

King Saud University

Arabian Journal of Chemistry

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ORIGINAL ARTICLE

Phytochemical, Antimicrobial, Antidiabetic, Thrombolytic, anticancer Activities, and *in silico* studies of *Ficus palmata* Forssk



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Received 6 September 2022; accepted 20 November 2022 Available online 24 November 2022

KEYWORDS

Ficus palmata Forssk; GCMS; Thrombolytic; Anticancer HepG2 cell line; Molecular docking; ADME **Abstract** *Ficus palmata* Forssk. (Moraceae family) is medicinally valuable plant that is mostly used as folk medicine for the treatment of different diseases. Phytochemical composition was evaluated by preliminary phytochemical investigation, GCMS analysis, and total bioactive contents (TPC and TFC). The antioxidant, enzyme inhibition, antimicrobial, thrombolytic and anticancer activities were performed for biological evaluation. The extract exhibited the maximum total phenolic (49.24 \pm 1.21 mg GAE/g) and total flavonoid contents (29.9 \pm 1.13 mg QE/g) which may be correlated to higher antioxidant potential of extract. The GCMS investigation identified the presence of 27 phytocompounds of different classes related to aldehydes, esters of fatty acids, triterpenes, steroids, triterpenoid. The extract possessed the strong α -glucosidase (73.4 \pm 4.65 %) and moderate α -amylase inhibition activity (47.1 \pm 3.29 %). Significant results were observed in case of antiviral, antifungal, and antibacterial activities. *F. palmata* extract inhibited the growth of HepG2 cancer cells in a dose-dependent manner. The extract also exhibited moderate *in vitro* thrombolytic activity. In addition, the phytocompounds identified by GCMS were subjected to *in silico* molecular docking studies to analyze the binding affinity between phytocompounds and

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Peer review under responsibility of King Saud University.



https://doi.org/10.1016/j.arabjc.2022.104455

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enzymes (α -glucosidase and α -amylase). Moreover, the best docked compounds were selected for ADMET studies which provide information about pharmacokinetics, physicochemical properties, drug-likeness, and toxicity of identified phytocompounds. The outcome of our research revealed that ethanolic extract of *F. palmata* possessed good antidiabetic, antimicrobial, thrombolytic and anticancer potential. This plant should be further explored to isolate the bioactive compounds for new drug development.

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1. Introduction

Ethnopharmacology and health sciences have observed a fast growing pharmaceutical and pharmacological studies towards the use of medicinal flora for potential therapeutic purposes (Ramalingum and Mahomoodally 2014). For many years, plant species have been medicinally utilized for therapeutic reasons (Hussain and Kumaresan 2014).

Despite the increasing prevalence of synthetic and semisynthetic derivatives for the treatment of infectious disorders, the number of resistant bacterial strains towards antibiotic drugs has been increasing day by day (Saklani and Chandra 2011). It continues to pose a threat to both underdeveloped and industrialized countries. Gram-positive bacteria produce resistance against antibacterial agents, has been a major international problem. A few effective antimicrobial agents are available for bacterial infections caused by these bacteria (Patel et al., 2012). Staphylococci that are multidrug-resistant (MDR) are becoming a bigger threat for the human physical and mental health. The resistant pathogenic Staphylococcus aureus strain, notably methicillin-resistant Staphylococcus aureus (MRSA), has posed severe difficulty in the treatment and control of Staphylococcal infections (Kaur and Chate 2015). Methicillin-resistant Staphylococci (MRS) have induced illnesses that are difficult to treat. MRSA has arisen simultaneous resistance to routinely used antibiotics, including macrolides, fluoroquinolones, tetracycline, aminoglycosides and chloramphenicol (Clinical and Institute 2009). In recent literature, plant polyphenols have been proved to exert antibacterial activity by inducing morphological and physiological damage to the bacterial cell membrane. Different studies have revealed that polyphenol rich plants possess many activities chiefly antioxidant and antibacterial activities (Takó et al., 2020). Additionally, plants polyphenols were reported to work by different mechanisms against viruses, including preventing virus entrance and affecting virus replication (Montenegro-Landívar et al., 2021). Due to these reasons, plants and their metabolites should be studied thoroughly for their potentials to treat and prevent bacterial and viral infections, and this was one of the aims of our current study.

Diabetes is a metabolic disease caused by decrease in production or impaired action of insulin (Chigurupati et al., 2022). Inhibiting the diabetic enzymes α -glucosidase and α -amylase is the first step for reducing postprandial hyperglycemia (Anjum and Tripathi 2019). Both of these diabetic enzymes can also be inhibited by antioxidants which help in reducing hyperglycemic conditions and these antioxidants include *N*acetylcysteine, vitamin C and α -lipoic acid and some others (Bajaj and Khan 2012, Alqahtani et al., 2019). In this context, we selected a plant that is traditionally used in Saudi Arabia to treat diabetes with hope of supporting global efforts to battle diabetes and its complications.

Ficus palmata belongs to family Moraceae which comprises very high tree, shrubs and rarely herbs (Galil 2019). The plant of this family are traditionally used in asthma, scabies, gonorrhea, diabetes, diarrhea, and as antiseptic (Khan et al., 2022). The traditional use of this genus include in variety of diseases such as diabetes, ulcer, fungal infections, tumor, anti-inflammatory, hepatoprotective, nephroprotective and anticoagulant activities (Kitajima et al., 1999). *F. palmata* is distributed in some countries of South Asia, Somalia, South Egypt, Iran, Sudan, Ethiopia and Saudi Arabia (Khan et al., 2011). *Ficus palmata* is used for the treatment of various diseases such as tumor, diabetes,

ulcer, gastrointestinal and fungal diseases (Sati et al., 2020). In numerous studies the important pharmacological activities of different parts of *F. palmata* have been reported; *F. palmata* latex have antibacterial activity while stem bark of this plant possessed antioxidant, cardioprotective and antimicrobial activity in addition to several *in vivo* therapeutic effects like hepatoprotective, nephroprotective, antiulcer and anticoagulant activities (Khajuria et al., 2018). In literature, phytochemical investigation of *F. palmata* revealed the presence of terpenes, coumarins, sterols, furanocoumarin glycosides, isoflavones, lignans, chromone, triterpene, bergapten, vanillic acid, psoralen and flavone glycosides (Alqasoumi et al., 2014).

The present research aims to evaluate numerous *in vitro* biological activities and *in silico* characteristics of ethanolic extract of *F. palmata*. Phytochemical tests were performed by qualitative phytochemical evaluation, determination of total phenolic and flavonoid contents and GCMS analysis. Additionally, *in vitro* biological assays including antioxidant (DPPH, ABTS, FRAP, CUPRAC), antibacterial (Gramnegative and Gram-positive bacterial strains), antiviral (H9, IBV and NDV strains), cytotoxicity and enzyme inhibition (α -glucosidase and α -amylase) activities were performed. The phytocompounds identified by GCMS in ethanolic extract of *F. palmata* were studied for *in silico* studies.

2. Materials and methods

2.1. Collection and Extraction of F. palmata aerial parts

Plant was collected from Jazan province in September 2020, Fayfa Mountains, Saudi Arabia. Plant was identified by botanist, King Saud University, Riyadh, Saudi Arabia with voucher number (24558), submitted specimen in college of science, king Saud University. The collected aerial part of plant (1 kg) was dried in shade and grounded to fine powder. The grounded plant powder was soaked in 80 % EtOH (3 L) (Javed et al., 2020). The aqueous alcoholic solution have better properties for extraction of polyphenols (Ghalloo et al., 2022). The extract was concentrated using a rotary evaporator at 40 °C, to yield a dark brown residue of 98.5 g.

2.2. Phytochemical analysis of extract

2.2.1. Preliminary phytochemical investigation

The *F. palmata* extract was investigated for qualitative phytochemical analysis for the determination of metabolites (primary and secondary) like carbohydrates, glycosides, amino acids, starch, lipids, proteins, saponins, steroids, terpenoids, phenols, flavonoids and tannins according to methods described in literature (Velavan 2015).

2.2.2. Quantification of total bioactive contents

2.2.2.1. Determination of total phenolic content. The total phenolic content was determined by Folin–Ciocalteu method

adapted from literature with slight modification (Sembiring et al., 2018, Ghalloo et al., 2022). 0.5 mg/mL of sample was prepared; 0.1 mL of Folin-ciocalteu reagent was mixed with 0.1 mL of sample in a 96-well microplate. Sodium carbonates 10 % (2.8 mL) was prepared then added into the mixture and leave it at room temperature for 2 h.Absorbance was taken by BioTek synergy HT (Winooski, VT, USA) microplate reader at 765 nm in a 96-well microplate. As a standard, Gallic acid dilutions (0.05–0.5 mg/mL) used for calibration curve.

Total phenolic content =

Gallic acid equivalent (GAE) mg / g of plant extract

2.2.3. Determination of total flavonoid content

The total flavonoid contents was determined by using Aluminum calorimetric assay from the literature (Sembiring et al., 2018) by made some modifications. Standard solution with different concentrations 30, 40, 50, 60, 70, 80, 90, 100 µg/mL was prepared in ethanol. 50 µL of extract was diluted from 1 mg/mL and added in 10 µL of Aluminum chloride (10 %) solution then add 150 µL of ethanol added. Sodium acetate (1 M) 10 µL also added in the mixture in a 96-well microplate. The entire mixture was incubated for 30 min at room temperature. Ethanol was taken as blank and Quercetin was used as standard. Absorbance was taken by using a BioTek Synergy HT microplate reader at 415 nm.

Total flavonoid content

= quercetin equivalent mg/g of plant extract.

2.2.4. Gas chromatography-Mass spectrometry (GCMS) Analysis

The extract of *F. palmata* was analyzed by GCMS analysis with a specification of instrument was Agilent 7890B with Mass Hunter acquisition software. The GCMS (instrument) has a HP-5MS ultra inert capillary with non-polar column 30 m \times 0.25 mm ID \times 0.25 µm film, containing inert gas Helium which runs at a speed of 1 mL/min. Sample injector started at 250 °C, and oven temperature was adjusted in this manner that for 5 min at 50 °C, then progressively rise to 250 °C at 100 °C/minute and at the end, goes to 3000°C at 70 °C / minute for 10 min. The identification of phytocompounds was determined by evaluation with the NIST library data and the % peak area was calculated from chromatogram total peak area (Dilshad et al., 2022).

2.2.5. Phenolic acids quantification by HPLC

The sample extract was quantified by HPLC Shimadzu, Japan. Briefly, dried sample (0.1 g) was mixed in a 1 mL methanol, and 10 μ L volume was injected in the HPLC system for quantification of the phenolic profile. Shim-pack CLC-ODS C-18 column (5 cm \times 4.5 mm, 5 μ m) Shimadzu was used for sample analysis and a mixture of glacial acetic acid and distilled water was utilized as a mobile phase with the ratio of 56:44 v/v.

2.3. In vitro biological activity

2.3.1. Antioxidant assay

2.3.1.1. Dpph radical scavenging Activity. DPPH assay was used to determine the anti-oxidant activity of plant extract

was performed according to literature with slight modification (Ratshilivha et al., 2014). DPPH solution 0.1 mM was prepared then sample solution (plant extract) was also prepared in methanol at 5 mg/mL. Inoculated 90 μ L of DPPH solution and 10 μ L of sample solution in 96-well microplate. The reaction mixture was incubated at 37 °C for 30 min. Ascorbic acid was used as standard. The absorbance was taken at 517 nm with a microplate reader synergy HT Biotek, USA. The following formula was used for determination of Radical scavenging activity

Inhibition% =

 $100 - (absorbance of sample/absorbance of control) \times 100$

2.3.1.2. Frap assay. Ferric-reducing anti-oxidant power assay was performed according to literature with little modification (Akinrinde et al., 2018). 100 μ L of *F.palmata*(1 mg/mL) and 2 mL FRAP reagent was mixed and incubated for 30 min. Absorbance was taken at 593 nm by using BioTek Synergy HT reader.

2.3.1.3. Cuprac. Cupric reducing antioxidant capacity was determined by method adopted in literature with slight modification (Benarfa et al., 2019). Firstly, 100 μ L of *F. palmata* extract was added into 10 mM (200 μ L) CuCl2, 7.5 mM (200 μ L) neocuprine and 1 M (200 μ L) ammonium acetate buffer (pH 7.5) reaction mixture. After incubation for 30 min, absorbance was taken at 450 nm by using microplate reader.

2.3.1.4. ABTS. ABTS activity was carried out by the modified method in literature (Ghalloo et al., 2022) in which 100 μ L of sample solution was added with 200 μ L of ABTS solution. The mixture was incubated for 30 min at room temperature. The absorbance was measured at 417 nm using a BioTek Synergy HT microplate reader.

2.3.2. Antibacterial assay

Antibacterial activity was performed according to literature (Ahmad et al., 2019).

2.3.2.1. Test organism. Micrococcus luteus (ATCC 10240), Staphylococcus epidermidis (ATTC 12228), Bordetella bronchiseptica (ATCC 4617), Bacillus subtilis (ATCC 6633), Bacillus pumilus (ATCC 14884), Staphylococcus aureus (ATTC 6538), Escherichia coli (ATCC 10536). These 7 bacterial strains were procured from microbiology laboratory of the Islamia University of Bahawalpur.

2.3.2.2. Agar well diffusion method. Fresh Inoculum was prepared from fresh bacterial cultures by picking the little amount of colonies and inoculates in sterile Muller Hilton nutrient broth and places it in incubator for 24 h at 37 °C for bacterial growth. Standardize inoculated bacterial nutrient broth with 0.5Macforland by taken absorbance in UV. Agar media was poured in petri plates and left it to solidify. Wells were formed in each petri plates respective to concentrations of plant extract used after diluting it in DMSO (1 mL for each) 25 mg/mL, 50 mg/mL, 75 mg/ mL, 100 mg/mL. Petri plated were Incubate for18- 24 h at 37 °C, After 24 h, zone of inhibition was measured. Ceftriaxone (250 mg) was used as standard. Dilution of standard was made 1 mg/mL in sterilized distilled water. The zone of Inhibition of standard and sample were noted to determine the antibacterial activity of plant.

2.3.3. Enzymes inhibition assay (α -glucosidase and α -amylase) 2.3.3.1. α -Glucosidase inhibitory activity. α -Glucosidase inhibition activity was performed according to literature (Rehman et al., 2018) with minor modification by changing the absorbance at 400 nm. Phosphate buffer solution 0.1 M of pH 6.8 was prepared. Ficus palmata extract (sample) 1 mg was diluted in 1 mL of methanol. Added 10 µL of sample, 70 µL of phosphate buffer solution, 10 µL of α -glucosidase in each well of 96-well microplate. The plate was incubated for 15 min at 30 °C. Added 10 µL of substrate ρ -Nitrophenyl- α -Dglucopyranoside and incubate for 30 min further. Absorbance was measured in %inhibition through the HT BioTek microplate reader. Methanol set as control and acarbose (100 µg/ mL) as a positive control. The enzyme inhibitory activity of sample was measured as

%inhibition =

 $100 - (absorbance of sample/absorbance of negative control) \times 100$

2.3.3.2. α -Amylase inhibitory activity. α -Amylase inhibitory activity of Ficus palmate was performed according to literature (Anyanwu et al., 2019) with slight modification by changing the incubation time. Assay was performed in 96-well microplate. Sodium phosphate buffer (contained 0.6 M NaCl) was prepared with maintained pH 6.9. The volume of 25 µL Sodium phosphate buffer, 20 µL of soluble starch (1 %W/V 20 mg/2mL), 20 µL of plant extract (1 mg/mL), 20 µL of Acarbose was added in each well of 96-well microplate and incubate at 37 °C for 7 min. After incubation, 15 µL of α-amylase (12 mg/2mL) was added in reaction mixture and incubate at 37 °C for 17 min for further. The volume of 20 µL of HCl (1 M) and 100 µL of iodine reagent was added. The volume of 20 µL methanol and acabose (100 µg/mL) as negative and positive control in separate wells were used. Absorbance was measured at 620 nm by HT BioTek microplate reader.

%*inhibition* = (*absorbanceofcontrol*)

aborbanceofsample/absorbanceofcontrol)
 * 100

2.3.4. Evaluation of antiviral activity

2.3.4.1. Viral strains. Three viral strains that was Infectious bronchitis virus, Newcastle disease virus, Avian Influenza A.

2.3.4.2. Inoculation of viruses. The protocol followed from literature (Shahzad et al., 2020). Eggs were brought from the government poultry farm, Model town A Bahawalpur. Eggs were free of pathogens. Chick embryonated eggs were taken which was 9–11 days old, were injected with viruses in chorioallantoic fluid. Before and after inoculations, eggs were candled. The wider side of eggs was swabbed with 70 % ethanol and moves it in tray place in Biosafety cabinet II. The viral inoculum injected through the 3 cc sterile needle along the wider side of eggs. The viral inoculum and drug was in same quantity. Molten wax was used to seal the hole after it, eggs were incubated at 37 °C. After 72 h of inoculation, allantoic

fluids were obtained from the harvested eggs and check for haemagglutination test (HA). The changes in HA titer was noted as the virus particles in sample increases the HA titer quantity increase. For negative control, Normal saline was used; DMSO used as solvent control, virus without drug as virus control. The standard used was acyclovir (Musaddiq et al., 2020).

2.3.4.3. Hemagglutination Test (HA). Test usually performs for quantification of titer of viruses except IBDV(Elizondo-Gonzalez et al., 2012) All steps followed for HA test was according to world organization of animal health manual of diagnostic test and vaccine for terrestrial animal (Stear 2005) 96-well flat bottom microplate was used. 50 μ L of 7.4 pH phosphate buffer solution was added in each well. Same quantity of allantoic fluid(from infected eggs) was added in first wells and then mix it and add 50 μ L from well 1 to well 2 then serially diluted it and mix it till 11 th well. Well 12th remain as negative control. 50 μ L of chicken RBC solution (1 %) put in each well. Afterthat the plate set at 37 °C for 1 h and note the HA variations.

2.3.5. Antifungal assay

2.3.5.1. Fungal strains. Three fungal strains Aspergillus niger, Fusarium avenaceum, Fusarium brachygibbosum were used to identify antifungal activity of extract.

2.3.5.2. Agar tube dilution method. Antifungal activity was conducted through agar tube dilution method according to method in literature with slight modification (Andleeb et al., 2020). Agar, dose of extract and fungi were inoculated in tubes and then incubated at 37 °C for 24 h. Terbinafine as standard and DMSO as control (negative) was used in this assay.

% Growth Inhibition = Growth in control / Growth in test *100

2.3.6. 2.3.6. Evaluation of thrombolytic assay

The thrombolytic assay of extract was determined according to the method reported in literature(Ahmed et al., 2022). The blood sample was taken out without affecting the blood clot in eppendorf tube and incubated at 37 °C for 45 min. After incubation, serum was carefully removed from the blood sample without affecting the blood clot then again weighed the tube. 1 mg/mL of plant extract was added to each preweight tube containing blood clot which was then incubated for 90 min at 37 °C. The Eppendorf tubes thrombolytic activity was observed after 90 min. The Eppendorf tubes released fluids which was taken out and weight again. The weight difference between eppendorf tubes was observed which demonstrated the extract antithrombotic activity. Streptokinase (standard) and distilled water (negative control) was used.

$$\% \ clot \ lysis = (Wr / Wc) * 100$$

Where; Wr = released clot weight. Wc = clot weight.

2.3.7. Determination of Cytotoxic activity of extract

Cytotoxicity of extract was determined by MTT assay on HepG2 liver cancer cells which was performed according to literature with slight modification in the absorbance at 490 nm (Khan et al., 2013). Dulbecco's Modified Eagle's Medium (DMEM) was used as medium for cancer cell growth. Penicillin and streptomycin set as standard. The Dulbecco's Modified Eagle's Medium (DMEM) was added in a 75 cm2 flask along with 10 % foetal bovine serum (FBS), 100 IU/mL penicillin, and 100 μ g/mL streptomycin. The cells were then kept in an incubator at 37 °C with 5 % CO2. Extracts diluted in DMSO (0.05 % concentration) were used to treat cells.

2.3.7.1. Determination of cell viability. HepG2 cell line was treated with different concentration of sample extract 20,40,60 and 100 μ g/mL for 48 h. After treatment, a 10 μ L MTT reagent was added to each well and incubated for an additional 4 h. After that, 150 μ L of DMSO was added to dissolve the formazan crystals, and the absorbance was measured at 490 nm. The percentage of cell viability was determined by the following equation;

$$Cell viability (\%) = (A sample - A blank / A control - A blank) * 100$$

Where;

A $_{sample}$ = absorbance of sample.

A $_{blank}$ = absorbance of blank.

A $_{\rm control}$ = absorbance of control.

A $_{blank}$ = absorbance of blank.

2.4. In Silico activities

2.4.1. Molecular docking studies

The binding affinities between the phytocompounds (by GCMS) with enzymes 5zcb (α -glucosidase) and 4w93 (α amylase) were assessed by using PyRx software. The 3D shapes of proteins were obtained from the RCSB Protein Data Bank (https://www.rcsb.org/) (Majid et al., 2022). First, the proteins were prepared using the Discovery studio by removing water and ligands, inserting polar hydrogen atoms, and saving them in PDB format. Then download the ligand in SDF format through PubChem (https://pubchem.ncbi.nlm. nih.gov/). The Open babel software converted the ligands to PDB files. The prepared ligands were docked blindly in the protein's grid box to allow them to find any suitable binding location (Khursheed et al., 2022). The molecular docking was confirmed by redocking α -glucosidase, α -amylase, and chosen ligands with Autodock 1.5.6. Furthermore, the binding affinity values yielded the same results. The Ligplot software was used for visualization of 2D structures of ligand-protein interactions (Aati et al., 2022).

2.4.2. ADMET studies

The best docked compounds were analyzed for ADMET studies by using the online Swiss ADME (<u>https://www.swissadme.ch/</u>) tool (Majid et al., 2022). The toxicity of docked compounds (best docked) were assessed through the online software ProTox-II (<u>https://tox-new.charite.de/</u>) (Banerjee et al., 2018).

2.5. Statistical analysis

The findings were all presented as mean standard deviation (mean \pm SD). The information obtained through quantitative analysis IBMSPSS (v20, Chicago, IL, USA) was used to per-

form ANOVA (one-way analysis of variance) followed by post-hoc test. Significant values were defined as p values less than 0.05.

3. Results

3.1. Phytochemical profiling of extract

3.1.1. Preliminary phytochemical investigation

Phytochemical tests of *Ficus palmata* ethanolic extract revealed the presence of carbohydrates, lipids, saponins and tannins as well as phenols, flavonoid, steroids/terpenes, alkaloids and cardiac glycoside while the resins, proteins and amino acids were absent in extract of *F. palmata* as confirmed by the various tests and presented in Table 1.

3.1.2. Total bioactive contents

The results of total phenolic and flavonoid contents are shown in Table 2. The total phenolic content in the extract of *Ficus palmata* was 49.24 mg GAE/g and the total flavonoid content was 29.9 mg QE/g.

3.1.3. Determination of phytochemicals by GCMS

GCMS analysis was conducted to determine the major and minor compounds in extract of Ficus palmata ethanolic extract showed various peaks in GCMS chromatogram as shown in Fig. 1 which were determined on fused silica capillary column. The total 27 tentative compounds were identified and the major compounds were recognized by their peak % Area which was determined by the total area of peaks. Major compounds of GCMS are 12-Oleanen-3-yl acetate (5.28 %), 11, 14, 17-eicosatrienoic acid methyl ester (1.99 %), Bergapten (1.82 %), Acetic acid, 4,4,6a,6b,8a,11,11,14b-octamethyl-13o xodocosahydropicen-3-yl ester (1.77)%), 9.12-Octadecadienoic acid (1.63 %), n-Hexadecenoic acid (1.57 %), Taraxasterol (1.56 %), Stigmasterol, 22,23dihydro-(1.30 %), Phytol (1.03 %), 3-Hydroxy-4methoxybenzoic acid (1.02 %), 3.5-Dihydroxy-6-methyl-2.3-d ihydro-4H-pyran-4-one (0.98 %), and 2,7-Naphthalenediol (0.89 %) in addition to many other minor compounds. Most

Table 1 Phytochemical analysis of the ethanolic extract of *F. palmate.*

Sr. No.	Metabolites	Tests	Results
1.	Carbohydrates	Molisch's Test	+
		Iodine Test	+
2.	Glycoside	Erdmann's Test	+
3.	Phenols	Ferric chloride test	+
4.	Proteins	Biurette Test	_
5.	Alkaloids	Hager's Test	+
6.	Tannins	Lead Acetate Test	+
7.	Lipids	Saponification Test	+
8.	Amino acids	Ninhydrin Test	-
9.	Saponins	Froth Test	+
10.	Flavonoids	Reaction with NaOH	+
11.	Resins	Acetic anhydride	-
12.	Steroids/	Salkowaski's test	+
	Terpenes		

"+" present; "-" absent.

Table 2 Results of total bioactive content and antioxidant activities of ethanolic extract of <i>F. palmata</i> .								
Total bioactive conte	ent	Antioxidant (mg Tl	Antioxidant (mg TE/g)					
TPC (mg GAE/g)	TFC (mg QE/g)	DPPH	FRAP	ABTS	CUPRAC			
49.24 ± 1.21	$29.9~\pm~1.13$	$75.62~\pm~2.31$	110.39 ± 4.53	$92.89~\pm~3.92$	$195.62~\pm~7.72$			
"TPC" Total phenolic content, and "TFC" Total flavonoid content.								



Fig. 1 GCMS chromatogram of ethanolic extract of *F. palmata*.

of the detected compounds belong to triterpenoid, fatty ester, aldehyde, organic acid, aromatic substances, diterpene alcohol, and sterols classes (Table 3).

3.1.4. Phenolic Acids Quantification by HPLC

In the HPLC quantification, six compounds; *p*-Coumeric acid, Kaempherol, Quercetin, Ferrulic Acid, Benzoic acid, and Rutin were quantified (Fig. 2 and Table 4).

3.2. In vitro biological activies

Ficus palmata extract was evaluated in vitro biologically for its antioxidant, enzyme-inhibiting, antimicrobial, anticancer, and thrombolytic potentials.

3.2.1. Antioxidant activities

Ficus palmata showed strong antioxidant results due to the presence of compounds such as flavonoids, diterpenes, triterpenes and phenolics (Karatoprak et al., 2020). These antioxidant activities are responsible for the antidiabetic potential of plant. Antioxidant activities were determined by DPPH, ABTS, FRAP and CUPRAC. Results are shown in Table 2. DPPH and ABTS scavenging activities of ethanolic extract of *F. palmata* were 75.62 mg TE/g, 92.89 mg TE/g respectively while the FRAP and CUPRAC results were 110.39 mg TE/g, 195.62 mg TE/g respectively.

3.2.2. In vitro enzyme inhibition assay

Antidiabetic activity was performed to confirm its traditional use in diabetes mellitus which our results also supported its use by indicating the strong antidiabetic potential against diabetic enzymes. The extract possessed α -glucosidase inhibitory activity of 73.4 % at a concentration of 1 mg/mL as shown in Fig. 3. The inhibitory effect of *F. palmata* extract against α -amylase was 47.1 % at 1 mg/mL (Fig. 3). The inhibitory effect of *F. palmata* against α -glucosidase enzyme was much higher than against α -amylase while the acarbose showed 89.91 % inhibition against α -glucosidase and 77.41 % against α -amylase. Acarbose was used as a positive control while methanol was used as a negative control.

3.2.3. Antibacterial assay of ethanolic extract of F. palmata

Antibacterial activity was performed to evaluate the potential of *F.palmata* extract against seven bacterial strains and our report confirmed that the *F. palmata* extract was active against many bacterial strains and could be employed to reduce different bacterial infections. The *F. palmata* possess many secondary metabolites like triterpenes, alkaloids, sterols, polyphenols, flavonoids, and saponins, which showed their antibacterial potential and make the plant significant for therapeutic purposes i.e. dysentery, diarrhea, skin diseases, lung complications, ulcers, cough, heart disease (Santos et al., 2013). The extract showed highest zone of inhibition (23 mm) against *Staphylococcus aur*

Retention time (minutes)	Peak area (%)	Phytochemical compounds identified	Molecular formula	Molecular weight (g/mol)	Chemical class
6.19	0.98	3,5-Dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4- one	$C_6H_8O_4$	144.1	Organic acid
8.00	0.25	Hydroquinone	$C_6H_6O_2$	110.1	Aromatic organic
8.11	0.58	2-Methoxy-4-vinylphenol	$C_9H_{10}O$	150.1	Phenolic compound
10.23	0.89	2,7-Naphthalenediol	$C_{10}H_8O_2$	160.1	Aromatic substance
11.52	1.02	3-Hydroxy-4-methoxybenzoic acid	$C_8H_8O_4$	168.1	phenolic acid
11.64	0.03	4-Hydroxy-3-methylbenzoic acid	$C_8H_8O_3$	152.1	Monohydroxybenzoic acid
15.06	0.12	Pentadecanoic acid, 14-methyl-, methyl ester	$C_{17}H_{34}O_2$	270.5	Fatty acid methyl ester
15.62	1.57	n-Hexadecenoic acid	$C_{16}H_{30}O_2$	254.4	Palmitic acid
15.91	0.38	Hexadecenoic acid, ethyl ester	$C_{18}H_{36}O_2$	284.5	Palmitic acid ester
17.12	1.82	Bergapten	$C_{12}H_8O_4$	216.1	Furocoumarin
17.35	0.25	9,12,15-Octadecatrienoic acid, methyl ester	$C_{19}H_{32}O_2$	292.5	Fatty acid
17.52	1.03	Phytol	$C_{20}H_{40}O$	296.5	Diterpene alcohol
17.91	1.63	9,12-Octadecadienoic acid	$C_{18}H_{32}O_2$	280.4	Fatty acid
18.03	1.99	11, 14, 17-Eicosatrienoic acid methyl ester	$C_{21}H_{36}O_2$	320.5	Fatty acid methyl ester
18.19	0.29	9, 12-Octadecadienoic acid ethyl ester	$C_{20}H_{36}O_2$	308.5	Fatty acid ester
18.22	0.14	Octadecanoic acid	$C_{18}H_{36}O_2$	284.5	Fatty acid
23.03	0.09	Urs-12-en-24-oic acid, 3-oxo-, methyl ester	$C_{31}H_{48}O_3$	468.7	Triterpenes
23.44	0.26	β-Amyrin	C ₃₀ H ₅₀ O	426.7	Triterpenes
25.26	0.14	Lupeol	C ₃₀ H ₅₀ O	426.7	Triterpenoid
25.83	0.64	cis, cis, cis-7, 10, 13-Hexadecatrienal	$C_{16}H_{26}O$	234.38	Fatty aldehyde
26.38	5.28	12-Oleanen-3-yl acetate, (3.α.)	$C_{32}H_{52}O_2$	468.8	Triterpenoid
31.00	0.19	Brucine	$C_{23}H_{26}N_2O_4$	394.5	Monoterpenoid indole alkaloid
34.51	0.12	Stigmasterol	$C_{29}H_{48}O$	412.7	Sterol
35.66	1.77	Acetic acid, 4,4,6a,6b,8a,11,11,14b-octamethyl-13- oxodocosahydropicen-3-yl ester	$C_{32}H_{52}O_3$	484.8	Ester of acetic acid/ picene
35.91	1.30	Stigmasterol, 22,23-dihydro-	C ₂₉ H ₅₀ O	414.7	Steroid
36.63	0.23	α-Amyrin	C ₃₀ H ₅₀ O	426.7	Triterpenoid
37.89	1.56	Taraxasterol	C ₃₀ H ₅₀ O	426.7	Triterpenoid

 Table 3
 Identification of metabolites in Ficus palmata by GCMS.



Fig. 2 HPLC chromatogram in *Ficus palmata*.

Table 4 Compounds quartered	ntified through HPLC.			
Compound Name	RT (Min)	Area	k-factor	Conc. (µg/g)
p-Coumeric acid	3.214	1,673,213.2	0.0003	207.47
Kaempherol	11.974	2,347,633.2	0.0000408	95.78
Quercetin	15.814	103,407.0	0.000696	71.97
Ferrulic acid	12.666	394,200.0	0.0000767	30.26
Benzoic acid	18.394	77,093.7	0.0000345	2.66
Rutin	24.839	332,476.7	0.000113	37.57



Fig. 3 α -Glucosidase and α -Amylase % inhibition assay.

 Table 5
 Antibacterial activity of extract of F. palmata against seven bacterial strains.

Bacterial strains	Zone of inhibition (mm)							
	Conc. 25 mg/mL	Conc. 50 mg/mL	Conc. 75 mg/mL	Conc. 100 mg/mL	Standard Ceftriaxone 1 mg/mL			
Staphylococcus aureus	17	20	21	23	26			
Staphylococcus epidermidis	14	16	18	21	26			
Escherichia coli	10	12	15	22	25			
Bordetella bronchiseptica	12	13	14	15	20			
Bacillus subtilis	15	16	17	18	22			
Bacillus pumilus	12	13	14	16	21			
Micrococcus luteus	11	13	15	17	21			

eus at a concentration of 100 mg/mL and showed minimum activity (15 mm) at concentration 100 mg/mL against *Bordetella bronchiseptica*. The extract showed least activity at concentration of 25 mg against few bacteria but little more for some other bacteria. The results revealed the direct relation between zone of inhibition (mm) and different concentrations of *F. palmata* extract. The detailed results of antibacterial activity are shown in Table 5 and Fig. S1.

3.2.4. Antiviral activity

The purpose to perform this experiment was that the extract possess different phytochemicals like flavonoids, polyphenolics, terpenoids, lignans, coumarins, sulphides, alkaloids, saponins, furyl compounds, as determined by GCMS, which have antiviral activity and our results confirm the antiviral potential, antiviral activity was performed (Bouazzi et al., 2018). Purpose of this study, Synthetic antiviral agents was used for the treatment of various viruses have reported adverse effects so less toxic agents are required, so the antiviral assay was performed to evaluate its activity against some specific viral strains (Fernandez et al., 1998). The results demonstrated a significant antiviral activity of *F. palmata* ethanolic extract of aerial parts against IBV, NDV and H9 with respective titer count 04, 04 and 00. Ethanol extract showed highly strong inhibitory effect against H9 virus and strong activity against IBV and NDV as depicted in Table 6.

Table 6	Antiviral values	s of ethanol extract	xtract of F. palmata.		
Strains	Titer count in control	Titer count with Acyclovir	Titer count with <i>F. palmata</i>		

IBV	1024	00	04	
NDV	2048	00	04	
H9	2048	00	00	

"IBV" Avian Infectious Bronchitis Virus, "H9" Influenza virus, and "NDV" Newcastle disease virus,

HA titer 0–8; highly strong, 16–32; strong, 64–128; moderate, and 256–2048; not active (Musaddiq et al., 2020).

Thrombolytic Activity



Fig. 4 Thrombolytic activity of ethanol extract of *F. palmata.at* 1 mg/mL.

3.2.5. In vitro thrombolytic assay

This activity was intended to be performed due to its excellent medicinal relevance and result which demonstrated its great thrombolytic potential and potential for use as an antiplatelet or thrombolytic agent. The thrombolytic potential of the plant *F. palmata* extract was evaluated, which are shown in Fig. 4. The percentage of clot lysis by ethanolic extract of *F. palmata* was 37.17 % at 1 mg/mL while Streptokinase showed 76.98 % clot lysis.

3.2.6. Antifungal activity

Ficus palmata traditionally used as antifungal agent and to confirm its antifungal potential, the antifungal activity was performed. Our results also confirmed the antifungal potential of plant because it possesses antimicrobial compounds likely 3,5-Dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4-one (Igwe et al., 2016), Octadecanoic acid (Jasim et al., 2015), and Urs-12-en-24-oic acid, 3-oxo-, methyl ester (Chenniappan et al., 2020). Table 7 represented the antifungal activity of *F. palmata*



Fig. 5 Cytotoxic effect of *F. palmata* ethanolic extract on HepG2 cell line.

extract. The results of this assay showed the maximum antifungal activity of extract against fungal strains *Fusarium avenaceum* (58.6 %) followed by *Fusarium brachygibbosum* (45.45 %) and *Aspergillus niger* (41.66 %).

3.2.7. Cytotoxic effect of F. Palmata extract

Because of its folkoric use, this activity was performed to confirm its potential as anticancer agent and great results of anticancer activity also justify its anticancer/antitumor phytocompounds in this plant. Results of cytotoxicity of *Ficus palmata* extract against HepG2 cell line are shown in Fig. 5. The plant extract showed significant variations in % cell viability at different concentrations. At 100 µg/mL, extract showed 17.46 % viability on a cell line HepG2. However, at 20 µg/ mL concentration of extract showed 57.67 % viability.

3.3. In silico studies

3.3.1. Molecular docking studies of phytocompounds against enzymes

Molecular docking of compounds identified by the GCMS was performed against both α -glucosidase and α -amylase. Binding affinity of 9 compounds with α -glucosidase and 8 compounds with α -amylase were greater than the binding affinity of acarbose (standard). Maximum binding affinity of phytocompound with α -glucosidase was by urs-12-en-24-oic acid, 3oxo-, methyl ester with a value of -9.8 Kcal/mol by making no hydrogen bond and hydrophobic interactions Leu287, Gly286, Asn258, Met229, Phe225, Pro223, Val222, Leu219, and binding affinity of taxasterol with α -amylase with a value -10.8 Kcal/mol was maximum by making hydrogen bond Glu233, Asp197 (Table 8, Fig. 6, Fig. 7, S2, S3, S4). The

Table 7 Antifungal activity of F. palmata extract by agar tube dilution method.

Sample	strains	growth test (mm)	Growth control (mm)	% inhibition
F. palmate	A. niger	40	96	41.66 ± 4.29
	F. avenaceum	51	87	$58.6~\pm~5.67$
	F. brachygibbosum	45	99	$45.45~\pm~5.23$

Sr.	Ligand	Structures	α-glucosidas	se		α-amylase		
no			Binding energy (Kcal/mol)	H-bond interactions	Hydro-phobic interactions	Binding energy (Kcal/mol)	H-bond interactions	Hydro- phobic interactions
1.	urs-12-en-24-oic acid, 3- oxo-, methyl ester		-9.8	-	Leu287,Gly286, Asn258,Met229, Phe225, Pro223, Val222, Leu219	-9.2	-	Gln63, Trp59, Trp58, Asp300, Ile235, Glu233, Asp197, Leu162, Tyr62, Thr163, Leu165
2.	β-Amyrin	HO	-8.4	-	Leu287, Met229, Glu226, Phe225, Pro223, Val222	-10.1	-	Asp300, His299, Glu233,, Thr163, Tyr62, Leu162, Asp197, Trp58, Trp59
3.	acetic acid, 4,4,6a,6b,8a,11,11,14b- octamethyl-13- oxodocosahydropicen-3-yl ester		-8.3	Asn371, Lys334	Glu408, Asn333, Ile329, Leu301, Glu300, Leu296, His284	_	-	-
4.	α-Amyrin	HO	-8.1	Asn275, Thr253	Asn277, Phe276, Gly274, Glu271, Ile251, Ala247, Phe246, Lys242, Lys7, Trp6, Lys4	-10.2	Asp197, Glu233	His305, Asp300, His299, Thr163, Leu162, Tyr62, Trp59, Trp58
5.	taraxasterol	HO	-7.8	Glu408	Phe335, Lys334, Ile304, Leu301, Asp297, Leu296, His284	-10.8	Glu233, Asp197	Asp356, Asp300, His299, Thr163, Leu162, Tyr62,, Trp59, Trp58,
6.	stigmasterol	HOCHEN	-7.6	-	Lys290, Trp288, Leu287, Gly286, Gly259, Asn258, Phe225, Pro223, Ile143, Glu141,	-9	Ile235, Lys200	His201, Asp197, Tyr62, Leu165, Gln63, Thr163, Trp59, Leu162, Asp300, Tyr151

Table 8 Binding affinity and interactions of best docked compounds of enzymes.

10

	Table 8	(continued)
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Sr.	Ligand	Structures	α-glucosidas	se		α-amylase		
10			Binding energy (Kcal/mol)	H-bond interactions	Hydro-phobic interactions	Binding energy (Kcal/mol)	H-bond interactions	Hydro- phobic interactions
7.	lupeol	HO	-7.5	Phe282, Asn258,	Gly286, Gly259, Met229, Glu226, Phe225, Pro223	-10	Asp197	Asp356, His305, Asp300, Thr163, Leu162, Tyr62, Trp59, Trp58,
8.	brucine		-7.5	-	Asn316, Asn277, Glu271, Val269, Tyr249, Ala247, Phe246, Lys242, Lys7, Trp6	-9.2	-	Tyr151, Ile235, Leu162, Trp58, Trp59, Tyr62, Asp300, Asp197, His201, Glu233
9.	stigmasterol, 22,23-dihydro-	-7.1	-7.1	Lys205	Phe210, Gly209, Ala208, Glu173, Asn171, Glu157, Glu130, His129, Trp128, Ile127, Lys118	-9.3	Glu240	Ile235, Tyr151, Trp59, His299, Leu162, Arg195, Asp197, Ty62, Trp58, Glu233, Ala198, His201, Lys200
10.	Acarbose		-6.9	Asn316, Asn277, Phe276, Glu271, Ala270, Thr253, Tyr249, Lys248, Phe246, Lys7, Trp6	Asn275, Val269, Ile251, Asp250, Lys242,	-7.7	Gly403, Arg421, Arg398, Gly334, Thr11, Ser3, Thr6, Arg252, Gln404	Pro405, Asp402, Gln8, Phe335, Pro332



Fig. 6 2D structures of best docked compounds with illustrated interactions (hydrogen and hydrophobic interactions) to α -glucosidase. Red color (interactions) represented the hydrophobic interactions and Green dotted line showed the hydrophilic interactions.



Fig. 7 2D structures of best docked compounds with illustrated interactions (hydrogen and hydrophobic interactions) to α-amylase. Red color (interactions) represented the hydrophobic interactions and Green (dotted) line showed the hydrophilic interactions.

molecular docking for the compounds quantified by the HPLC was also performed. Amongst these compounds, Rutin showed best binding affinity with binding affinity -9.0 Kcal/mol against α -amylase and -8.9 Kcal/mol again

st α -glucosidase (Figs. S5 and S6, and Table S1). This revealed the potential of *F. palmata* extract to inhibit diabetic α -amylase and α -glucosidase.

3.3.2. ADMET analysis

The compounds (best docked) were further investigated using online software SwissADME which provide more knowledge about pharmacokinetic, physiochemical behavior and druglikeness. In case of Lipinski's rule of five, nearly all compounds violate the lipinski's rule except brucine while stigmasterol violate the lipophilicity rule and acetic acid, 4,4,6a,6b,8a,11,11,14b-octamethyl-13-oxodocosahydropicen-3 -yl ester violate the molar refractory rule. When any drug fails to meet two or more criteria of Lipinski's rule that drug considered as non-orally drug. All compounds violate one rule except brucine (follows all rules) indicated that these compounds are bioavailable or orally available. All the best docked compounds possessed orally active likeness characteristics. Due to the many benefits it offers, the oral administration route is favoured over the many other medication delivery methods. These benefits include dependability to accommodate a range of medications, safety, excellent patient compliance, convenience of swallowing, pain avoidance, and simplicity of use. Table S2 and S3 represents various characteristics of best docked compounds including lipophilicity, pharmacokinetic behavior, molecular weight, no. of rotatable bond, no. of hydrogen bond acceptor and donor, Lipinski rule and other properties while Fig. 8 presented the bioavailability radar of compounds (best docked). Table S4 represented the compounds with predicted toxicity and LD_{50} performed by software ProTOX-II. In this, one compound predicted as carcinogenic and immunotoxic agent as urs-12-en-24-oic acid, 3oxo-, methyl ester while other compounds showed low toxicity level.

4. Discussion

Phytocompounds investigation of extract of F. palmata exhibited phytocompounds such as carbohydrate, amino acids, lipids, tannins, phenols, flavonoid, saponins, steroid/ terpenes and alkaloids. Our results were in agreement with previous studies which have reported the presence of phytochemicals including carbohydrates, alkaloids, flavonoids saponins, tannins, resins and phenolic compounds in F. palmata extract while protein and amino acids were absent in all extract (Saklani and Chandra 2012). Most of these phytochemicals possessed pharmacological properties; for example alkaloids were found to possess analgesic and antimicrobial activities; flavonoids act as antioxidant and antibacterial while saponins possessed antibacterial, anticancer, and antidiabetic activities (Ghalloo et al., 2022). Thus, the presence of these phytochemicals in extract of F. palmata makes it likely to possess promising various therapeutical activities.

In literature, total bioactive components of the *F. palmata* plant parts were previously reported (Abbasi et al., 2015, Tewari et al., 2021). Our study revealed that the ethanolic

extract of *F. palmata* possessed total phenolic content (49.24 \pm 1.21 mg GAE/g extract) and total flavonoid content (29. 9 \pm 1.13 mg QE/g extract). The presence of high phenolic contents may increase its antioxidant potential (Balamurugan et al., 2020). From the vast literature review of whole plant of *F. palmata* which confirmed that there was no comprehensive data available about total bioactive components.

Antioxidants play a key role in the defense mechanism of organism against various pathological diseases (Litescu et al., 2011). Antioxidant activities of plants are attributed mainly to phenolic compounds which are necessary plant secondary metabolites. Basic mechanism of antioxidants is to neutralize reactive oxygen species and prevent conversion to free radicals. In our study, the plant *F. palmata* showed a significant antioxidant activities due to the presence of some polyphenolic acids which have antioxidant activities. These compounds are *p*-Coumeric acid (Boz 2015), Ferulic acid (Graf 1992), and Quercetin (Azeem et al., 2022). These results support of previous data reported in literature (Tewari et al., 2021).

The antidiabetic effect of *F. palmata* was previously reported and was found to happen through the inhibition of α -amylase and α -glucosidase enzymes leading to reduction of blood glucose level (Tewari et al., 2021). Our study confirmed the antidiabetic property of *F. palmata* extract which was more effective against α -glucosidase as compared to α -amylase; this is also in support of previous literature (Singh et al., 2014, Anjum and Tripathi 2019, Sati et al., 2020). This activity could be explained by antioxidant effect of flavonoid and triterpenoids (Singh et al., 2014), and also may be due to the compounds Kaempherol (Imran et al., 2019), Quercetin (Bule et al., 2019), and Rutin (Mel et al., 2020).

In our study, determination of antimicrobial potential of F. palmata extract was conducted by agar tube dilution and agar well diffusion method for antifungal and antibacterial activity respectively, against three fungal strains and seven bacterial strains. The results revealed maximum inhibition against fungal strains and significant inhibition zone (>10) towards many bacterial strains used in assay. Phytochemicals identified tentatively by GCMS and HPLC quantified compounds namely Hydroquinone, 2-Methoxy-4-vinylphenol, Bergapten, 3,5-Dih vdroxy-6-methyl-2,3-dihydro-4H-pyran-4-one, Phytol, Octadecanoic acid, 9, 12-octadecadienoic acid ethyl ester, Urs-12-en-24-oic acid, 3-oxo-, methyl ester, Ferulic acid, Benzoic acid, and Quercetin (Formica and Regelson 1995) are likely to be responsible for this antimicrobial activity. Phytochemical compounds (phenolic compounds, alkaloids, terpenoids, carotenoids, and some sulfur-containing compounds) are known to inhibit pathogenic strains. Such phytochemical compounds could be involved in the discovery of new antibacterial medications to combat various multidrug-resistant bacterial infections (Barbieri et al., 2017). Previous studies revealed the antibacterial potential of F. palmata bark, leaf, root and fruit extracts which showed significant zone of inhibition (Saklani and Chandra 2011, Kothiyal and Saklani 2017).

Most of phytochemicals as flavonoid, polyphenols, saponins, coumarins, alkaloids, proteins and peptides have been reported to possessed antiviral activity against different viruses (Jassim and Naji 2003, Shahzad et al., 2019). There is no literature data found about the antiviral activity of *F. palmata* extract. Results of our study showed for the first time that



Stigmasitosterol 22,23-dihydro

Fig. 8 Bioavailability RADAR of best docked compounds.

ethanolic extract of *F. palmata* is strongly effective against viruses H9, NDV and IBV. These findings are promising and can be further investigated to develop new antiviral agent.

Thrombosis or blood clot is formed by the deposition of fibrin, tissue factors and platelets at the site of injury in endothelial cell surface. Platelets is a key factor in the formation of clot, as the thrombosis process begins when active platelets establish platelet-to-platelet linkages which leads in the development of plaque (Bhowmick et al., 2014). Streptokinase is a major thrombolytic agent with severe side effects that emphasize the researcher in the discovery of an alternative antithrombotic agent. Several studies reported that the tannins, saponins and alkaloids are involved in clot lysis and these phytochemicals are responsible for disrupting the fibrin and fibrinogen which leads to fibrinolysis (Uddin et al., 2020). In this present study, compounds detected by GCMS and quantified by LCMS; mainly 11, 14, 17-Eicosatrienoic acid methyl ester (Hansch and Von Kaulla 1970), α-amyrin, β-amyrin, lupeol (Mohammad et al., 2015), Kaempherol (Formica and Regelson 1995), Rutin (Dar and Tabassum 2012), Benzoic acid (Hansch and Von Kaulla 1970) are likely to be responsible for thrombolytic activity. There is no data available in literature about thrombolytic activity of F. palmata extract. Our study reported a significant thrombolytic potential of ethanolic extract of aerial part of F. palmata.

To best of our knowledge, there is no literature found about the cytotoxic activity of *F. palmata* extract against HepG2 cell line. We are reporting cytotoxicity of *F. palmata* extract against HepG2 cell line for the first time, in which we observed remarkable decline in the viability of cancer cells. This effect could be attributed to the high phenolic content of *F. palmata* with antioxidant potential. These antioxidants work by scavenging free radicals which are responsible for many pathological diseases including cancer, heart disease, aging, alzheimer's disease, inflammation, neurodegenerative disorder (Menaga et al., 2021).

In literature, there is no *in silico* studies data available on F. palmata. The docking study of F. palmata have been employed for the prediction of binding affinity and to find the best pose of the respective compounds to bind within the active sites of diabetic enzymes. For better understanding about the inhibition ability of the studied bioactive phytochemical compounds, 9 phytochemical compounds of ethanolic extract (Taraxasterol, α-amyrin, β-amyrin, stigmasterol 22,23-dihydro, urs-12-en-24-oic acid, 3-oxo-methyl ester, lupeol, brucine, acetic acid, 4,4,6a,6b,8a,11,11,14b-octamethyl-13-oxodocosahydropi cen-3-yl ester and stigmasterol) along with acarbose were docked against α-glucosidase and 8 compounds (Taraxasterol, α-amyrin, β-amyrin, stigmasterol 22,23-dihydro, urs-12-en-24oic acid, 3-oxo-, methyl ester, lupeol, brucine, stigmasterol) along with a arbose were docked against α -amylase enzyme. ADME and toxicity studies were also performed for the evaluation of its pharmacokinetic behavior and toxicity level of major compounds analyzed by GCMS.

5. Conclusions

The present research exploring the *in vitro* and *in silico* activities, GCMS and phytochemical composition of *F. palmata* extract. The extract exhibited the high total phytochemical compounds which support the high antioxidant and significant enzyme (α -amylase and α -glucosidase) inhibition activities. The extract was evaluated for

antibacterial activity (against seven bacterial strains) which, due to certain antibacterial phytochemicals; revealed good antibacterial activity. Strong antiviral and anticancer activities were exerted by the plant against certain viral strains and cell line (HepG2) because of presence of some important phytocompounds in the extract, justifying pharmacological activities of plant. The inhibition activity of F. palmata extract against diabetic enzymes were further evaluated by in silico studies such as molecular docking of all GCMS phytocompounds, from which the selected compounds (best docked) as Taraxasterol, α-amyrin, β-amyrin, stigmasterol 22,23-dihydro, urs-12-en-24-oic acid, 3-oxo-, methyl ester, lupeol, acetic acid, 4,4,6a,6b,8a,11,11,14b-octame thyl-13- oxo docosahydropicen -3-yl ester, brucine, and stigmasterol, undergo ADME and toxicity studies. ADME and toxicity study indicated pharmacokinetic behavior, drug likeness and toxicity level of best docked compounds. The results of biological assays of F. palmata ethanolic extract presented this plant to be uncovered treasure for the discovery of some lead compounds that can be applied against several disorders. In-depth in vivo studies to evaluate the therapeutic potentials of this plant are highly recommended.

Funding

The authors are grateful to the King Saud University, Riyadh, Saudi Arabia for funding this study through Project number RSP2022R504.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Acknowledgements

The authors acknowledge to the King Saud University, Riyadh, Saudi Arabia for funding this study through Project number RSP2022R504.

Appendix A. Supplementary material

Supplementary data to this article can be found online at https://doi.org/10.1016/j.arabjc.2022.104455.

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