

Research Article

Burn Skin Pathogens: Isolation, Identification, Antimicrobial Activity Pattern Against *C. longa* Extract and Computational Studies of its Components

Essa Ajmi Alodeani

College of Medicine-Aldawadmi, Shaqra Univerity, Kingdom of Saudi Arabia

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ABSTRACT

C. longa extract was obtained and tested against the burn skin pathogens such as *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *acinetobacter baumannii*, *Klebsiella* spp, *Proteus vulgaris* and *Escherichia coli*. The finding of antimicrobial screening disclosed that the methanol extract produced more effect than hexane extract. All the components in the extracted were examined for the drug likeness physicochemical properties, which depicted that all are in compliance with the Lipinski Rule of five except component six and all possessed bioactivity score in most active zone except component four that corresponds to moderately active zone.

Key words: *C. longa* extract, antimicrobial screening, drug likeness and physicochemical property.

INTRODUCTION

Medicinal plants has been proved as the best source for obtaining drugs and also used as traditional medicine in developed countries, as declared by WHO¹. *Curcuma longa*, normally known as turmeric is the member of Zingiberaceae family and a perennial rhizomatous shrub². It's flavouring in spicy quality and digestive ability for foods prompting its continuous use. The plant comprises the funnel shape yellow flower, oblong pointed leaves. The part rhizome, which is used as medicine followed boiling, cleaning drying and giving rise to a yellow powder^{2,3}. It has been known to possess variety of biological activities antioxidant, hepatoprotective, anti-inflammatory, gastrointestinal, anticarcinogenic, cardiovascular, antimicrobial. It is also found to have potential for the treatment of *cholelithiasis*, *and cholestasis*, *hyperlipidemia*, *gastric ulcer*, *chronic anterior uveitis*, treatment of flatulence, jaundice, menstrual difficulties, hematuria, hemorrhage, and colic. The *C. longa* extract possesses desmethoxy curcumin, disdesmethoxycurcumin, curcumin, turmerone, atlantone, zingiberine, sugars, proteins, and resins. The most investigated constituent is curcumin, that occupied 0.3-5.4 percent in turmeric^{4,25}. *C. longa* extraction, phytochemical screening and antimicrobial studies has been widely investigated by the researchers^{16,17,22-24}. Infection is significant reason for mortality and morbidity in burn patients³⁸. The rate of nosocomial infections is more in burn patients because of various factors such as nature of burn and immune system³⁸⁻⁴⁰. Our previous investigation, we also have carried out the isolation, identification of Burn skin pathogens and antimicrobial activity pattern against pyrazole derivatives²⁶. It is also

important to find out computationally that which component is more potent biologically active³⁰⁻³⁷. Keeping in mind the pharmacological potential of *C. Long*, and the necessity for the new anti microbial agents that can be utilized to diminish the infections caused by burn skin pathogens, we designed our investigation in such a way that deals with the extraction, antimicrobial assay and also the computational studies.

MATERIALS AND METHODS

Pathogens from Burn skin

Fifty pus samples with an average age of 10-55 years were collected, isolated maintained and preserved using 1% glycerol stock. The phenotypic characterization of the pathogens from burn skin was performed applying the procedure reported in our previous study²⁶.

Phytochemical extraction

The extraction *C. long* was achieved employing Soxhlet extraction assembly, methanol and n-hexane solvent. The Fresh rhizomes of *C. longa* were grounded and fixed inside a thimble loaded into the soxhlet extractor for 15 hr in the solvent (300 mL) with refluxing at the boiling temperature of solvent. Finally the solvent was evaporated by vacuum evaporator. Obtained extract was incubated at 40 °C for 24 hr, after that stored at 4 °C evaluation for antimicrobial screening^{4,25}.

Antimicrobial screening

Organism culture and in vitro screening for antimicrobial activity of *C. longa* extract against the isolated pathogens, was done by the disk diffusion method with minor modifications. The isolated and biochemically characterized pathogens: *Pseudomonas aeruginosa*,

Table 1: Representing the zone of inhibition of the methanol extract of *Curcuma longa*, against the isolated pathogens from burn patient.

Pathogens	Effect of methanol extract on Microorganism				
	50 µg/ml	25 µg/ml	12.5 µg/ml	6.25 µg/ml	3.125 µg/ml
<i>P. aurigenosa</i>	24.20±0.24	18.26±0.26	28.22±0.12	12.34±0.42	10.86±0.61
<i>S. aureus</i>	20.12±0.22	18.32±0.62	16.64±0.36	11.30±0.72	9.82±0.45
<i>P. vulgaris</i>	21.16±0.32	20.36±0.43	21.44±0.35	16.64±0.45	15.32±0.54
<i>E. coli</i>	20.22±0.39	23.28±0.32	20.43±0.30	20.22±0.33	21.42±0.36
<i>Klebsiella spp.</i>	17.92±0.20	18.30±0.52	19.14±0.34	12.20±0.62	11.23±0.34
<i>A. baumannii</i>	13.26±0.22	17.52±0.24	16.44±0.74	10.24±0.53	8.43±0.22

Table 2: Representing the zone of inhibition the n-hexane extract of *Curcuma longa*, against the isolated pathogens from burn patient.

Pathogens	Effect of n-hexane extract on Microorganism				
	50 µg/ml	25 µg/ml	12.5 µg/ml	6.25 µg/ml	3.125 µg/ml
<i>P. aurigenosa</i>	20.12±0.32	15.22±0.14	13.08±0.10	12.10±0.22	10.08±0.14
<i>S. aureus</i>	18.34±0.32	13.64±0.20	10.16±0.42	10.12±0.18	-
<i>P. vulgaris</i>	15.34±0.15	11.06±0.21	10.21±0.20	-	-
<i>E. coli</i>	17.14±0.39	15.43±0.30	12.23±0.30	10.13±0.12	-
<i>Klebsiella spp.</i>	16.62±0.26	14.23±0.42	10.17±0.20	-	-
<i>A. baumannii</i>	11.14±0.30	10.12±0.34	-	-	-

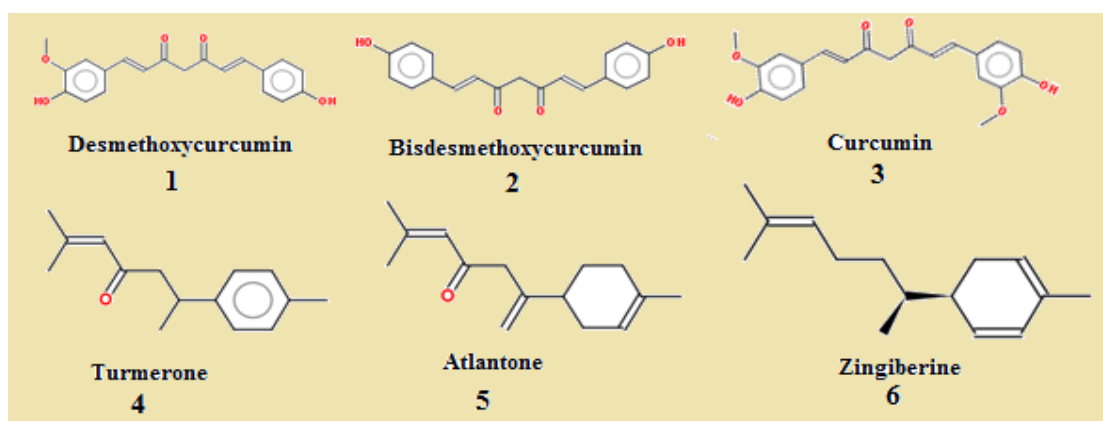
Figure 1: Exhibiting the components present in *C. longa* extract.

Table 3: Representing the zone of inhibition of the standard, methanol and hexane, against the isolated pathogens from burn patient.

Microorganism	Ciprofloxacin (10 µg/ml)	Methanol	Hexane
<i>P. aurigenosa</i>	34.24 ±0.31	-	-
<i>S. aureus</i>	21.46 ±.31	-	-
<i>P. vulgaris</i>	24.56±0.27	-	-
<i>E. coli</i>	23.82±0.47	-	-
<i>Klebsiella spp.</i>	21.34±0.42	-	-
<i>A. baumannii</i>	18.76±0.30	-	-

Staphylococcus aureus, *Acinetobacter baumannii*, *Klebsiella spp.*, *Proteus vulgaris* and *Escherichia coli* were subcultured in nutrient agar medium and incubated for 18 h at 37 °C. Following the incubation the bacterial cells were suspended, according to the McFarland protocol in saline solution to produce a suspension of about 10⁵ CFU/mL. About 10 mL of this suspension was mixed with 10 mL of sterile antibiotic agar at 40 °C and poured on to an agar plate in a laminar flow cabinet. Five paper disks (6.0 mm diameter) were fixed onto nutrient agar

plate. Ciprofloxacin was used as a standard drug (positive control). Methanol and hexane poured disk was used as negative control. The susceptibility of the bacteria to the test extract was determined by the formation of an inhibitory zone after 18 h of incubation at 36 °C. The zone of inhibition was calculated by antibiotic zone scale. The results were compared with the negative and positive controls²⁷⁻²⁹.

Physicochemical properties

The drug likeness score was calculated considering partition coefficient (log P), molar refractivity, molecular weight, number of heavy atoms, number of hydrogen donor, number of hydrogen acceptor and number of violation. All the parameters were checked with the help of software Molinspiration drug-likeness score online (www.molinspiration.com)³⁰⁻³⁷.

Bioactivity score

The drugs are also checked for the bioactivity by calculating the activity score for GPCR ligand, ion channel modulator, kinase inhibitor, nuclear receptor ligand. All the parameters were checked with the help of software Molinspiration drug-likeness score online

Table 4: Representing the physicochemical properties of all the components found in *Curcuma longa* and ciprofloxacin.

Physicochemical property score	Components						Standard
	1	2	3	4	5	6	
miLogP	2.30	2.48	2.67	4.48	4.12	5.12	-0.701
TPSA	93.07	83.83	74.60	17.07	17.07	0.00	74.569
Natoms	27	25	23	16	16	15	24.0
MW	368.38	338.36	308.33	216.32	218.34	204.36	331.347
nON	6	5	4	1	1	0	6
nOHNH	2	2	2	0	0	0	2
Nviolations	0	0	0	0	0	1	0
Nrotb	8	7	6	4	4	4	3
Volume	332.18	306.64	281.09	230.32	237.06	234.35	285.46

Table 5: Representing the bioactivity score of all the components found in *Curcuma longa* and ciprofloxacin.

Bioactivity score	Components						Standard
	1	2	3	4	5	6	
GPCR ligand	-0.06	-0.04	0.00	-0.68	-0.32	-0.39	0.12
Ion channel modulator	-0.20	-0.20	-0.14	-0.46	0.14	-0.10	-0.04
Kinase inhibitor	-0.26	-0.26	-0.26	-1.36	-1.20	-0.84	-0.07
Nuclear receptor ligand	0.12	0.18	0.25	-0.14	0.42	0.23	-0.19
Protease inhibitor	-0.14	-0.14	-0.08	-0.80	-0.49	-0.71	-0.21
Enzyme inhibitor	0.08	0.10	0.15	-0.25	0.47	0.29	0.28

(www.molinspiration.com)³⁰⁻³⁷.

RESULTS AND DISCUSSION

Antimicrobial activity

Antibacterial assessment was done against all burn skin pathogens, subcultured in Nutrient Agar medium using ciprofloxacin as a standard drug, Methanol and n-hexane as negative control. The susceptibility of the bacteria to the tested extract was determined by the formation of an inhibitory zone after 18 h of incubation at 36 °C. The obtained results are reported in Table-1, Table-2 and Table-3, that represented that both the portion possessed antimicrobial sensitivity but the antimicrobial potential of methanol extract is more than the n-hexane extract.

Phytochemistry

Phytochemical evaluation of *C. longa*, confirmed the presence of components such as Desmethoxy curcumin, Bisdesmethoxycurcumin, curcumin, turmerone, atlantone, zingiberine⁴⁻²⁵, the structure for all the components are exhibited in figure-1.

Physicochemical properties

Lipinski's rule of five states that, in general, an orally active drug has not more than 5 hydrogen bond donors (OH and NH groups), not more than 10 hydrogen bond acceptors (notably N and O), molecular weight under 500 g/mol, partition coefficient log P less than 5, number of violation less than 4. All the components of *C. long* were found in compliance with Lipinski's rule of five except component six that exhibited the partition coefficient 5.12, the detailed results are reported in Table 4.

Bioactivity score

The bioactivity score was calculated for GPCR ligand, ion channel modulator, kinase inhibitor, nuclear receptor ligand, protease inhibitor and enzyme inhibitor. For average organic molecule the probability of bioactivity

score is more than 0.00 then it is active, -0.50 to 0.0 then moderately active and if less than -0.50 then inactive. The results for bioactivity score are presented in Table 5. The results portrayed that the bioactivity score for all the components mostly lying under the zone of active drugs while the component four belongs to moderately active region.

CONCLUSION

C. longa extract was achieved and analysed for antimicrobial activity against all the pathogens isolated from burn skin. The phytochemical components of the extract were subjected for computational studies such as physicochemical and drug likeness properties. The observed results stated that methanol extract of *C. longa* possessed more antimicrobial potential than n-hexane extract, out of six only one component that is component-6 deviated from Lipinski's rule of five. All components lying under the zone for active drugs, except the component 4 that belongs to moderately activity zone for drugs.

Conflict of Interest

The authors have no conflict of interests

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