Tazobactam 100/10 ug/disc, Cefotaxime 30 ug/disc, Cefepime 30 ug/disc, Imipenem 10 mg/disc and Meropenam 10mg/disc), Fluoroquinolones (Ciprofloxacin 5ug/disc Levofloxacin 5 and μg), antimetabolites (Trimethoprim/Sulfamethxazole 1.25/23 ug/disc) and Tetracycline (Minocycline 30ug/disc). MIC was carried out in triplicate and average MIC was calculated, as recommended by the (CLSI, 2018). A. baumannii isolates with the same phenotype and antibiotic pattern profile were excluded, only different isolates were considered for further experiments.

IV. Plant material and extraction

Curcuma longa (Turmeric) was purchased from local market in Nag hammadi center, Egypt. Three solvents including distilled water, ethanol, and ethyl acetate (Piochem, Egypt), were used as follow. 20 grams of plant powder was soaked in 200 ml of each solvent under stirring conditions (150 rpm), for three consecutive days at room temperature by using bigger bill shaker, USA. The solutions were filtred through Whatman # 1 sterile filter paper and collected in a preweighed beaker. The solvent was allowed to evaporate at room temperature tills complete evaporation except for aqueous extract that was concentrated in a water bath until a constant dark sticky residue was obtained. The dried precipitates were dissolved in di-methylsulphoxide (DMSO), (10 % v/v) and, then stored in airtight sterilized bottles at - 4°C.

V. Determination of the minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) by INT formazone assay

The determination of MIC and MBC were assayed as described earlier (Salem *et al.*, 2017). The freshly prepared culture of *A. baumannii* isolates was adjusted to OD595 of 0.01. 100 μl of each isolate culture was put into sterilized 96-well plates. Then 20 μl of the original *C. longa* extracts (100 mg ml⁻¹) was added (serial dilutions of 10⁻¹ -10⁻¹⁰ were used, 8 replicates were made for each dilution into complete raw of the 96-well plate). Imipenem (512 mg ml⁻¹) was tested as a positive

control; un-inoculated media tested as negative control after 24 hrs incubation at 37°C. MIC was determined by the addition of 40 μl of p-iodonitrotetrazolium violet chloride (INT) (0.2 mg/ml, Sigma-Aldrich) to the plates and re-incubated at 37°C for 30 min., the lowest concentration which banned color change is the MIC (Eloff, 1998 & Lall *et al.*, 2013). The MBC was determined by transferring 50 ul from each well of an overnight MIC plate (and/or higher) to sterile (TSA) fresh plates. Viable colonies were counted after 24 h at 37°C. The limit of detection for this assay was 101 CFU/ml.

VI. Gas chromatography-mass spectrometry (GC-MS) analysis

Three different extracts of *C. longa* were subjected to gas chromatography-mass spectrometer technique (GC-MS) (Thermo scientific technologies, Trace 1310-ISQ) with capillary column TG-5 (30m×250µm×0.25µm) system were used. The mass detector used in split mode and helium gas with a flow rate of 1.5 ml/min was used as a carrier. The injector was operated at 230°C and the oven temperature for the initial setup was 60°C for 2 min. ramp 10/min. to 300°C for 8 min. Mass spectra were taken at 70eV, total GC running time was 35 min.

VII.Statistical analysis

The variability degree of results was expressed in form of Means \pm Standard Deviation (Mean \pm S.D). GraphPad Prism 6.0 (GraphPad Software, San Diego, CA, USA, https://www.graphpad.com) was used to calculate one way analysis of the variance (ANOVA) with multiple comparison tests (Tukey's) to evaluate the effect of the different extract. Differences were considered significant at p values of \leq 0.05.

Result

Patients and Clinical Characteristics

A total of 90 patients with clinical evidence of nosocomial infection (elevated liver and/ or kidney functions, respiratory diseases, etc.) were enrolled in the study (Table 1). Of them, 51 (56.66) were females and 39 (43.3%) were males. The majority (48.9 % - 29 female; 15 male) of participants were found in the age group between 30- 60 years. Twinty six patients (7 female; 19 male) with a percentage of approximately (28.9 %) of the study participants were above 60 years old. While a percentage of (22.2 % - 18 female; 6 male) were below 30 years old. Table (1) depicts the clinical characteristics of the study participant.

The prevalence of A. baumannii isolates among Isolated clinical samples

Among the 90 clinical specimens, the incidence of the isolated bacteria in the clinical samples was illustrated in Table 2 and Figure 1. The most common prevalent organism (from each corresponding clinical samples) was Klebsiella pneumonia with a percentage of 45.5% followed by Escherichia coli 39.9%, A.baumannii 11.1% and P. aeruginosa 4.5% (Figure 1A). The total bacterial count in urine culture showed that 70% (21/30) of female patients had a range of 10⁴ to 10⁵ CFU/ml compared to 65% (13/20) of male patients. While 5% of female and 2% of male patients had a CFU values ≥ 106 (data not shown). Isolates from urine samples showed that E. coli seems to be the predominant with (32/50; 64.0%), followed by K. pneumonia (11/50; 22.0%), P. aeruginosa (4/50; 8.0%) and A. baumannii with (3/50; 6.0%), (Figure 1B). The respiratory tract isolates with K. pneumonia showed to be predominant with (30/40; 75.0%), A. baumannii (7/40; 17.5%) followed by E. coli (3/40; 8.82 %) (Figure 1C).

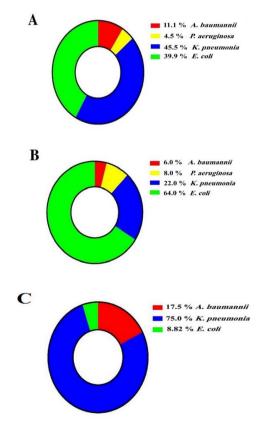


Fig. 1. The prevalence of A. baumannii isolates among Isolated clinical samples. (**A**) the most common prevalent bacterial isolates from each corresponding clinical samples; (**B**) Isolates from urine samples; and (**C**) respiratory tract isolates.

Table 1. Data and patients characteristics for examined clinical samples

	Patient character	Ma	le	Female		
	i attent character	No = 39	43.3%	No = 51	56.66%	
Ago	> (30)	5	12.82	15	29.41	
Age (year)	between (60 – 30)	15	38.46	29	56.86	
(year)	< (60)	19	48.71	7	13.72	
	Prevalence of underl	ying diseas	se			
Diabetes m	ellitus	5	10.63	5	9.8	
Diabetic an	d Hypertension	8	21.3	16	31.37	
Respiratory	disease	2	6.38	6	11.76	
Diabetic an	d respiratory disease	5	10.63	9	17.64	
Diabetic w	abetic with respiratory disease and hypertension		17.02	7	13.74	
Patients wi	tients with elevated liver function test		10.63	6	11.76	
Patients wi	th elevated kidney function test	7	19.16	2	4.1	
Catheter pr	resence	2	10.63	5	9.8	

Table 2. The incidence of Acinetobacter baumannii isolates among examined clinical samples

	Type of	No. of isolates						
No. of specimen	sample	Acinetobacter baumannii	Pseudomonas aeruginosa	Klebsiella pneumonia	Escherichia coli			
Urine	50	3 (6.0 %)	4 (8.0 %)	11 (22.0%)	32 (64.0%)			
Respiratory	40	7 (17.5 %)	0 (0.0 %)	30 (75.0%)	3(8.82 %)			
Total No. of isolates	90	10 (11.1%)	4 (4.5%)	41 (45.5%)	35 (39.9%)			

Minimum inhibitory concentration MIC (mm)

Ten isolates of *A. baumannii* (11.1%) were recovered from sputum samples (n= 7; 70%), and urine (n = 3; 30%). Antibiotic resistance of *A. baumannii* represented by (Table 3; Figure 2) against 15 antibiotics with different disc potency, was determined that all *A. baumannii* isolates were resistance to Ticarcillin, Ticarcillin/Clavulanic acid, Piperacillin, Piperacillin/Tazobactam, Cefotaxime, Cefepime, Levofloxacin, Ciprofloxacin and Amikacin. While 80

are resistant to Gentamicin, Tobramycin respectively, 70 % are against Imipenem Meropenam. Resistance decreased to 60% Trimethoprim/Sulfamethoxazole and 40 % with Minocycline respectively. For out of 10 A. baumannii isolates (codes A4, A7, A9 and A10) were found to share the same phenotype and antibiotic pattern profile. Hence, these isolates were excluded. The rest isolates that showed different phenotype and antibiotic pattern are A1, A2, A3, A5, A6 and A8 were considered for further experiments.

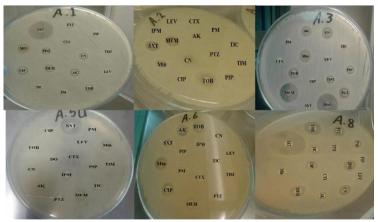


Fig. 2. Antibiotic susceptibility test of Acinetobacter baumannii.

*TIC, Ticarcillin: TIM, Ticarcillin/Clavulanic Acid: PIP, Piperacillin: PTZ, Piperacillin/ Tazobactam: CTX, Cefotaxime: PM, Cefepime: IPM, Imipenem: MEM, Meropenam: AK, Amikacin: CN, Gentamicin: TOB, Tobramycin: LEV, Levofloxacin: CIP, Ciprofloxacin: Min, Minocycline: CT, Colistin SXT, Trimethoprim/Sulfamethxazole. A1: A8 are the A.baumannii isolates code. A1 and A2 for urine isolates, sample no: A3, A5, A6 and A8 for respiratory isolates.

Antibiotics Isolates No.* $A\overline{3}$ used* **A2 A4 A7 A8 A9** A10 A1 **A5** TIC 0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)TIM 0(R)0(R)0(R)0(R)0(R)0(R) 0(R)0(R)0(R)0(R)PIP 0(R)0(R)0(R)0(R) 0(R) 0(R) 0(R)0(R)0(R)0(R)0(R) 0(R) PTZ 0(R) 0(R)0(R)0(R) 0(R)0(R)0(R)0(R) CTX 0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)0(R)PM 0(R)0(R) 0(R) 0(R)0(R) 0(R) 0(R) 0(R) 0(R) 0(R) IPM 0(R)0R26(S)23(S) 0(R)0(R)0(R)0(R)0(R)19(S) **MEM** 0(R)12(R) 15(I) 0(R)15(I) 0(R)0(R)16 (I) 0(R)0(R)0(R)0(R) AK 9(R) 0(R)0(R)0(R)19(S) 0(R)0(R)0(R)CK 0(R)0(R)14(I) 0(R)0(R)0(R)0(R) 14(I) 0(R)0(R)TOB 0(R) 0(R) 0 (R) 0(R)0(R) 16(S) 0(R) 16(S) 0(R) 0(R) LEV 0(R) 0(R) 0 (R) 0(R) 0(R) 16(S) 0(R) 16(S) 0(R) 0(R) CIP 0(R)0(R) 0(R)0(R)0(R) 0(R)0(R) 0(R)0(R) 0(R)23(S) 12(I) 22(S) Min 0(R)0(R)21(S) 0(R)0(R)14(I) 23(S) SXT 0(R)26(S) 0(R)0(R)27(S) 0(R)23(S) 0(R)0(R)19(S)

Table 3. Minimum inhibitory concentration MIC (mm).

Gas Chromatography-Mass Spectrometry (GC/MS) Analysis

The main bioactive components obtained by GC-MS analysis of *C. longa* aqueous extract, revealed the presence of 26 compounds, were identified and accounted for 95.78 % of the total area. The mean compounds identified was Turmerone (55.42%). While intermediate compound was B-myrcene (8.12%), and D-Limonene (8.04%). The minor compounds identified in *C. longa* aqueous extract were p-cymene (4.14%), and other compounds were presented in (Table 4;

Figure 3). Ethanolic extract were identified to possesses 18 signal peaks for 18 compounds present and accounted for 99.81% of the total. Turmerone (59.42%), alpha pinene (12.07%) and Citral (11.68%), were identified as a major compound. Ethylacetate extract, illustrated the presence of 24 compounds, were accounted 97.58% of the total area obtained by analysis of *Curcuma longa* extract. Caryophyllene (16.66%), Caryophyllene oxide (15.62%), (-)-isolongifolol (12.14%), Curdione (11.0%), B-Sesquiphellandrene (10.47%), and identified as a major compound.

Table 4. Bioactive components identified by GC-MS of Curcuma longa extract *

Aqueous extract										
No	RT* (min)	Compound Name	M. formula*	M. wt*	Area (%)					
1	5.33	β-pinene	C ₁₀ H ₁₆	136	1.55					
2	7.28	B-myrcene	C ₁₀ H ₁₆	136	8.12					
3	9.08	D-Limonene	C ₁₀ H ₁₆	136	8.04					
4	13.74	eucalyptol	C ₁₀ H ₁₈ O	154	1.89					
5	14.42	Linalyl propionate	C ₁₃ H ₂₂ O ₂	210	0.74					
6	16.08	Cubebenes	C ₁₅ H ₂₄	204	0.20					
7	16.67	Decanoic acid, methyl ester	C ₁₁ H ₂₂ O ₂	185	0.18					
8	17.12	beta – basbolene	C ₁₅ H ₂₄	204	0.66					
9	17.79	Beta - elemene	C ₁₅ H ₂₄	204	1.01					
10	18.54	Phytol ketone	C ₁₈ H ₃₆ O	268	0.10					
11	21.10	beta -sesquiphellandrene	C ₁₅ H ₂₄	204	6.80					
12	21.19	Beta -farnesene	C ₁₅ H ₂₂	202	0.68					
13	22.03	Curcumene	C ₁₅ H ₂₄	204	1.42					
14	22.23	Calarene epoxide	C ₁₅ H ₂₄ O	220	0.16					
15	23.75	Farnesol (E), methyl ether	C ₁₆ H ₂₈ O	236	0.47					
16	24.32	Globulol	C ₁₅ H ₂₆ O	222	0.14					
17	24.81	Geranyl-2-methylbutyrate	C ₁₅ H ₂₆ O ₂	238	1.76					
18	26.20	Cubenol	C ₁₅ H ₂₆ O	222	0.15					
19	28.78	Caryophyllene oxide	C ₁₅ H ₂₄ O	220	2.01					
20	30.15	trans-Valerenyl acetate	C ₁₇ H ₂₆ O ₂	262	0.17					
21	30.41	Corymbolone	C ₁₅ H ₂₄ O ₂	236	0.94					
22	34.84	Turmerone	C ₁₅ H ₂₂ O	218	55.42					
23	36.35	Eupatoriochromene	C ₁₃ H ₁₄ O ₃	218	0.76					
24	37.13	Phytol	C ₂₀ H ₄₀ O	296	0.20					
25	42.93	Anthraquinone	C ₁₄ H ₈ O ₂	208	1.6					
26	53.48	Heptacosane	C ₂₇ H ₅₆	380	0.65					
Total 95.78										

Table 4 (Continued). Bioactive components identified by GC-MS of Curcuma longa extract *

Ethanolic extract										
NO	RT* (min)	Compound Name	M. formula*	M. wt*	Area (%)					
1	17.79	Limonene	C ₁₀ H ₁₆	136	0.26					
2	18.04	alpha phellandrene	$C_{10}H_{16}$	136	0.44					
3	19.54	alpha-pinene	C ₁₀ H ₁₆	136	12.07					
4	20.14	Eucalyptol	C ₁₀ H ₁₆	136	3.44					
5	21.01	Citral	C ₁₀ H ₁₆ O	152	11.68					
6	26.07	Terpinen-4-ol	C ₁₀ H ₁₈ O	154	2.02					
7	26.63	Ascaridole	C ₁₀ H ₁₆ O ₂	168	0.57					
8	31.21	7-epi-cis-sesquisabinene hydrate	C ₁₅ H ₂₆ O	222	0.13					
9	32.54	Caryophyllene	C ₁₅ H ₂₄	204	0.17					
10	33.74	cis-Farnesol	C ₁₅ H ₂₆ O	222	2.45					
11	34.36	Butylated Hydroxytoluene	C ₁₅ H ₂₄ O	220	3.30					
12	34.83	α-cedrene	C ₁₅ H ₂₄	204	0.18					
13	35.09	γ-curcumene	C ₁₅ H ₂₄	204	0.09					
14	36.82	7-epi-cis-sesquisabinene hydrate	C ₁₅ H ₂₆ O	222	0.97					
15	37.47	Caryophyllene oxide	C ₁₅ H ₂₄ O	220	0.64					
16	37.74	β-curcumene	C ₁₅ H ₂₄	204	0.38					
17	41.78	germacrone	C ₁₅ H ₂₂ O	218	1.60					
18	42.33	Tumerone	C ₁₅ H ₂₂ O	218	59.42					
		Total 99.		•	•					
Ethyl acetate extract										
NO	RT* (min)	Compound Name	M. formula*	M. wt*	Area (%)					
1	8.48	1- dimethyl(isopropyl)silyloxy-2- phenylethane	C ₁₃ H ₂₂ OSi	222	0.13					
2	15.67	1,8-cineole	C ₁₀ H ₁₈ O	154	1.44					
3	25.59	alfaCopaene	$C_{15}H_{24}$	204	1.81					
4	25.79	Zingiberene	C ₁₅ H ₂₄	204	1.57					
5	25.95	Corymbolone	$C_{15}H_{24}O_2$	236	0.40					
6	26.73	Beta-Caryophyllene	C ₁₅ H ₂₄	204	16.66					
7	26.88	Ylangene	C ₁₅ H ₂₄	204	0.77					
8	27.11	Alloaromadendrene	C ₁₅ H ₂₄	204	0.44					
9	27.44	B-Curcumene	C ₁₅ H ₂₄	204	2.36					
10	27.68	B -Sesquiphellandrene	C ₁₅ H ₂₄	204	10.47					
11	28.08	agarospirol	C ₁₅ H ₂₆ O	222	1.67					
11 12	28.08 28.26	agarospirol B-bisabolene	C ₁₅ H ₂₆ O C ₁₅ H ₂₄	222 204	1.67 0.81					
11 12 13	28.08 28.26 28.69	agarospirol B-bisabolene Copaene	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄	222 204 204	1.67 0.81 5.13					
11 12 13 14	28.08 28.26 28.69 28.91	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O	222 204 204 220	1.67 0.81 5.13 3.40					
11 12 13	28.08 28.26 28.69	agarospirol B-bisabolene Copaene	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₄	222 204 204	1.67 0.81 5.13					
11 12 13 14 15 16	28.08 28.26 28.69 28.91 29.20 29.57	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ C ₁₅ H ₂₆ O	222 204 204 220 204 222	1.67 0.81 5.13 3.40 1.14 0.24					
11 12 13 14 15 16 17	28.08 28.26 28.69 28.91 29.20 29.57 30.61	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O	222 204 204 220 204 222 222	1.67 0.81 5.13 3.40 1.14 0.24 12.14					
11 12 13 14 15 16 17	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O	222 204 204 220 204 222 222 236	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0					
11 12 13 14 15 16 17 18 19	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96 31.27	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione (-)-Spathulenol	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₄ O ₂ C ₁₅ H ₂₄ O ₂	222 204 204 220 204 222 222 222 236 220	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0 6.18					
11 12 13 14 15 16 17 18 19 20	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96 31.27 32.55	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione (-)-Spathulenol Elemene	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₄ O ₂ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O	222 204 204 220 204 222 222 236 220 204	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0 6.18 1.62					
11 12 13 14 15 16 17 18 19 20 21	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96 31.27 32.55 33.20	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione (-)-Spathulenol Elemene cis-11-Eicosenoic acid	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₄ O ₂ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O C ₁₅ H ₂₄ C	222 204 204 220 204 222 222 236 220 204 310	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0 6.18 1.62 0.70					
11 12 13 14 15 16 17 18 19 20 21 22	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96 31.27 32.55 33.20 33.37	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione (-)-Spathulenol Elemene cis-11-Eicosenoic acid Caryophyllene oxide	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₄ O ₂ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O	222 204 204 220 204 222 222 236 220 204 310 220	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0 6.18 1.62 0.70 15.62					
11 12 13 14 15 16 17 18 19 20 21	28.08 28.26 28.69 28.91 29.20 29.57 30.61 30.96 31.27 32.55 33.20	agarospirol B-bisabolene Copaene Cis-Z-a-bisabolene epioxide B-CADINENE Cubedol (-)-isolongifolol Curdione (-)-Spathulenol Elemene cis-11-Eicosenoic acid	C ₁₅ H ₂₆ O C ₁₅ H ₂₄ C ₁₅ H ₂₄ C ₁₅ H ₂₄ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₆ O C ₁₅ H ₂₄ O ₂ C ₁₅ H ₂₄ O C ₁₅ H ₂₄ O C ₁₅ H ₂₄ C	222 204 204 220 204 222 222 236 220 204 310	1.67 0.81 5.13 3.40 1.14 0.24 12.14 11.0 6.18 1.62 0.70					

*RT: Retention time per minute; active compound detected by GC/MS; area (%) percentage of compound; *M. Formula: Molecular formula; *M. wt: molecular weight of the compound (g/mol).

In vitro assay for the antibacterial activity of Curcuma longa extract

Antibacterial activity of *C. longa* different extracts and Imipenem against *A. baumannii* isolates was analyzed by minimal inhibitory concentrations (MIC) by determining the bacterial viability using a colorimetric

INT-formazan assay. As a result, we determined the minimal bactericidal concentrations (MBC) which confirmed the killing of *A. baumannii* isolates over time (24 hrs). The individual use of *C. longa* aqueous and/or ethyl acetate extracts against *A. baumannii* isolates (A1, A2, A3, A5, A6, and A8) exhibited MBC values

varying from 40 to 100 mg mL⁻¹, respectively in (Table 5 and figure 4). Within isolates A1 and A8, ethanol extracts revealed MBC values (40 to 90 mg ml⁻¹) respectively when compared to imipenem figure 4A, B

and C. It is worth mentioning that Imipenem showed a great bactericidal efficacy against all isolates with a concentration of 0. 2 mg mL⁻¹ are adequate to kill all the tested *A. baumannii* isolates.

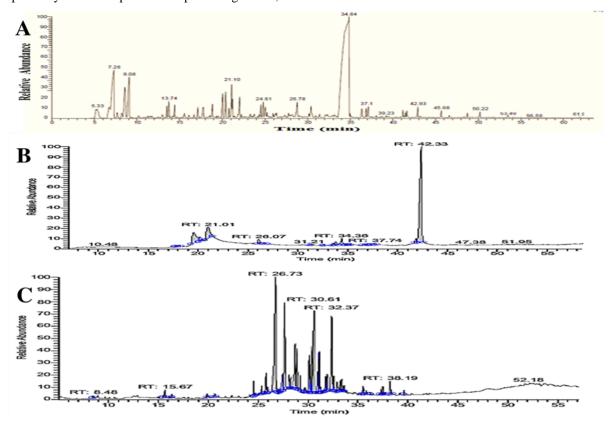


Fig. 3. Chromatogram obtained from GC/MS of *Curcuma longa*. (A): for aqueous extract, (B): for ethanolic extract and (C): for ethyl acetate extract.

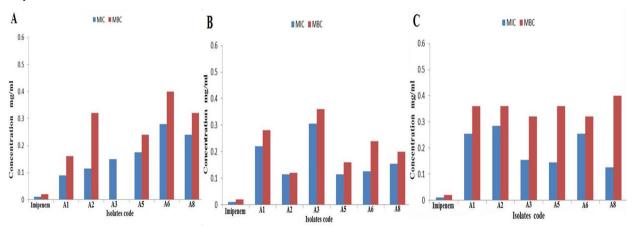


Fig. 4. MIC and MBC of Imipenem and *Curcuma longa* against *A. baumannii* isolates. MIC (blue) and MBC (red). *Curcuma longa* different extracts (100 mg ml⁻¹) (A): aqueous extract; (B): ethanolic extract; and (C): ethyl acetate extract. (Imi): Imipenem (0.512 mg ml⁻¹). A1: A8 are the *A. baumannii* isolates code. A1 and A2 isolates from urine sample no: A3, A5, A6 and A8 isolates for respiratory sample.

Table 5. MIC and MBC of *Curcuma longa* extracts*.

Tested	*MIC and MBC of Curcuma longa aqueous extract (100 mg ml ⁻¹)												
strains No.	R1	R2	R3	R4	R5	R6	R7	R8	MIC	SD	MIC+SD	MIC-SD	MBC
A1	20	20	20	20	20	20	30	30	22.5	4.6291	27.1291	17.8709	40
A2	30	20	30	30	30	30	30	30	28.75	3.535534	32.28553	25.21447	80
A3	30	30	30	40	40	40	40	50	37.5	7.071068	44.57107	30.42893	100
A5	40	40	40	40	40	50	50	50	43.75	5.175492	48.92549	38.57451	60
A6	70	70	70	70	70	70	70	70	70	0	70	70	100
A8	60	60	60	60	60	60	60	60	60	0	60	60	80
Tested	MIC and MBC of Curcuma longa ethanolic extract (100 mg ml ⁻¹)												
strains No.	R1	R2	R3	R4	R5	R6	R7	R8	MIC	SD	MIC-SD	MIC+SD	MBC
A1	50	50	50	60	60	60	60	50	55	5.345225	49.65478	60.34522	70
A2	20	30	30	30	30	30	30	30	28.75	3.535534	25.21447	32.28553	30
A3	70	70	70	80	80	80	80	80	76.25	5.175492	71.07451	81.42549	90
A5	20	30	30	30	30	30	30	30	28.75	3.535534	25.21447	32.28553	40
A6	30	30	30	30	40	30	30	30	31.25	3.535534	27.71447	34.78553	60
A8	30	40	40	40	40	40	40	40	38.75	3.535534	35.21447	42.28553	50
Tested strains				MIC	and M	BC of (Curcum	a longa	ethylace	etate extract	(100 mg ml ⁻¹)		
No	R1	R2	R3	R4	R5	R6	R7	R8	MIC	SD	MIC-SD	MIC+SD	MBC
A1	60	60	60	60	60	70	70	70	63.75	5.175492	58.57451	68.92549	90
A2	60	70	70	70	70	70	80	80	71.25	6.408699	64.8413	77.6587	90
A3	30	40	40	40	40	40	40	40	38.75	3.535534	35.21447	42.28553	80
A5	30	30	30	40	40	40	40	40	36.25	5.175492	31.07451	41.42549	90
A6	60	60	60	60	60	70	70	70	63.75	5.175492	58.57451	68.92549	80
A8	20	30	30	30	30	30	40	40	31.25	6.408699	24.8413	37.6587	100
Tested strains						MIC	and ME	Cof Im	ipenem (5	512 mg ml ⁻¹)			
No	R1	R2	R3	R4	R5	R6	R7	R8	MIC	SD	MIC-SD	MIC+SD	MBC
A1	0.1	0.2	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.035	0.147	0.077	0.2
A2	0.1	0.1	0.11	0.1	0.1	0.1	0.1	0.1	0.1	0.003	0.104	0.097	0.4
A3	0.1	0.1	0.1	0.1	0.1	0.11	0.1	0.1	0.1	0.003	0.104	0.097	0.2
A5	0.2	0.2	0.2	0.2	0.2	0.2	0.22	0.2	0.2	0.007	0.209	0.195	0.2
A6	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.11	0.1	0.003	0.104	0.097	0.4
A8	0.1	0.1	0.1	0.1	0.1	0.1	0.11	0.1	0.1	0.003	0.104	0.097	0.2

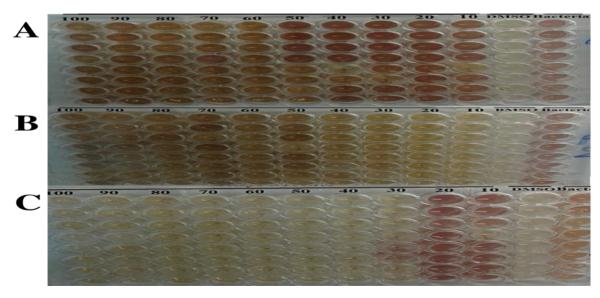


Fig. 5. MIC of *Curcuma longa* extract against *A. baumannii* isolates.(A):for aqueous extract; (B):for ethanolic extract; (C): for ethyl acetate extract. A1: A8 are the *A. baumannii* isolates code. A1 and A2 isolates from urine sample no: A3, A5, A6 and A8 isolates for respiratory sample.

Discussion

Antibiotic-resistance risks in particular of nosocomial infections with *A. baumannii*, *P. aeroginosa* are increasing and treatment of bacterial infection especially multi-drug resistant become the major of concern (Ibrahim *et al.*, 2021). The current study showed the proportion of nosocomial infections due to the gram-negative bacilli *A. baumannii* isolated from clinical samples as urine (6.0%), and sputum (17.5%) among the patients (Figure 1). A total 90 patient were included, categorized to male 43.12%, female 56.88% patients.

Demographic data, risk factors for A. baumannii (Table 1). A. baumannii are a common cause of urinary tract infections (UTI) that upset the kidney, leading to pyelonephritis, as well as the bladder, resulting in cystitis (Thomson & Armitage, 2010). UTI symptoms included elevated kidney function and high levels of A. baumannii in the patients' sputum (70%). This result is conceivable since UTI symptoms are not a reliable indication of illness, the presence of bacteria must be confirmed by urine culture in order to diagnose UTI (Elamary et al., 2020). The urine culture in our results indicated the presence of bacteriuria in 70% of female patients and 65% of male patients (Table 1). The percentage of A. baumannii in our study agreed with (Al-Agamy et al., 2014). The incidence of A. baumannii isolates among examined clinical samples was (10/110; 9.09%), K. pneumonia was the predominant with (49/110; 44.5%), E. coli (46/110; 41.8%) and P. aeruginosa (5/110; 4.5%) (Fouad et al., 2013; Sarangi et al., 2017). The highest prevalence reported for A. baumannii isolates (Table 3; Figure 1) were recovered from sputum samples (7/40; 17.5%), followed by urine (2/50; 4.0%) and pus (1/20; 5.0%) (Abdulzahra et al., 2018). The antibiotic susceptibility test for A. baumannii isolates indicated that all isolates (n = 10; 100%) were multi-drug resistant to Ticarcillin, Ticarcillin/Clavulanic Acid, Piperacillin, Piperacillin/Tazobactam, Cefotaxime, Cefepime, Ciprofloxacin, and Amikacin, while 70-80% are extensively drug-resistant. Resistance decreased to 40% with Minocycline and all isolates were sensitive to Colistin (Figure 2). Our findings are in covenant with earlier results concerning the MDR isolates (Younis et al., 2019). Bactericidal activity of C. longa A very promising the best result were obtained with ethanolic extract with average MBC (43.12 mg ml-1), aqueous and ethyl acetate extract of C. longa (43.75 – 50.83 mg ml-1) (Table 5) (Ahmed et al., 2021).

(GC-MS) of *C. longa* extract (Table 4; Figure 3A, B and D) revealed the presence of 68 compounds, turmerone (55.42%), B-myrcene (7.45%), P-cymene (4.14%), D-

limonene (3.9%), β -sesquiphellandrene (2.19%). Aromatic turmerone and turmerone ($C_{15}H_{20}O$ and $C_{15}H_{22}O$), were identified as two major sesquiterpenes ketonic compound responsible for the aroma of turmeric by (Nair, 2019).

Antibacterial activity of C. longa, aromatic turmerone and turmerone were reported together (Negi et al., 1999; Lee et al., 2011). The minor compounds identified in our results p-cymene (4.14%), 1,8-cineole (1.05%), β -pinene (0.84%) and cis-sabinol (0.27%)agreed with (Jayaprakasha et al., 2005). Antibacterial activity of P-cymene, 1, 8-cineole, β-pinene and cissabinol against E.coli isolated from UTI patients were reported by (Lagha et al., 2019). Moghaddam et al. (2018) were reported antimicrobial activity of βocimene, other compound present in (Table 4) of Ferulago angulata essential oil. The major compounds identified by GC-MS of C. longa ethyl acetate extract were caryophyllene, caryophyllene oxide, (-)isolongifolol, curdione, B-sesquiphellandrene (Chowdhury et al. 2008). Rania et al. (2002) and Naz et al. (2010) reported Caryophyllene, susquiphellandrene as a major constituent of C. longa rhizome. Mehra & kumar. (2019) were reported antibacterial activity of compounds (-)-Spathulenol, Copaene, Cis-Z-abisabolene epioxide, β-curcumene, alfa-copaene agarospirol, zingiberene, 1,8-cineole and oleic acid in methanol extract of C. longa. Moo et al. (2020) revealed antibacterial activity β-caryophyllene against Bacillus cereus due to changing in membrane permeability and integrity leading to membrane damage and intracellular content leakage, which eventually caused cell death. Santos et al. (2019) regarded the mechanism of bactericidal of limonene, caryophyllene oxide, citral, trans caryophyllene, and 1, 8-cineole to the direct damage caused to the integrity of the cell membrane by the lipophilic components of the essential oil. Woo et al. (2020) revealed the inhibitory effects of β caryophyllene on Helicobacter pylori infection because of interruption of bacterial replication via down regulation genes by β-caryophyllene.

Conclusion

Nosocomial infection with *A. baumannii* has increased year by year in human medicine and it is also considered as a major cause of mortality worldwide, so there is a need to decrease the prevalence of nosocomial infections while improving the healthcare quality and patient safety. Although ethanol or/aqueous extract of *C. longa* was promising for the in vitro study against *A. baumannii*. GC/MS of C. longa reveal many active compounds that may act as antibacterial agent against MDR *A. baumannii*.

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