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The Anti-Diabetic potential of the African *Adansonia digitata* L. Plant Extract

A thesis submitted in partial fulfillment of
the requirements for the degree of
Master of Science in Biotechnology

By
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2020

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بسم الله الرحمن الرحيم

In the Name of Allah, the Most Gracious, the Most Merciful

Abstract

Diabetes Mellitus (DM) is a severe chronic disease affecting a large population worldwide. The Middle Eastern region has been gifted with lots of native medicinal plants. However, there is less information on development of these medicinal plants into nutraceuticals for modulating specific diseases such as diabetes of common occurrence regionally. Towards this end, there is a crucial need to investigate other types of treatments to modulate diabetes. This study is conducted to fill in this gap by investigating medicinal plant used as folk medicine, which is Baobab (*Adansonia digitata* L.). This study is performed in two phases: *in-vitro* and *in-vivo*. The *in-vitro* part included extraction, assessment, and characterization of the baobab fruit pulp crude extract. The cytotoxic activity was assessed using MTT assay on L-929 fibroblast cells, with an IC₅₀ value of 105.7 µg/mL, whereas DPPH was used to assess the antioxidant activity with calculated IC₅₀ at 114.8 µg/mL. To characterize metabolites mediating for these effects, ultrahigh performance (UHPLC) analysis coupled to MS revealed for a total of 77 metabolites belonging to different chemical classes including organic acids, sugars, alcohols, phenolics, coumarins, and fatty acids. Almost 50% of the identified metabolites are reported for the first time in *A. digitata* fruits. In the second phase, the extract was tested *in-vivo* in an experimental streptozotocin (STZ) induced Sprague Dawley male rat model of diabetes. *A. digitata* L. low dose (150 mg/kg) injected i.p. twice a week showed potential hypoglycemic activity as revealed from several biochemical parameters such as fasting blood glucose, alkaline phosphatase, and blood urea nitrogen levels in comparison to high dose (300 mg/kg). Overall, the current study proves that *Adansonia digitata* L. has good potential as an antidiabetic, hepatoprotective, and reno-protective drug with a good safety margin.

Keywords: Diabetes Mellitus, Middle Eastern plants, *Adansonia digitata* L., Diabetic animal model, metabolomics, metabolites, streptozotocin, STZ, Sprague Dawley rats

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List of abbreviations

DM: Diabetes Mellitus

β-cells: Beta Cells

STZ: Streptozotocin

ALT: Alanine Amino Transferase

AST: Aspartate Transaminase

TG: Triglycerides

MTT: (3-[4,5-dimethylthiazol-2-yl]-2,5 diphenyl tetrazolium bromide)

DPPH: (2,2-diphenyl-1-picryl-hydrazyl-hydrate)

UPLC: Ultra Performance Liquid Chromatography

HPLC: High Pressure Liquid Chromatography

GC/MS: Gas Chromatography Mass Spectrometry

LC/MS: Liquid Chromatography Mass Spectrometry

UHPLC-MS: Ultrahigh-pressure (or performance) liquid chromatography – Mass Spectrometry

BPC: Base Peak Chromatogram

m/z: mass-to-charge ratio

DMEM: Dulbecco's Modified Eagle Medium

FBS: Fetal Bovine Serum

Pen-strep: Penicillin Streptomycin

MeOH: Methanol

DMSO: Dimethyl Sulfoxide

H&E: Hematoxylin & Eosin Stain

Dedication

To my beloved father and Mother

*Any words will not be enough for your efforts with me since primary school. Thank you for always encouraging, supporting, and believing in me. Every single achievement in my life is because of you and will always be for both of you. **Mr. Tawfik Badawy and Mrs. Eisha Abdel-Kreem**, I will always make you proud of me!*

To my twin sister, my brother, and my brother in law

My soulmate Maha Tawfik she is the second living part of me, and Mahmoud Tawfik the wisest and supportive brother. Thank you both for always being by my side. My brother in law, my sister's Fiancé, Mahmoud Elfakhrany thanks a million for everything and for being by my side.

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Chapter One: Introduction and Literature Review

1.1. Introduction:

Diabetes Mellitus (DM) is a chronic disease characterized by disturbed glucose metabolism, affecting an estimated 8.8 percent of the people worldwide, with an expected increase in this percentage to 9.9 percent in 2045. Egypt is among the top countries in the diabetic population in 2019 (Figure 1), with 8.9 million diabetic people (Elflein, 2019).

In 1550 BC, the Ebers papyrus contained a description of several diseases, including polyuric syndromes (Ebbell, 1937). Araetus (81-138 BC), another great scientist, also noted that diabetes is a sort of polyuric waste disorder (Araetus, 1856). The word “diabetes” means a urine fountain. Araetus stated that “Diabetes is a wonderful affliction being “melting” down of the flesh and limbs into urine” (Araetus, 1856). Furthermore, diabetes mellitus was well known for its association with “honey urine.”

Diabetes (DM) has three major types, which are; type I, type II, and gestational diabetes (Colberg et al., 2016). Diabetes type I may be caused by autoimmune reactions destroying the pancreatic β -cells, resulting in a lack of insulin. Therefore, type I is insulin-dependent, and patients rely on the external supply of insulin for life. On the other hand, diabetes type II, the most common type, is non-insulin-dependent. Gestational diabetes occurs during pregnancy; however, the mothers could develop type II DM later on (Dyck, 2003). Pre-diabetes is a term commonly used to describe conditions where blood glucose level is higher than the normal, but not high enough for DM diagnosis (figure 2) to be made (Prevention, 2020). Some risk factors are associated with diabetes, such as obesity, hypertension, and hyperlipidemia (Steyn & Damasceno, 2006).

To date, there is no cure for diabetes (Prevention, 2020). The management of DM is not without disadvantages and complications such as drug resistance and toxicity. Additionally, uncontrolled diabetes could result in chronic complications such as blindness, heart disease, and renal failure (Mamun-or-Rashid et al., 2014). Hence, there is a crucial need to investigate other types of treatment to modulate this chronic disease.

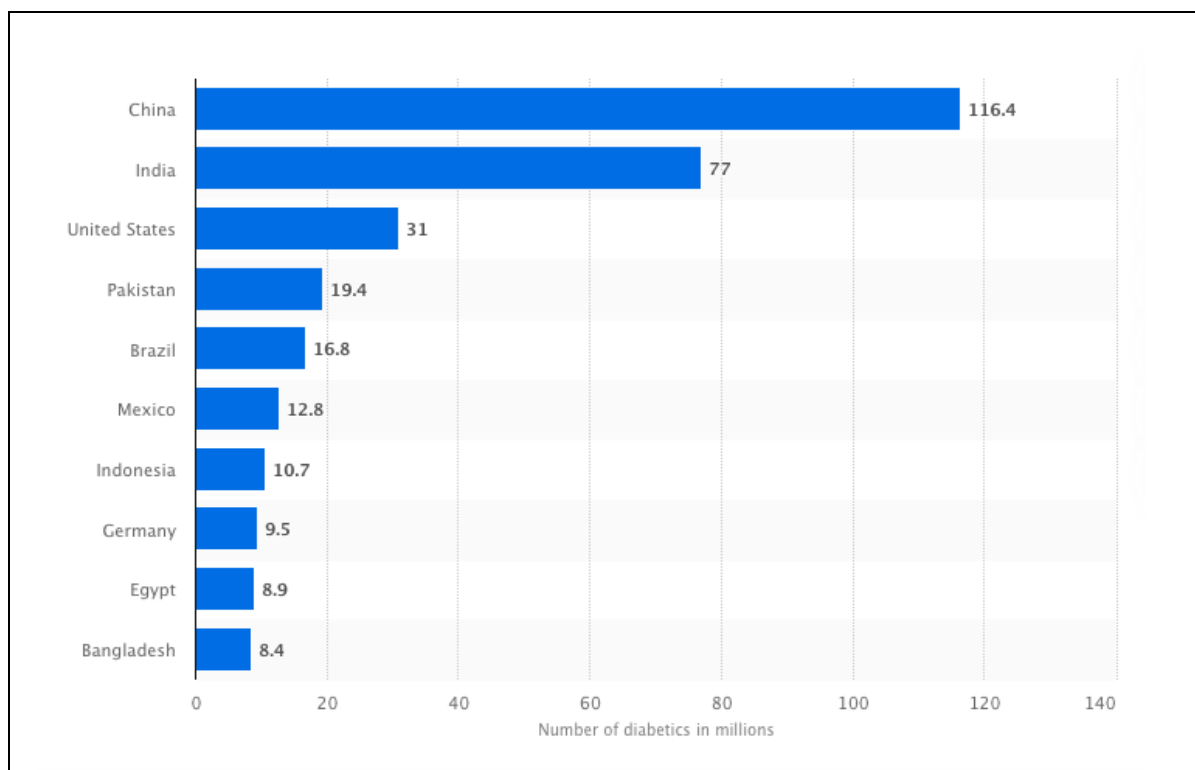


Figure 1: Estimated number of diabetic population worldwide in 2019, (Elflein, 2019).

The Middle East region is rich with medicinal plants used as folk medicines that have a great impact on diseases like DM (Öztürk, Altundag, & Gücel, 1999). However, their side effects must be considered before use (P. Gupta, 2012; Kooti, Farokhipour, Asadzadeh, Ashtary-Larky, & Asadi-Samani, 2016; Mamun-or-Rashid et al., 2014). Although there is an abundance of literature worldwide on the use of plant extracts in diabetes, there are few reliable studies on Baobab (Tabaldi) “*Adansonia digitata L.*,” a plant native to Africa and traditionally used to treat DM.

Therefore, the aim of the current study is to investigate this crude plant extract and characterize its effects on diabetic complications in a well established experimental animal model of diabetes (Srinivasan & Ramarao, 2007). Animal models may be considered cruel by animal rights activists. We believe that such models, when humanely and appropriately used, provide an excellent opportunity to study diseases and provide significant pre-clinical studies that would improve patient care worldwide. (Rees & Alcolado, 2005).

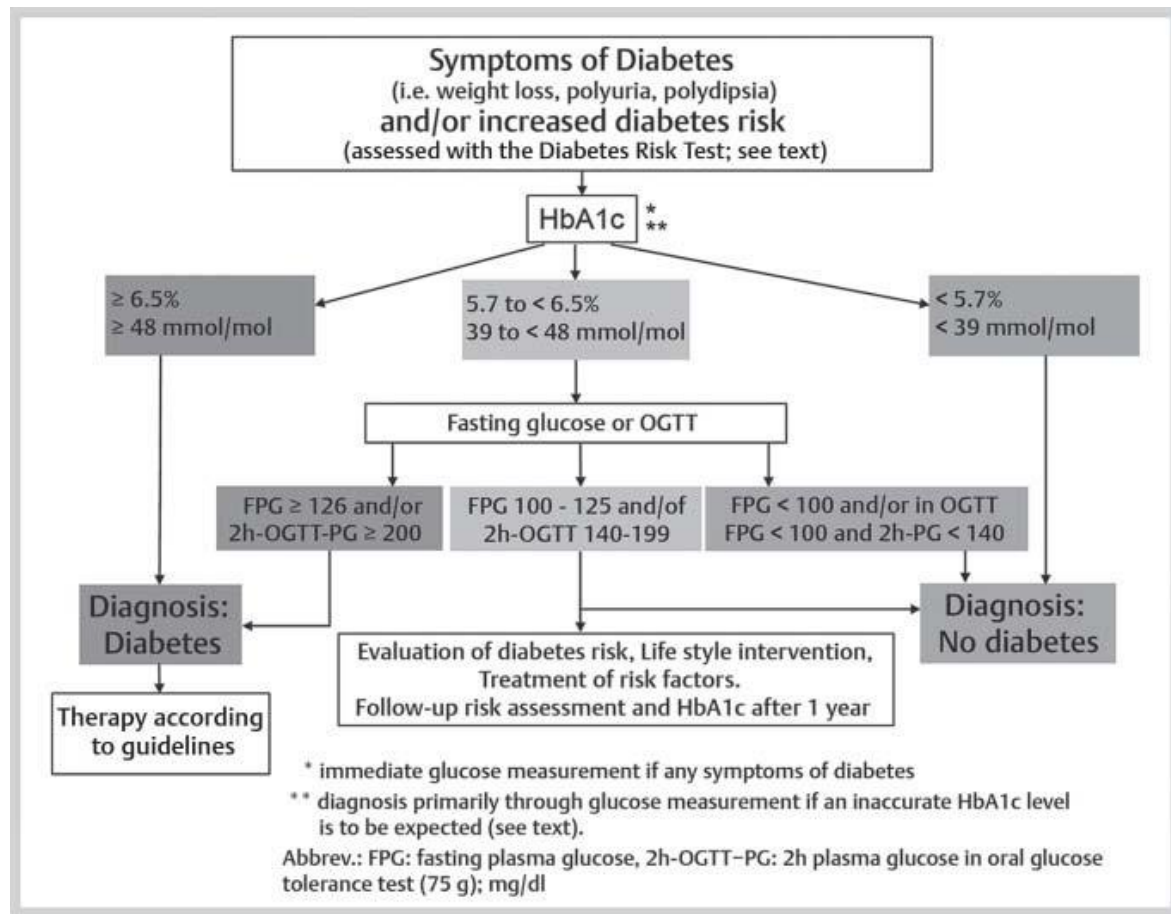


Figure 2: *Diabetes Mellitus* Diagnostic flowchart (Kerner & Brückel, 2014).

1.2 Hypothesis

The *Adansonia digitata* L. fruit pulp crude extract will have beneficial antidiabetic effects in STZ-diabetic rats.

Aims:

- Extraction and characterization of *Adansonia digitata* L. plant.
- *In vitro* testing of the cytotoxic and antioxidant activities of the plant extract.
- *In vivo* testing of the plant extract, in an animal model of *Diabetes Mellitus*.

1.3 Animal Models of Diabetes

Experimental animal models of diabetes mellitus (DM) can be divided into three main groups; chemical, surgical, or genetically-induced diabetes (S. Kumar, Singh, Vasudeva, & Sharma, 2012). Chemical induction of diabetes is achieved with diabetogenic agents, such as Streptozotocin (STZ), Alloxan (ALX), Goldthioglucose (GTG), as well as other drugs such as antipsychotic drugs. These chemicals either cause entire damage of the β - cell, develop an inhibition of insulin production and secretion, and decrease or minimize the metabolic productivity of insulin in the targeted tissues (M. P. Singh & Pathak, 2015). The most commonly used chemicals for diabetes mellitus induction in animals are Streptozotocin (STZ) and Alloxan (ALX), which are usually injected either intraperitoneal (IP), subcutaneous (SC), or intravenous (IV) injections.

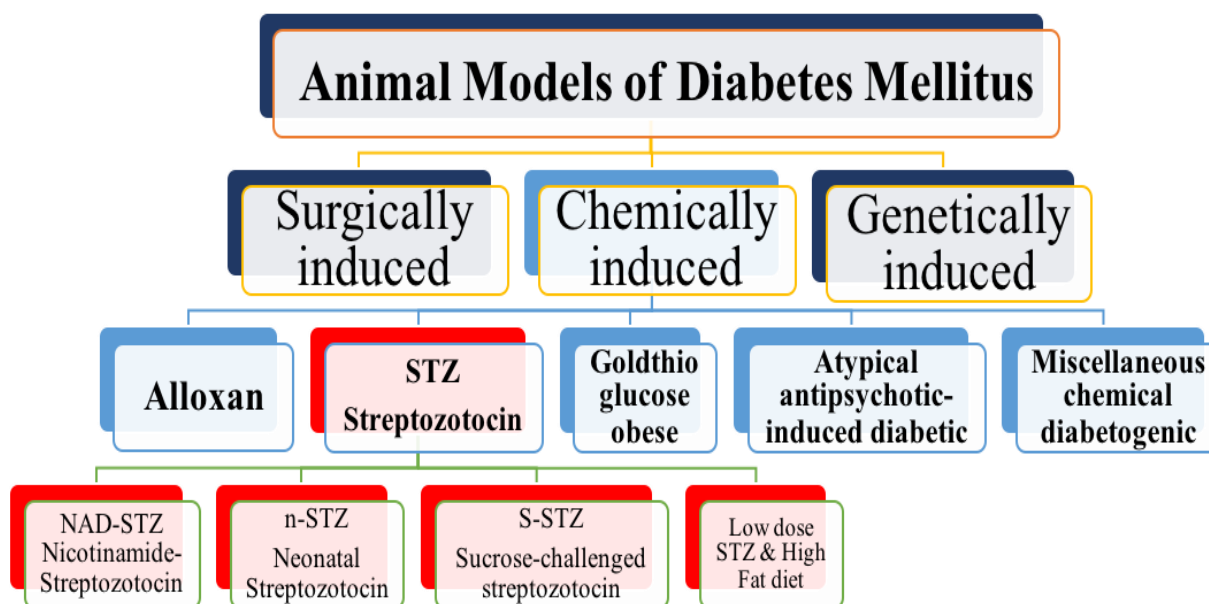


Figure 3: Schematic flow chart of the induced animal models of *diabetes mellitus*.

The second type of diabetic model is surgically induced by either partial (to induce type II DM) or complete removal of the pancreas to induce type I DM (Ozturk, Atlan, & Yildizoglu-Ari, 1996).

However, pancreatectomy is not widely accepted nowadays by researchers for diabetic investigations (Rees & Alcolado, 2005).

The third type is genetically induced diabetes in animals, using several mutations such as (db/db mice), or induction from non-diabetic outbred animals by the repeated breeding. The general concept behind the mutation induction in animals is that a single or multigene defect cause diabetes. Other genetically induced diabetic models are Zucker diabetic fatty rat, Goto-Kakizaki rat, LEW.1WR1 rats, NONcNZO10 mouse, C57BL/6J mice, Kuo Kundo mice, Tsumara Suzuki Obese Diabetes mice, db/db mice, and Obese rhesus monkey (*Macaca mullata*) (Ktorza et al., 1997).

There are also other unclassified models of diabetes, such as virus-induced diabetic animal model, Oral glucose loading animal model, Insulin Antibodies-induced diabetes, and In-vitro models for diabetes (Ktorza et al., 1997).

1.4 Streptozotocin-Induced Diabetic Model

Streptozotocin (STZ) is a glucosamine-nitrosourea complex initially marketed under the name of Zanosar® (R. B. Weiss, 1982). It is the most commonly used diabetogenic drug in inducing diabetes to all species (Junod, Lambert, Stauffacher, & Renold, 1969). Rats, mice, hamsters, dogs, monkeys, and sheep STZ-induced models of diabetes using a wide range of doses in single or multiple injections have been reported widely in the literature (P. Kumar, Kale, & Baquer, 2012; Srinivasan & Ramarao, 2007). In large animals, STZ can be combined with partial pancreatectomy to develop diabetes.

Streptozocin (STZ) is a selective cytotoxic agent that specifically destroys pancreatic β cells. Streptozotocin enters the pancreatic β cells through the glucose transporter 2 (GLUT2). Since STZ consists of two parts, which are glucose and methyl nitrosourea, it splits, and the methyl nitrosourea moiety acts as an alkylating agent, causing DNA alkylation and will destroy the cells (Szkudelski, 2001). However, the cells producing insulin in the pancreas could have some resistance from STZ if they do not have GLUT2 (Ledoux & Wilson, 1984).

There are several modifications of the STZ induced model, e.g., Neonatal STZ induced diabetes rat model (n-STZ), Nicotinamide-Streptozotocin (NAD-STZ) induced model, Sucrose-challenged STZ-induced rat model (STZ-S), and the Low dose STZ with high-fat diet rat model (M. P. Singh & Pathak, 2015). The low dose STZ / high-fat diet rat model is the closest to DM type II in humans and is commonly used for pharmacological screening (Srinivasan, Viswanad, Asrat, Kaul, & Ramarao, 2005).

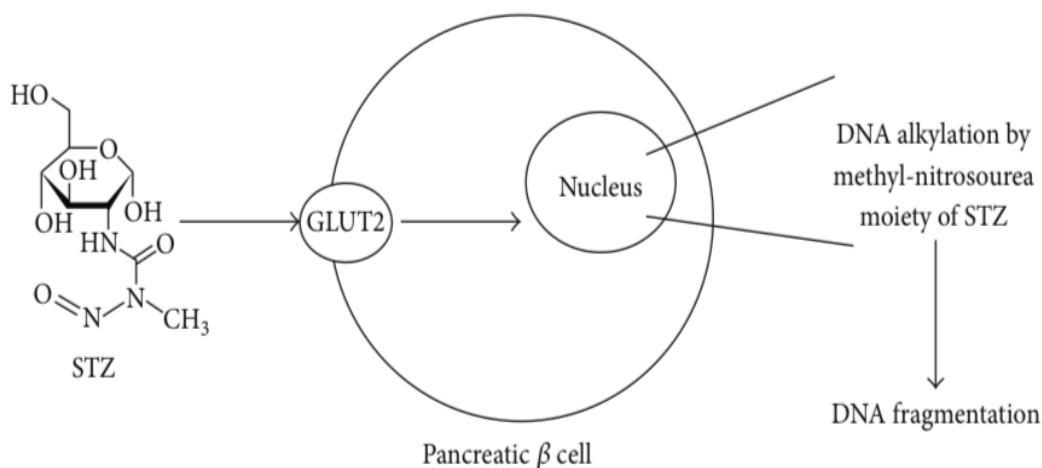


Figure 4: Mechanism of action of streptozotocin (STZ).

STZ enters the β -cells by the GLUT2 receptor (Glucose Transporter 2), causing DNA fragmentation and cell death (Al-Awar et al., 2016).

1.5 Commonly used anti-diabetic medicinal plants

Commonly used medicinal plants in folk medicine, as anti-diabetic plants, are available in the Middle Eastern region such as Tabaldi, (Baobab) (*Adansonia digitata* L.), Kaff Maryam (*Anastatica hierochuntica* L.), Argel (*Solenostemma Argel*), Fenugreek (*Trigonella foenum-graecum*), Hanzel (*Citrullus colocynthis*), Curcumin (*Turmeric*), Cinnamon (*Cinnamomum Zeylanicum*), and Mango (*Mangifera indica*). In this study, we will focus on Baobab (*Adansonia digitata* L.) plant extract, which is native to Sudan and is commonly used in folk medicine as an anti-diabetic treatment.

Fenugreek has some antibacterial, gastric stimulant, anti-diabetic, anticancer, and hepatoprotective effects (Srinivasan, 2006). Caffeic acid & Ellagic acid have some effects in decreasing the plasma glucose levels in diabetic model mice, which are present in fenugreek (Chao, Hsu, & Yin, 2009). The methanolic extract of Argel, it has antioxidant and hypoglycemic effects without any reported toxicity. However, there is little research conducted to investigate its mechanism of action (Deen & Al-Naqeb, 2014). Kaff Maryam methanolic extract significantly increased IgG levels after a dose of 50 mg/kg, which was believed to have some effects on β -cell function (Abdulfattah, 2013).

In a clinical trial, 125 mg per day of Hanzel for 2 months showed an increase in HbA1c and Fasting BS with patients diagnosed with type II diabetes without any side effects reported (Barghamdi et al., 2016). In another clinical trial using Curcumin, lowering of triglycerides, alanine aminotransferase (ALT), and beta-amyloid was seen (S. C. Gupta, Patchva, & Aggarwal, 2013).

Cinnamon showed a very good ability to normalize glucose metabolism, weight changes, and lipid abnormalities in an STZ animal model of diabetes (Anand, Murali, Tandon, Murthy, & Chandra, 2010). Mango can decrease hyperglycemia in diabetic rats at concentrations of 5, 10, and 15% (Perpetuo & Salgado, 2003). Phytochemical analyses were performed for the methanolic mango extract, and it showed the presence of glycosides, flavonoids, tannins, saponins, terpenoids, and steroids (Gwarzo & Bako, 2013; Zekry, Badawy, Ezzelarab, & Abdellatif, 2020).

There are currently few reliable studies of the *Adansonia digitata* L. effects on diabetes and its complications such as (Bako, Mohammad, Waziri, Bulus, & Zubairu., 2014) . As a result, this study is designed to provide a further investigation of its effects as well as its mode of action.

1.6 Origin and history of the *Adansonia* genus

The tree of *Adansonia* genus occurs mainly in Africa, especially in Madagascar, and in Australia (Baum, 1995). It was named *Adansonia* after the french botanist Micheal Adanson who discovered it in Senegal in 1749. It is also known as the upside-down Baobab tree or monkey bread tree. The name is based on the Arabic word *bu hobab*, meaning “fruit with multiple seeds.” There

are eight different species of Baobab, including six that are native to Madagascar, one to Australia, and another one to the African mainland (Baum, Small, & Wendel, 1998). The Baobab tree has diverse uses by the African people (Baum, 1995).

The first species is *Adansonia digitata* L. (*Malvaceae*) or African baobab, which is distributed across the African continent. *Adansonia digitata* s.l. is another one consisting of the diploid *A. kilima* and the tetraploid *A. digitata* s.s. (Pettigrew et al., 2018). The *A. digitata* s.s. is distributed across the African Sub-Sahara extending from the northern Sahel to South Africa. However, it is not likely to be present in the center of Africa, especially in the rainforest areas, as well as the high elevation above 800 meters of the eastern African area (Gerald E. Wickens & Lowe, 2008).

The phylogeographic analyses showed that the *Adansonia digitata* s.l. Populations are native to eastern and western Africa. Moreover, they are genetically distinct from these particular areas (Pock et al., 2009). The *A. digitata* and *A. kilima* show high genetic similarity. These analyses also indicate that the evolution of the tetraploid species is occurring recently (Pettigrew et al., 2018; Gerald E. Wickens & Lowe, 2008).

Baobabs are known for their high nutritional value as a cultural fruit. The fruit is consisting of seeds surrounded by the powdery pulp in a hard endocarp. The fruit taste is sweet and sour, which is liked by many people who consume it on a regular basis. However, the seeds are not edible until they are roasted or ground (Gebauer, El-Siddig, & Ebert, 2002). People used to consume it by discarding the seeds or allowing them to pass as they are in the digestive system after eating the fruit pulp. Furthermore, some genetic analyses are suggesting that the spread and distribution of the baobab tree over the continent of Africa is due to human practices (Pock et al., 2009).

The African baobab species spread to other regions located around the Indian Ocean, such as Yemen, Comoros, and southern Iran (Parsa, 1959). The immersion of the baobab fruit in seawater showed tolerance and no effect on the viability of the fruit itself even after six months of immersion in that salt condition (Pock et al., 2009). This might provide an explanation of the presence of the baobabs in the coastal regions across the Indian subcontinent (Gerald E. Wickens & Lowe, 2008).

Some specific populations sharing the same genetic ancestors of baobabs were grouped in the Mascarene Islands, southeast India, Malaysia, and West Africa because of the German, French,

and English people and their movements between the Asian and African continents (Burton-Page, 1968).

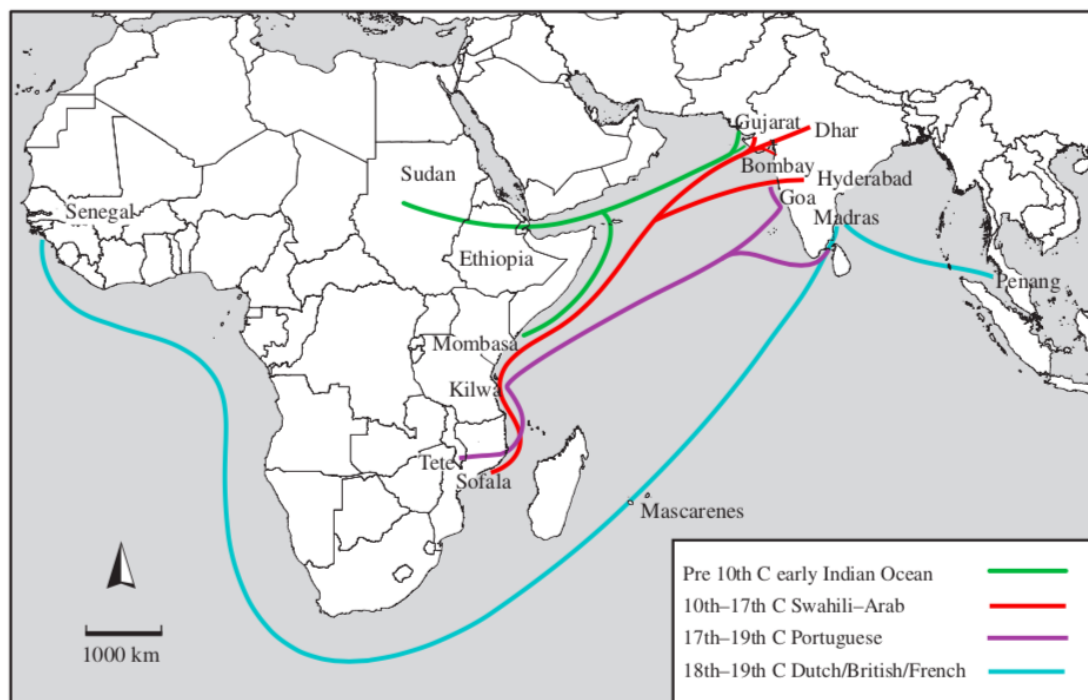


Figure 5: The spread of the *Adansonia digitata* L. tree from Africa.

The introduction of the Baobab species from Africa to the Indian subcontinent. Due to the trade through the ocean during various periods (Bell, Rangan, Kull, & Murphy, 2015).

1.7 *Adansonia digitata* L. plant in Folk Medicine

Tabaldi, Baobab., is called “The small Pharmacy” because of its huge contribution to folk medicine (Obizoba & Anyika, 1994). The fruit pulp is very rich in nutrients as well as carbohydrates, calcium, potassium, thiamine, nicotinic acid, and vitamin C (G. E. Wickens, Goodin, & Field, 1985). Phytochemical analyses for the methanolic extract showed the presence of glycosides, flavonoids, tannins, saponins, terpenoids, and steroids. In addition, this methanolic extract from

the fruit pulp showed anti-diabetic effects in Alloxan induced diabetes models (Gwarzo & Bako, 2013).

All parts of the baobab are edible, e.g., seeds, roots, flowers, leaves, bark, and fruit pulp. The trees of Baobab can be grown even in high temperatures and be resistant to it for a long period of time. Baobabs are known for their excellent antioxidant and anti-inflammatory properties since ancient times. It is also used to treat several microbial infections as traditional medicine. Soups can be prepared from the baobab leaves, and these soups can be thickened by the addition of the seeds. Furthermore, the seeds can be roasted in order to be used as snacks (Caluwé, Halamová, & Damme, 2010). Baobab fruit includes pulp with large seeds embedded in the acidic pulp. The pulp is used for food recipes as well as making some beverages (Obizoba & Anyika, 1994).

The bark, leaves, and fruits of *Adansonia digitata* are used as medicinal plants in South Africa. Fresh bark beverage is used for cold and influenza symptoms (Sulaiman et al., 2011). Furthermore, the seeds, wood, and bark of the baobab are used as anti-inflammatory agents (Selvarani & James, 2009). The leaves are famous for treating fevers, kidney, and bladder diseases. In addition to their effects on asthma and diarrhea. A mixture of the flour of pulp mixed with millet flour and water is given to the children for the treatment of the infant diarrhea (Lockett, Calvert, & Grivetti, 2000). *Adansonia digitata* L. plant showed potential antiviral effect against three tested viruses including herpes simplex, polio, and Sindbis (Ananil, Hudson, & Souzal, 2000).

In West Africa, the seeds, fruit pulp, and the leaves are considered as essential components in sauce preparation (Caluwé et al., 2010). Baobabs are also called “super fruit” thanks to their high nutrition value of vitamins, minerals, and fatty acids contents (Gruenwald, 2009). The baobab products are important because of their dietary fibers and ascorbic acid components. The fruit is considered as one of the richest fruits in Vitamin C, which is present in a range of 2.8 to 3 g/kg in the fruits (Silvia Vertuani, Braccioli, Buzzoni, & Manfredini, 2002). Another study revealed that the percentage of vitamin C present in the fruit pulp is (280-300 mg/100 g), which is seven times the vitamin C content found in the oranges (51 mg/100 g) (Manfredini, Vertuani, & Buzzoni, 2002). A recent study stated that the baobab pulp could provide 100% of the recommended daily intake of vitamin C for pregnant females when consuming 40 g of the fruit (Chadare, Linnemann,

Hounhouigan, Nout, & Van Boekel, 2008). Regarding the actual content of the ascorbic acid, it was found that the fruit pulp contains 337 mg/100 g (Eromosele, Eromosele, & Kuzhkuzha, 1991).

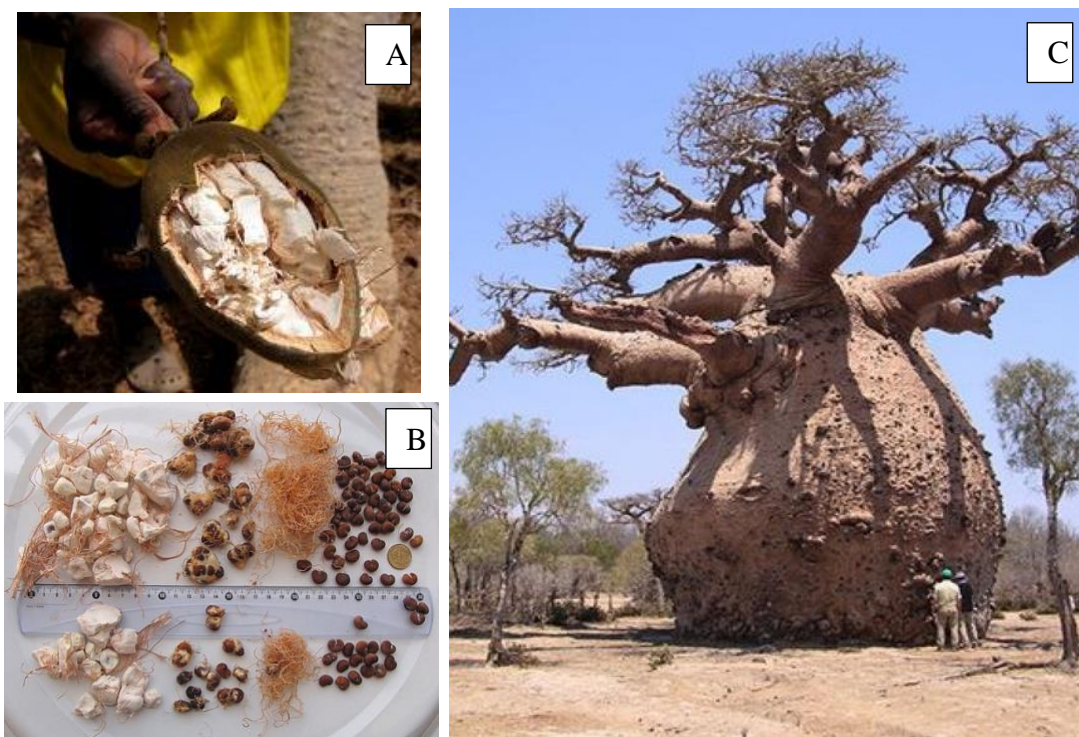


Figure 6: *Adansonia digitata* L. Plant

(A): The Baobab fruit pulp. **(B)**The Baobab fruit pulp and seeds. **(C)** The upside-down Baobab tree picture. Source: wo.wikipedia.org.

1.8 Chemical content of *Adansonia digitata* L.

The seed of the *Adansonia digitata* L. contains high amounts of fat ($12.2 \pm 0.2\%$), crude fiber ($16.3 \pm 9.3\%$), and protein ($18.4 \pm 0.5\%$). The *Adansonia digitata* L. seed oil is a great source of mono as well as polyunsaturated fatty acids. The composition of the oil is mostly saturated fatty acids, monounsaturated fatty acids, and polyunsaturated fatty acids at 31.7%, 37%, and 31.7% respectively. The oleic acid is the major fatty acid with 35.8%, then the linoleic 30.7%, and finally the palmitic 24.2% (Lockett et al., 2000; Ralaimanerivo, Gaydou, & Bianchini, 1982).

The fruit pulp is considered a great source of carbohydrates ($76.2 \pm 1.0\%$), low fat ($0.3 \pm 0.0\%$), and low protein ($8.2 \pm 0.1\%$). The fruit pulp and seed are great sources of calcium, magnesium, and potassium.

Amino acids content in both seed and fruit are presented by high amounts of aspartic acid, arginine, and glutamic acid (Glew et al., 1997).

The leaf extract contains succinic, lactic, malic, citric, formic, and fumaric acids were identified as well as the fruit pulp. The RP-HPLC-PDA-ESI-MS/MS (High Performance Liquid Chromatography-Photo Diode Array-Electrospray Ionization-Mass Spectrometry) analysis also confirmed some organic acids including citric acid at 77.3%, malic acid (17.6%), ascorbic acid (3.2%), and gallic acid (1.3%) (Tsetegho Sokeng et al., 2019).

Phenolic analysis revealed variety of classes such as tannins, flavones, and flavonols. Flavonoids (flavanols, flavones, flavonols, and proanthocyanidins) as well as the non-flavonoids (hydro-xycinnamic derivatives) were detected in the baobab fruit pulp (Tsetegho Sokeng et al., 2019). Furthermore, hydroxy cinnamates were identified by (N. X. Li et al., 2017) and proanthocyanidins by (Shahat, 2006).

Table 1: Summary of the previously reported work on the anti-diabetic and chemical composition of the *Adansonia digitata* L.

Plant Part	Type of extraction	Route of administration	Chemical composition	Chromatographic analysis - Fractionation	Reference
Fruit pulp	Methanol and Millipore grade water acidified with formic acid (0.1%)	N/A	<ul style="list-style-type: none"> - Sugars and derivatives - Organic acids - Amino acids - Phenolic compounds 	RPHPLC-PDA-ESI-MS/MS	(Tsetegho Sokeng et al., 2019)
	Ethyl Acetate	N/A	<ul style="list-style-type: none"> - hydroxycinnamic acid glycosides (HAGs) - iridoid glycosides (IGs) - phenylethanoid glycosides (PGs) 	UHPLC-DAD-HRMS	(N. X. Li et al., 2017)
	Ethyl Acetate	N/A	<ul style="list-style-type: none"> - proanthocyanidin 	TLC	(Shahat, 2006)
	Methanol	Intraperitoneally/ Orally	N/A	N/A	(Bako et al., 2014)
	Aqueous	Orally	N/A	N/A	(Muhammad, Jarumi, Alhassan, Wudil, & Dangambo, 2016)
Leaves	Methanol and Millipore grade water acidified with formic acid (0.1%)	N/A	<ul style="list-style-type: none"> - Sugar and derivatives - Organic acids - Amino acids - Phenolic compounds 	RP-HPLC-PDA-ESI-MS/MS	(Tsetegho Sokeng et al., 2019)

Chapter two: Materials and Methods

2.1. Extract preparations

Adansonia digitata L. was purchased from the local market in Sudan. The fruit pulp was crushed and ground into powder. Alpha Chemika Methanol reagent M.W.= 32.04 was used in preparing the crude extract. This powder was dissolved in methanol at a ratio of 2:10, 100 grams were soaked in 500 ml methanol for 48 hours. The extract was filtered using Whatman filter paper No. 1. The rotary evaporator Rotavapor® BÜCHI was used to evaporate the methanol from the extract, and the sample was left to air dry for 3-5 days in a hood. The extract was prepared freshly for further use (modified from (Bako et al., 2014)).

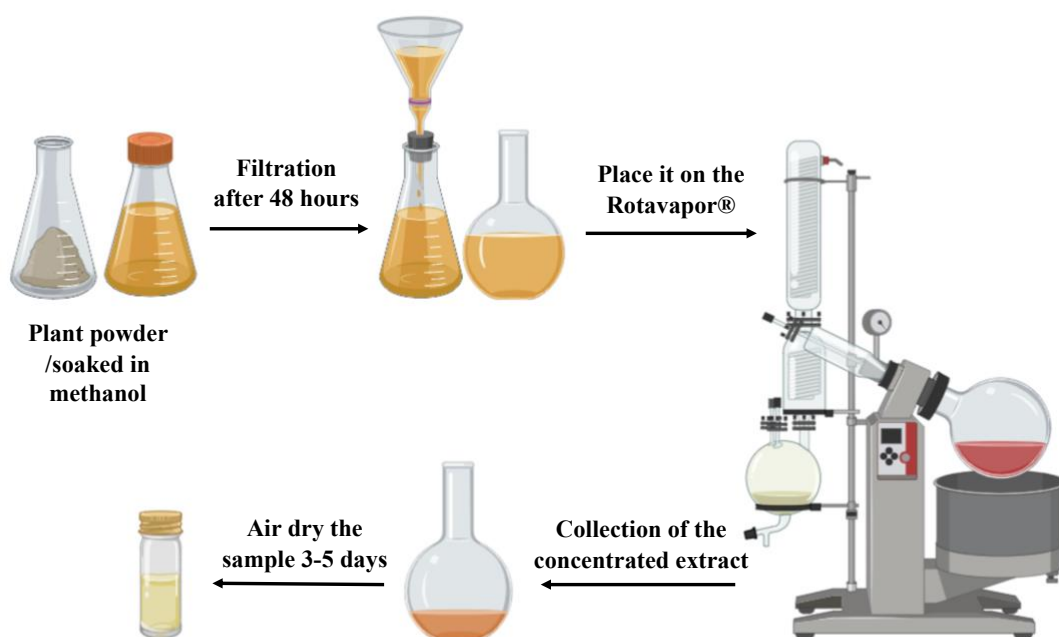


Figure 7: *Adansonia digitata* L. plant crude extract preparation procedure

2.2. Cytotoxicity Assay (MTT)

3-[4,5-dimethylthiazol-2-yl]-2,5 diphenyl tetrazolium bromide assay was used to measure the cell viability. Cell's mitochondrial activity can be indicated by the conversion of the tetrazolium salt MTT into formazan crystals. The viable cell number is measured by detecting the formazan optical density (OD) at 570 to 720 nm. The MTT assay is widely used for the measurements of the drug sensitivity in cell lines and primary cells (van Meerloo, Kaspers, & Cloos, 2011).

L929 fibroblast cells (ATCC® CCL-1™, 100 old-day male rat) were seeded in 96-well plates in Dulbecco's Modified Eagle Medium (DMEM) supplemented with 1% Penicillin-Streptomycin antibiotic and 5% Fetal Bovine Serum (Canfield, 2010). The cells were incubated for 24 hours in Thermo Scientific™ Heracell™ VIOS CO₂ incubator with 37° C and 5% CO₂. The media was removed, and the control, blank, and extract were added to the cells. After the second incubation of 24 hours, the media was removed, and the MTT (3-[4,5-dimethylthiazol-2-yl]-2,5 diphenyl tetrazolium bromide) reagent was added and incubated for 4 hours. Finally, the MTT was removed, and Dimethyl sulfoxide (DMSO) was added to the wells. The optical density (OD) was measured using SPECTROstar® Nano plate reader at 570 nm (modified from (van Meerloo et al., 2011)).

2.3. Antioxidant Activity Using (DPPH)

DPPH is the abbreviated term of free radical α , which is 2,2-diphenyl-1-picryl-hydrazyl-hydrate (DPPH; C₁₈H₁₂N₅O₆, M=394.33). This assay aims at the determination of the antioxidant activity by the use of the stable free radical of α . DPPH depending on the scavenging capacity measurement of the antioxidant agents (Blois, 1958). It works by the odd electron of the DPPH nitrogen atom is being reduced by receiving the hydrogen atom from the tested antioxidant material (Contrerasguzman & Strong, 1982).

The DPPH powder is prepared by weighing 3.94 mg and dissolving it in 100 ml methanol or ethanol for 0.1 mM solution preparation. This assay was performed in a 96-well plate. In brief, 50 μ L of the DPPH solution are added to 150 μ L of the tested extract (ratio of 1 ml DPPH to 3 ml of the tested sample). The mixture is allowed to stand in the dark with vigorous shaking at room

temperature for 30 minutes. Finally, the absorbance of the different concentrations was obtained using a SPECTROstar® Nano plate reader at 517 nm optical density. Ascorbic acid was used as a strong antioxidant material (modified from (Chiabchalard, 2013)).

2.4. (UHPLC/MS-MS) Analysis

2.4.1. Preparation of fruit extracts for UHPLC-MS analysis

Briefly, the freeze-dried fruits were ground with a pestle in a mortar, then 2 g of each fruit powder was homogenized with 5 mL 100% MeOH containing 10 µg/mL umbelliferone (an internal standard used for relative quantification of UHPLC-MS features) (Guillarme & Veuthey, 2017). Extracts were then vortexed vigorously and centrifuged at 3000 g for 30 min to remove plant debris. For solid-phase extraction, 500 µL were aliquoted and loaded on a (500 mg) C18 cartridge, which was preconditioned with methanol and water. Samples were then eluted with 3 mL 70% MeOH and 3 mL 100% MeOH. The eluents were evaporated to dryness under a gentle nitrogen stream. The obtained dry residue was re-suspended in 500 µL methanol for further UHPLC-MS analysis (modified from (Guillarme & Veuthey, 2017)).

2.4.2. High-resolution UHPLC-MS analysis

The UHPLC analysis was performed on an Acquity UHPLC System (Waters) equipped with a HSS T3 column (100 × 1.0 mm, particle size 1.8 µm; Waters). The analysis was carried out by applying the following binary gradient at a flow rate of 150 µL min⁻¹: 0–1 min, isocratic 95% A (water/formic acid, 99.9/0.1 [v/v]), 5% B (acetonitrile/formic acid, 99.9/0.1 [v/v]); 1–16 min, linear from 5 to 95% B; 16–18 min, isocratic 95% B; and 18–20 min, isocratic 5% B (M. Farag, Mahrous, Lübken, Porzel, & Wessjohann, 2013).

The injection volume was 3.1 µL (full loop injection). Eluted compounds were detected from m/z 90 to 1000 using a MicroTOF-Q hybrid quadrupole time-of-flight mass spectrometer (Bruker Daltonics) equipped with an Apollo-II electrospray ion source in negative and positive (deviating values in brackets) ion modes using the following instrument settings: nebulizer gas, nitrogen, 1.4 (1.6 bar); dry gas, nitrogen, 6.1 min⁻¹, 190 °C; capillary, –5000 V (+4000 V); endplate offset, 500

V; funnel 1 RF, 200 Vpp; funnel 2 RF, 200 Vpp; in-source CID energy, 0 V; hexapole RF, 100 Vpp; quadrupole ion energy, 5 eV (3 eV); collision gas, argon; collision energy, 7 eV (3 eV); collision RF, stepping 150/350 Vpp (200/300 Vpp), (timing 50/50); transfer time, 58.3 μ s; prepulse storage, 5 μ s; pulser frequency, 10 kHz; and spectra rate, 3 Hz. Internal mass calibration of each analysis was performed by infusion of 20 μ L 10 mM lithium formate in isopropanol[thin space (1/6-em)]:[thin space (1/6-em)]water, 1[thin space (1/6-em)]:[thin space (1/6-em)]1 (v/v), at a gradient time of 18 min using a diverter valve. For Auto-MS/MS analysis, precursor ions were selected in Q1 with an isolation width of ± 3 –10 Da and fragmented at collision energies of 15–70 eV using argon as collision gas. Product ions detection was performed using the same settings as above, but with funnel 2 RF 300 Vpp in negative mode (Mohamed A. Farag et al., 2016).

2.5. Animals and Induction of Diabetes

All experiments were conducted in accordance with the NIH guidelines for Animal Care and Use. Animal use protocol was approved by MSA University, Faculty of Biotechnology Animal Care and Use Committee (Courtesy of Dr. Mona Saadeldin, MSA University). Male Sprague Dawley rats weighing 150 - 200 g were maintained in a temperature-controlled environment at approx. 24 °C with a 12-h light/dark cycle and ad libitum access to food and water.

Streptozotocin (STZ) was injected intraperitoneally at a dose of 65-75 mg/kg (Furman, 2015) (Bonnie-Nielsen, Steffes, & Lernmark, 1981; Srinivasan & Ramarao, 2007). Confirmation of diabetes was performed by measuring blood glucose levels using samples collected from the tail vein 3-5 days after the injection. Animals with glucose concentration > 250 mg/dL were considered diabetic and included in further experimentation. Pancreatic tissue histological examination after animal termination confirmed Diabetes mellitus (Qinna & Badwan, 2015).

The rats were randomized into six groups (Altman, 2002); each group consisted of 6 rats. The groups were:

1st group: normal control healthy (negative control),

2nd group: diabetic control untreated (positive control),

3rd group: diabetic + metformin (administered orally in the drinking water),

4th group: normal healthy + High dose (300 mg/kg) of *adansoina digitata* L. extract (intraperitoneally, twice a week),

5th group: diabetic + low dose (150 mg/kg) of *adansoina digitata* L. extract (intraperitoneally, twice a week),

6th group: diabetic + high dose (300 mg/kg) of *adansoina digitata* L. extract (intraperitoneally, twice a week) (modified from (Bako et al., 2014)).

The study duration was 4 weeks. Blood glucose and body weight were monitored twice a week. After four weeks, the rats were sacrificed, and blood and tissue samples were collected for biochemical (renal function, liver function, and lipid profile) and histological testing.

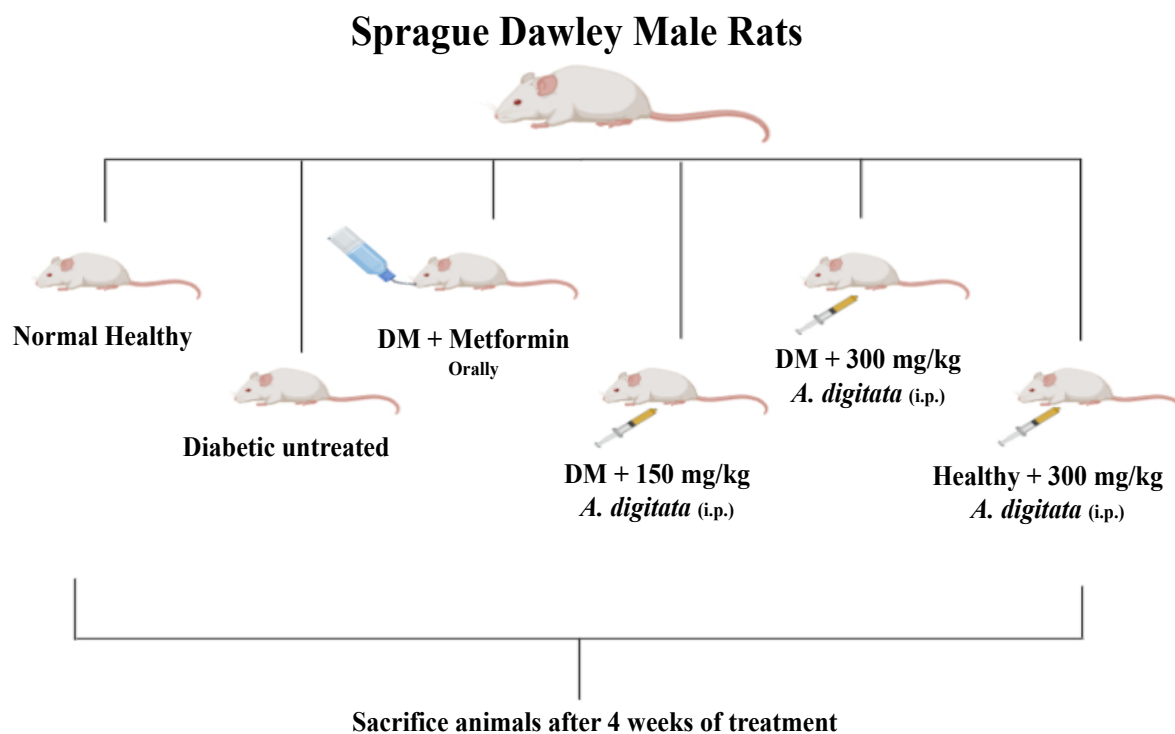


Figure 8: Sprague Dawley Male Rats' groups flow chart

2.6. Biochemical Analysis

Blood samples (2-3 mL) were collected following animal sacrifice for liver, kidney, and lipid profile assessment. Blood was collected in and centrifuged at 3000 r.p.m. for 5 minutes. The blood

biochemical parameters were measured using the commercially available kits (SPECTRUM company, Shanghai, China). Statistical significance was determined using one-way ANOVA, and $p < 0.05$ was considered significant.

2.7. Histopathological Analysis

After four weeks of treatment, rats were sacrificed, and samples collected. The liver, kidney, and pancreas samples were washed in Saline solution and post-fixed for 24 hours in 4% paraformaldehyde. Samples were dehydrated in alcohol (70%, 80%, 90%, 95%, and 100%), cleared in terpineol for 24 hours, and infiltrated in Paraplast at 60 °C for 2 hours. Finally, the tissues are embedded in paraffin wax then sectioned using Leica Rotary Microtome™ (Model: 1512). 5-6 µm thick sections are mounted on clean glass slides. Paraffin sections were deparaffinized and stained with Mayer's Hematoxylin and counterstained with 1% Eosin (H&E) (Lillie & Fullmer, 1976).

The diameter of pancreatic islets was evaluated according to the methods described by (Glaharn, Punsawad, Ward, & Viriyavejakul, 2018). An average of 40 islets for each animal was measured under 100 x using image analysis software (ImageJ, National Institutes of Health, version 1.5a). Data were presented as mean \pm SD applying one-way analysis of variance (ANOVA) followed by Tukey's comparison test using GraphPad Instat (Prism). A P- a value of < 0.05 was considered to be statistically significant.

The scoring of liver and kidney damage was performed according to previous studies (Klopfleisch, 2013; Knodell et al., 1981), the grade of damage was obtained by adding the entire score of the types of histopathological lesions (4-5 field/rat/200x), as described in the appendix **Table 4**. Differences between the data were analyzed using Kruskal–Wallis test, followed by Dunn's Multiple Comparison Test. Data were expressed as median \pm SD.

2.8. Statistical Analysis

Statistical analysis was performed using R software version 3.5.2 and GraphPad PRISM version 5. R software was used for the MTT analysis to determine the toxicity level for the different

concentrations. One-Way ANOVA (Analysis of Variance) for multiple comparisons among samples/groups and t-test was used to determine significance. Data are presented as mean \pm SD.

Bruker Daltonics Data Analysis 4.4 was used for chromatograms inspection. While Metaboscape (Bruker Daltonics) 3.0 was used for data pre-processing using T-ReX 3D (Time aligned Region Complete eXtraction) algorithm for retention time alignment. Where isotopes, adducts, and fragments belonging to the same compound are automatically detected and combined into one feature. Detected features are displayed as a bucket table with their corresponding Rt, measured m/z , molecular weight, and detected fragments. The Bucket table was created with intensity threshold $10e4$ for both negative and positive ionization modes with a retention time range from 0 to 18 min and mass range from 120 to 1600 m/z (Olmo-García et al., 2019).

For the molecular networking and metabolites annotation, the features list created by the Metaboscape was exported as MGF file for both of the positive and negative measurements. The negative and positive mode MGF files were uploaded independently to the GNPS online platform (<https://gnps.ucsd.edu>), where two molecular networks were created following the online workflow (GNPS 2.0). A molecular network was created with a cosine score above 0.65 and 0.7 for positive and negative modes, respectively, and the number of shared fragments was adjusted to 4. The networks were visualized using Cytoscape 3.5.1 (Smoot, Ono, Ruscheinski, Wang, & Ideker, 2011).

Chapter Three: Results

3.1. Results

3.1.1. (MTT) Cytotoxicity results

The methanolic extract of *Adansonia digitata L.* plant extract was tested on L-929 fibroblast cell line. The results showed very low cytotoxicity levels for the tested concentrations (10 µg/mL, 20 µg/mL, 50 µg/mL, 100 µg/mL, 150 µg/mL, and 250 µg/mL). All experiments were conducted in triplicates with 5% Dimethyl Sulfoxide (DMSO) as a positive control. The negative control consisted of cells only with no extract. A third control consisted of 0.05% DMSO, since it was the solvent used to reconstitute the extracts. Cells were examined and photographed using Olympus Inverted System Microscope Model IX70 (**Figure 9**).

The results showed that the *Adansonia digitata L.* methanolic extract is safe and did not cause cell death on the L-929 fibroblast cell line (**Figure 10**), compared to the 5% DMSO which was very toxic. The observed IC₅₀ for the extract was 105.7 µg/mL.

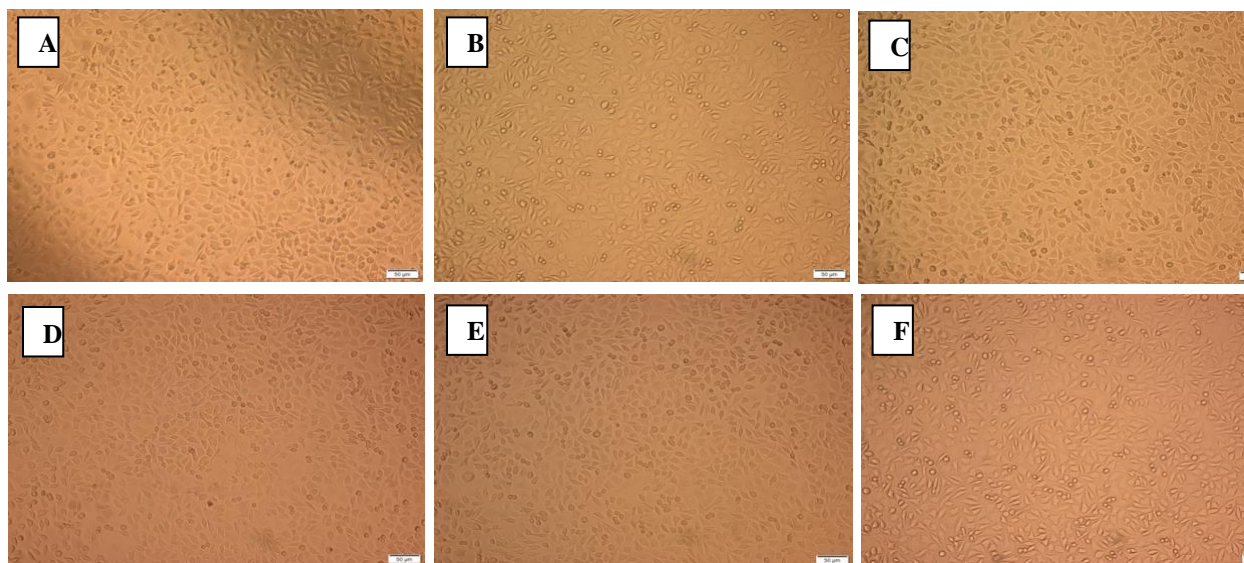
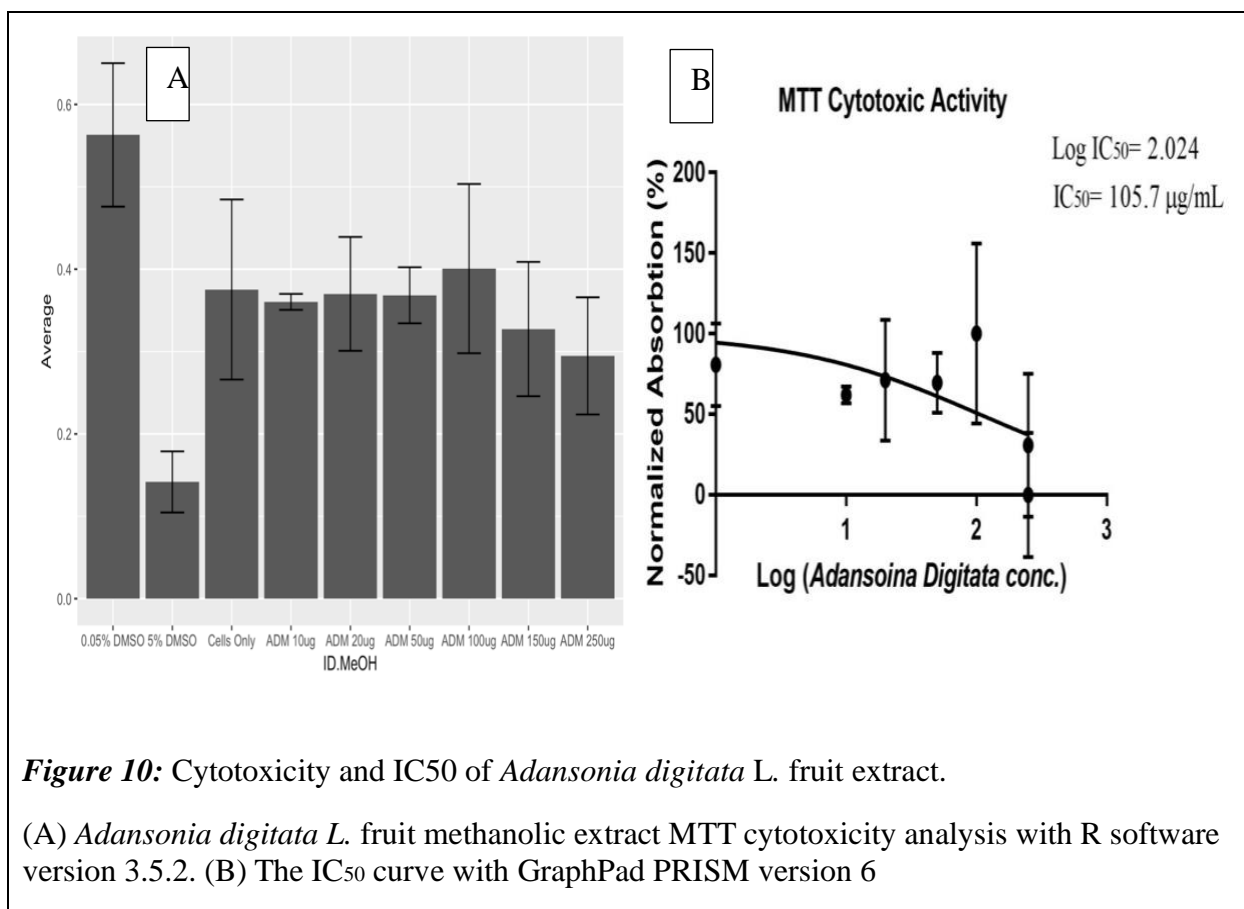


Figure 9: Cytotoxicity of *Adansonia digitata* L. fruit methanolic extract.

Photomicrographs showing L-929 cells after 24 h incubation with, under 10X magnification with a picture scale of 50 μm (Olympus Inverted System Microscope Model IX70)

- A. AD/MeOH 250 $\mu\text{g/mL}$
- B. AD/MeOH 150 $\mu\text{g/mL}$
- C. AD/MeOH 100 $\mu\text{g/mL}$

- D. AD/MeOH 50 $\mu\text{g/mL}$
- E. AD/MeOH 20 $\mu\text{g/mL}$
- F. AD/MeOH 10 $\mu\text{g/mL}$



3.1.2. Antioxidant Activity Using (DPPH) Results

The methanolic extract of *adansonia digitata L.* plant was examined to determine which extract has a higher antioxidant activity. The results showed that the methanolic extract showed high antioxidant activity. Six different concentrations of methanol extracts were tested (10 $\mu\text{g/mL}$, 20 $\mu\text{g/mL}$, 50 $\mu\text{g/mL}$, 100 $\mu\text{g/mL}$, 150 $\mu\text{g/mL}$, and 250 $\mu\text{g/mL}$). Ascorbic acid was used as a strong antioxidant agent.

The IC_{50} value, which is the sample concentration required for the 50% inhibition of the DPPH free radical, was measured by the DPPH absorbance versus the log of the concentrations curve as described previously (Garcia et al., 2012). The observed IC_{50} for the methanolic extract of the *Adansonia digitata L.* is 114.8 $\mu\text{g/mL}$ (**Figure 11**).

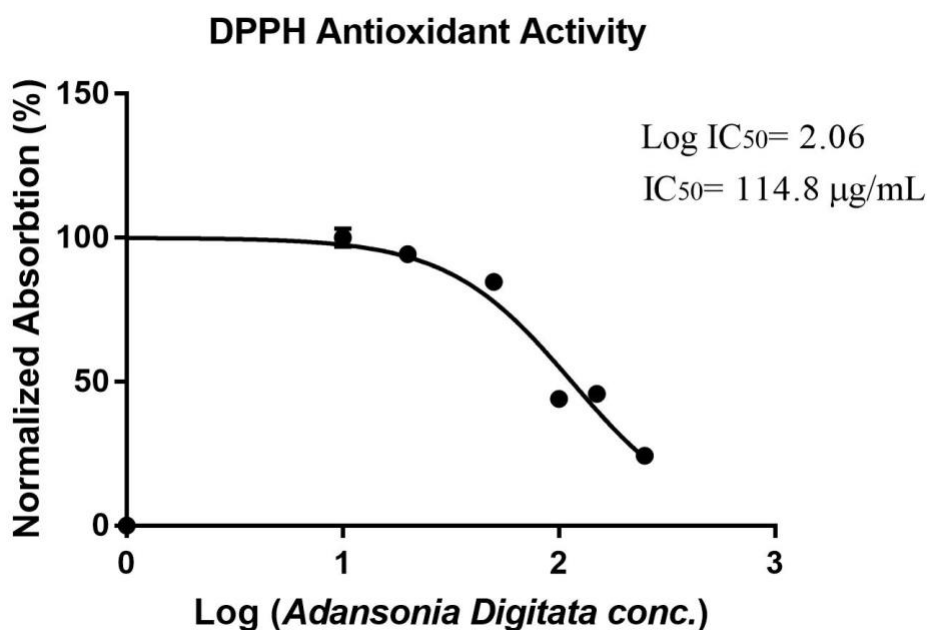


Figure 11: Antioxidant activity of *Adansonia digitata L.* methanolic extract.

DPPH antioxidant analysis IC_{50} curve with GraphPad PRISM version 6

3.1.3. UPLC-HRMS/MS profiling of *Adansonia digitata* L. fruit

Phytochemical constituents of *Adansonia digitata* L. fruits were analyzed via reversed-phase UPLC/PDA/ ESI-qTOF-MS and eluted with a gradient mobile phase, which permitted a holistic investigation of its analytes under both positive and negative ionization modes. Different ionization modes allowed for the detection of different metabolite classes through amending competitive ionization and suppression effects (M. A. Farag et al., 2016). Overlaid UPLC-MS base peak chromatogram (BPC) of *Adansonia digitata* L. fruit extract in both ionization modes is shown in **Figure 12**.

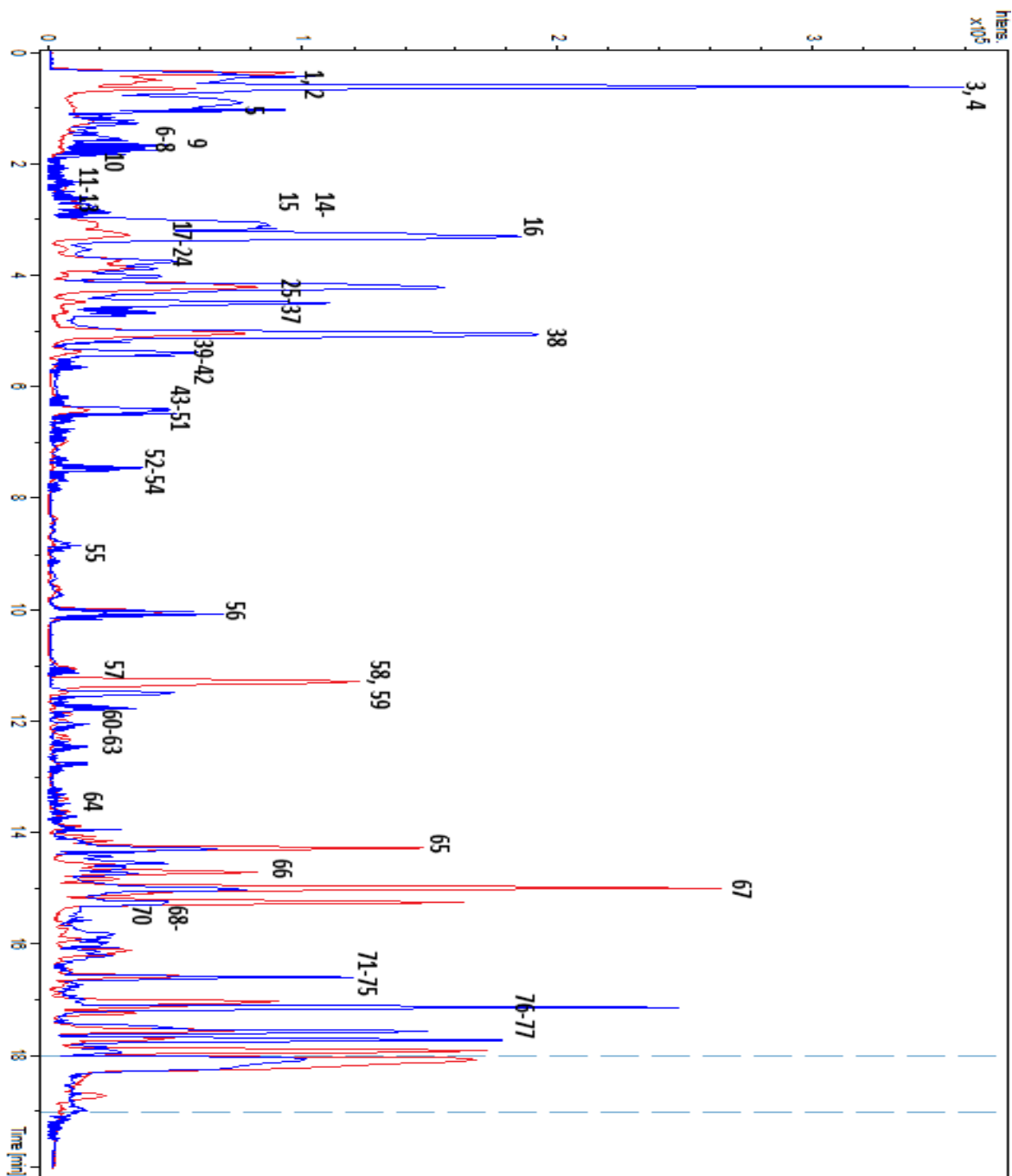


Figure 12: Overlaid BPC of *Adansonia digitata* L. fruit extract in both ionization modes.
Red and **blue** colors are corresponding to **positive** and **negative** ionization modes, respectively.

To aid in metabolite identification, two Molecular networks were generated for the positive and negative ionization modes individually through GNPS 2 platform to group peaks based on similar spectra. The spectral networks permitted the visual review of different metabolite classes, similarities among detected metabolites, and allowed isomers detection.

In contrast, the positive network exhibited 287 nodes grouped into 26 clusters and 109 single nodes, whereas the negative afforded 161 nodes in 13 clusters and 47 isolated nodes (**Figure 13**).

Molecular networking aided dereplication of *Adansonia digitata* L. metabolites

Metabolite dereplication was carried out based on their retention times, molecular formula, UV absorption maxima, and their fragmentation pattern in comparison to previously reported data. A total of 77 compounds were dereplicated belonging to different classes, including organic acids, sugars, alcohols, phenolics, coumarins, and fatty acids. Almost 50% of the identified metabolites (40 compounds) are reported for the first time in *Adansonia digitata* fruits (Table 1). Organic acids were detected in peaks 1, 2, 5, 6, & 13, while saccharides were detected in peaks 3, 4, 7, 14, 15, 25, 29, 31, & 32. Aromatics were detected in peaks 8, 12, 36, 57, 60, 61, 62, benzoic acid was detected in peaks 9 & 10, and the nitrogenous compound was detected in peaks 11 & 18.

In addition, flavan-3-ol was detected in peaks 16 & 28 and cinnamates were detected in peaks 17, 19, 21, 24, 26, 27, 30, 35, 37, 39, 40, 43, 50, 51, 52, 53, 63, & 64. Moreover, proanthocyanidin was detected in peaks 20, 22, 23, 33, 34, 38, & 45, furanochromones were detected only in peak 41, flavonol was detected in peaks 42, 44, 46, 47, 48, & 49, and flavonoid was detected only in peak 54. Terpenes were detected in peaks 55 & 56 and coumarin was detected in peaks 58, 59, 65, & 66. Finally, different fatty acids were detected in peaks 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, & 77.

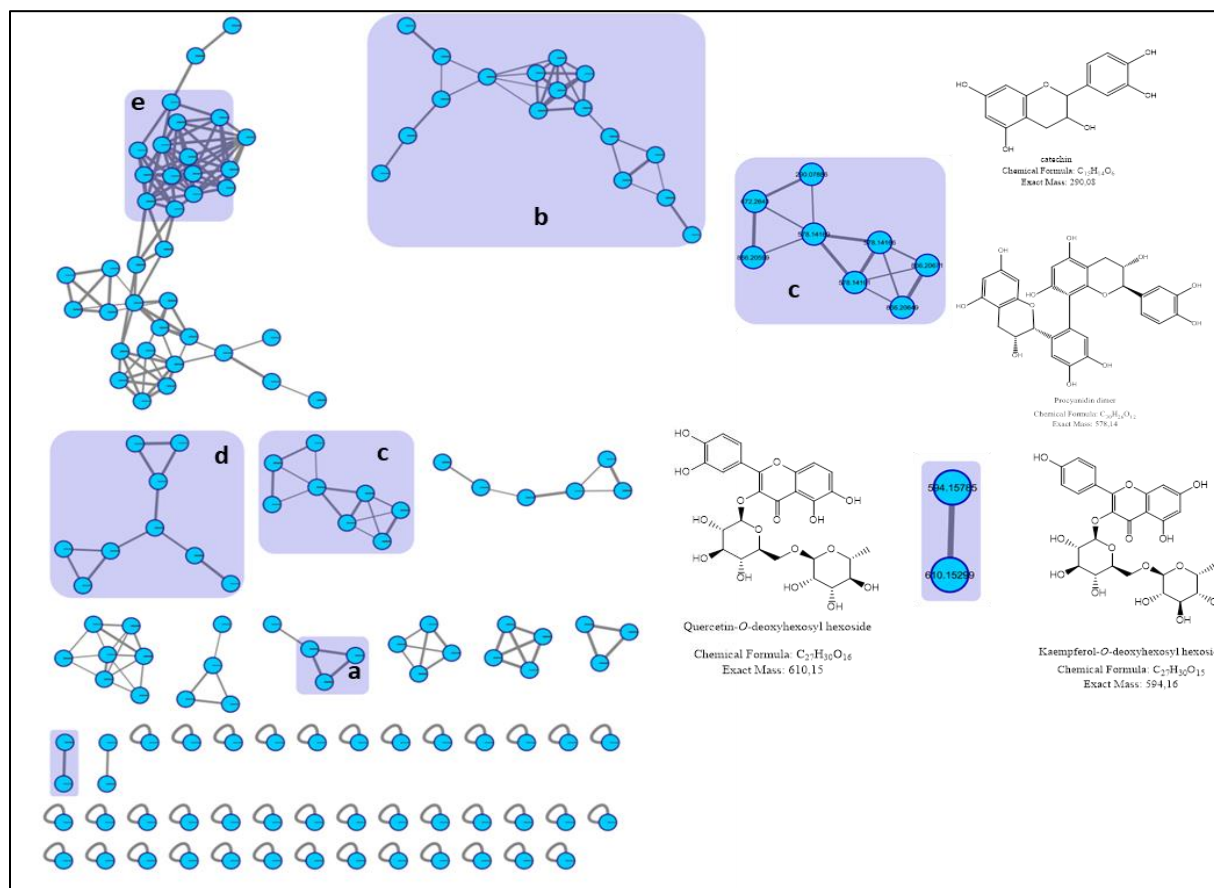


Figure 13: Negative molecular network of MS/MS data of *Adansonia digitata* L. fruit extract revealing main metabolite classes.

Table 2: The detected metabolites from the *Adansonia digitata* L. fruit using UHPLC-MS.

Peak	Rt min	Class	M-H	MS ₂	M+H	MS ₂	formula	Error ppm	Name	References
1.	0.48	Organic acid	191.0198	129, 111			C ₆ H ₈ O ₇	-0.14	Citric acid	(Braca et al., 2018)
2.	0.48	Organic acid	133.0142	115			C ₄ H ₆ O ₅	0.3	Malic acid	(Arold Jorel Tsetegho Sokeng et al., 2019)
3.	0.62	Saccharides	341.1091	179, 133	343.1234	163	C ₁₂ H ₂₂ O ₁₁	0.06	Disaccharide (sucrose)	(Arold Jorel Tsetegho Sokeng et al., 2019)
4.	0.66	Saccharides	371.1196	325, 193			C ₁₃ H ₂₄ O ₁₂	0.16	Unknown diasaccharide*	
5.	0.75	Organic acid	191.0200	111			C ₆ H ₈ O ₇	-1.87	Citric acid isomer	(Braca et al., 2018)
6.	1.0	Organic acid	205.0354	111			C ₇ H ₁₀ O ₇	-0.7	Citric acid methyl ester*	(Leenamajja Makila et al., 2016)
7.	1.3	Saccharides			385.1699	163, 145	C ₁₅ H ₂₈ O ₁₁	1.36	Butandiol- <i>O</i> - pentoside- <i>O</i> -hexoside*	(Hui Sun et al., 2016)
8.	1.39	Aromatics			127.0393	109	C ₆ H ₆ O ₃	-4.05	Pyrogallol*	(Hatice Tohma et al., 2017)
9.	1.53	Benzoic acid	167.0353	123			C ₈ H ₈ O ₄	2.05	Vanillic acid*	(Gruz, Novák, & Strnad, 2008)
10.	1.82	Benzoic acid	153.0193	109	155.0342	111	C ₇ H ₆ O ₄	-2.02	Dihydroxy benzoic acid* (protocatechuic acid)	(Ismail, Pu, Guo, Ma, & Liu, 2019)
11.	2.48	Nitrogenous compound			188.0701		C ₁₁ H ₉ NO ₂	2.51	Unknown	
12.	2.6	Aromatics	137.0243	109			C ₇ H ₆ O ₃	-2.6	Dihydroxybenzaldehyde*	(Lutter, Clark, Prenzler, & Scollary, 2007)
13.	2.84	Organic acid	175.0615	131, 157			C ₇ H ₁₂ O ₅	-1.8	Hydroxyglutaric acid*	(Peter Lorenz et al., 2012)
14.	3.32	Saccharides			369.1753	295, 163, 145, 133, 115	C ₁₅ H ₂₈ O ₁₀	0.38	Butanol- <i>O</i> -hexopyranosyl- <i>O</i> -deoxypyranoside*	(Fujimatu, Ishikawa, & Kitajima, 2003)
15.	3.32	Saccharides			295.1012	163, 145, 133	C ₁₁ H ₁₈ O ₉	2.55	Dihydroxy-methylene-butanoate- <i>O</i> -hexoside *	Tuliposide B (I. M. Abu-Reidah, Arráez-Román, Al-Nuri,

Peak	Rt min	Class	M-H	MS ₂	M+H	MS ₂	formula	Error ppm	Name	References
										Warad, & Segura-Carretero, 2019)
16.	3.39	Flavan-3-ol	289.0717	245, 203	291.0861	207, 139	C ₁₅ H ₁₄ O ₆	2.42	Catechin/ Epicatechin	(Braca et al., 2018; Ismail et al., 2019)
17.	3.5	Cinnamates	353.0891	191			C ₁₆ H ₁₈ O ₉	-2.1	<i>O</i> -Caffeoyl quinic acid ester (Clorogenic acid)	(Ismail et al., 2019)
18.	3.53	Nitrogenous compound	352.1034	191			C ₁₆ H ₁₉ NO ₈	0.85	Unknown*	
19.	3.72	Cinnamates	281.0660	193, 163, 145			C ₁₃ H ₁₄ O ₇	4.39	<i>O</i> -feruloyl lactic acid ester*	(S. Wu, Wilson, Chang, & Tian, 2019)
20.	3.79	Proanthocyanidin	577.1345	407, 289			C ₃₀ H ₂₆ O ₁₂	-0.98	Procyanidin dimer isomer	Xx (Braca et al., 2018; Ismail et al., 2019)
21.	3.8	Cinnamates	327.1082	165, 147			C ₁₅ H ₂₀ O ₈	2.8	Dihydrocoumaroyl <i>O</i> -hexoside (dihydromelilotoside)*	(Yang et al., 2007)
22.	3.9	Proanthocyanidin	577.0346	407, 289	579.1472	409, 301, 291	C ₃₀ H ₂₆ O ₁₂	4.25	Procyanidin dimer isomer I	(Braca et al., 2018; Ismail et al., 2019)
23.	3.91	Proanthocyanidin	865.1987			289	C ₄₅ H ₃₈ O ₁₈	0.2	Procyanidin trimer isomer I	(Braca et al., 2018; Ismail et al., 2019)
24.	3.98	Cinnamates	341.0893	179, 161			C ₁₅ H ₁₈ O ₉	4.5	Caffeoyl- <i>O</i> -hexoside	(X.-N. Li et al., 2017)
25.	4.03	Saccharides			295.1018	163, 145, 133	C ₁₁ H ₁₈ O ₉	1.88	Dihydroxy—methylenebutanoate- <i>O</i> -hexoside isomer	(I. M. Abu-Reidah et al., 2019)
26.	4.1	Cinnamates	487.1457	193			C ₂₁ H ₂₈ O ₁₃	-1.3	Feruloyl <i>O</i> -pentosyl - <i>O</i> -hexoside*	(Ibrahim M Abu-Reidah, Arráez-Román, Segura-Carretero, & Fernández-Gutiérrez, 2013)
27.	4.23	Cinnamates			165.0544	119, 107	C ₉ H ₈ O ₃	3.67	Coumaric acid	(Ismail et al., 2019)
28.	4.24	Flavan-3-ol	289.0716	245, 203	291.0860		C ₁₅ H ₁₄ O ₆	-0.27	Catechin/ Epicatechin	(Braca et al., 2018; Ismail et al., 2019)
29.	4.33	Saccharides			295.1018	163, 145, 133	C ₁₁ H ₁₈ O ₉	1.9	Dihydroxy-methylenebutanoic acid- <i>O</i> -hexoside	(I. M. Abu-Reidah et al., 2019)

Peak	Rt min	Class	M-H	MS ₂	M+H	MS ₂	formula	Error ppm	Name	References
30.	4.47	Cinnamates	325.0925	163			C ₁₅ H ₁₈ O ₈	-0.29	Coumaroyl- <i>O</i> -hexoside*	(Ibrahim M Abu-Reidah et al., 2013)
31.	4.52	Saccharides			383.1905	163, 145, 133, 155	C ₁₆ H ₃₀ O ₁₀	1.7	Methylbutyl - <i>O</i> -pentosyl- <i>O</i> -hexoside*	(I. M. Abu-Reidah et al., 2019)
32.	4.52	Saccharides			295.1008	163, 145, 133	C ₁₁ H ₁₈ O ₉	0.9	Dihydroxy-methylenebutanoic acid- <i>O</i> -hexoside	(I. M. Abu-Reidah et al., 2019)
33.	4.52	Proanthocyanidin	865.1992	577, 407, 289			C ₄₅ H ₃₈ O ₁₈	1.03	Procyanidin trimer Isomer II	Xx (Braca et al., 2018; Ismail et al., 2019)
34.	4.7	Proanthocyanidin	865.1994		867.1867	579, 289	C ₄₅ H ₃₈ O ₁₈	0.77	Procyanidin trimer isomer	Xx (Braca et al., 2018; Ismail et al., 2019)
35.	4.73	Cinnamates	367.1024	191, 173, 133			C ₁₇ H ₂₀ O ₉	-2.77	<i>O</i> -feruloyl quinic acid	Xx (Braca et al., 2018)
36.	4.8	Aromatics	461.1653	149			C ₂₀ H ₃₀ O ₁₂	4.5	Homovanillyl alcohol- <i>O</i> -pentosyl hexoside*	(Yang Zhang et al., 2012)
37.	4.8	Cinnamates	163.0401	119			C ₉ H ₈ O ₃	-0.1	Coumaric acid isomer	(Ismail et al., 2019)
38.	5.02	Proanthocyanidin	1153.2595	865, 577, 289			C ₆₀ H ₅₀ O ₂₄	-2.04	Procyanidin tetramer	(Ismail et al., 2019)
39.	5.1	Cinnamates			163.0391	135	C ₉ H ₈ O ₃	-0.5	Unknown*	
40.	5.49	Cinnamates	367.1031	191, 173, 133			C ₁₇ H ₂₀ O ₉	-0.52	<i>O</i> -feruloyl quinic acid isomer	(Braca et al., 2018)
41.	5.59	Furanochromones			409.1111	247	C ₁₉ H ₂₀ O ₁₀	4.37	Khellol- <i>O</i> -hexoside*	(Schuster, 1980)
42.	5.66	Flavonol	609.1457	301			C ₂₇ H ₃₀ O ₁₆	-0.65	Quercetin- <i>O</i> -deoxyhexosyl hexoside	(Braca et al., 2018)
43.	6.2	Cinnamates	325.0916	163			C ₁₅ H ₁₈ O ₈	-3.57	Coumaroyl- <i>O</i> -hexoside	(X.-N. Li et al., 2017)
44.	6.27	Flavonol	609.1467	301			C ₂₇ H ₃₀ O ₁₆	1.04	Quercetin- <i>O</i> -deoxyhexosyl hexoside	(Braca et al., 2018)
45.	6.43	Proanthocyanidin	577.1346	407, 289			C ₃₀ H ₂₆ O ₁₂	-0.93	Procyanidin dimer isomer III	(Braca et al., 2018; Ismail et al., 2019)
46.	6.46	Flavonol			449.1073	287	C ₂₁ H ₂₀ O ₁₁	1.17	Kaempferol- <i>O</i> -hexoside	(Braca et al., 2018)

Peak	Rt min	Class	M-H	MS ₂	M+H	MS ₂	formula	Error ppm	Name	References
47.	6.44	Flavonol	593.1505	447, 285	595.1617	449, 287	C ₂₇ H ₃₀ O ₁₅	1.04	Kaempferol- <i>O</i> -deoxyhexosyl hexoside*	(Ferrerres et al., 2017)
48.	6.47	Flavonol			287.0543	217, 151	C ₁₅ H ₁₀ O ₆	3.07	Kaempferol	(Braca et al., 2018)
49.	6.52	Flavonol	593.1505	447, 285			C ₂₇ H ₃₀ O ₁₅	-2.46	Kaempferol- <i>O</i> -deoxyhexosyl hexoside isomer*	(Ferrerres et al., 2017)
50.	6.62	Cinnamates	293.0671	163, 145, 119			C ₁₄ H ₁₄ O ₇	1.28	<i>O</i> -coumaroyl malic acid methyl ester*	(M. K. Lee et al., 2007)
51.	6.88	Cinnamates	355.1022	193			C ₁₆ H ₂₀ O ₉	-3.48	Feruloyl - <i>O</i> - hexoside	(X.-N. Li et al., 2017)
52.	7.3	Cinnamates	326.1032	206, 163, 145			C ₁₈ H ₁₇ NO ₅	-0.52	Coumaroyl - <i>N</i> -tyrosine*	(Clifford & Knight, 2004)
53.	7.5	Cinnamates	279.0870	163			C ₁₄ H ₁₆ O ₆	1.6	Unknown*	
54.	7.63	Flavonoid	447.09314	285			C ₂₁ H ₂₀ O ₁₁	0.34	Kaempferol- <i>O</i> -hexoside	(Braca et al., 2018)
55.	8.9	Terpenes	405.2123	225			C ₁₉ H ₃₄ O ₉	3.8	Megastigmane- <i>O</i> -hexoside*	(K. Singh, Dubey, Tripathi, Singh, & Saraf, 2014)
56.	10.1	Terpenes	507.2582	345			C ₂₇ H ₄₀ O ₉	0.9	Unknown	
57.	11.2	Aromatics	149.0605	122			C ₉ H ₁₀ O ₂	-1.68	Unknown	
58.	11.32	Coumarin			261.0753	230	C ₁₄ H ₁₂ O ₅	0.36	Khellin*	(Vanachayangkul, Butterweck, & Frye, 2009)
59.	11.38	Coumarin			231.0648	203, 216	C ₁₃ H ₁₀ O ₄	1.35	Visnagin*	(Vanachayangkul et al., 2009)
60.	12.29	Aromatics			137.0600	122, 109	C ₈ H ₈ O ₂	-2.2	Methoxy benzaldehyde	(Razanamaro et al., 2015)
61.	12.31	Aromatics			133.0651	105	C ₉ H ₈ O	2.94	Cinnamaldehyde	(Seukep et al., 2013)
62.	12.32	Aromatics			165.0907	137, 124	C ₁₀ H ₁₂ O ₂	1.63	Eugenol	(Razanamaro et al., 2015)
63.	13.6	Cinnamates			207.1017	165, 150	C ₁₂ H ₁₄ O ₃	-0.9	Unknown *	
64.	13.72	Cinnamates			211.0961	193, 181, 165	C ₁₁ H ₁₄ O ₄	1.55	Unknown*	

Peak	Rt min	Class	M-H	MS ₂	M+H	MS ₂	formula	Error ppm	Name	References
65.	14.97	Coumarin			411.1410 [M+Na]	329, 245, 227, 199	C ₂₁ H ₂₄ O ₇	0.81	Isovaleryl-acetyl khellactone (dihydrosamidin)*	(Olennikov, Fedorov, Kashchenko, Chirikova, & Vennos, 2019)
66.	14.98	Coumarin			329.1377	245, 227	C ₁₉ H ₂₀ O ₅	1.9	Decursin*	(H. J. Lee et al., 2009)
67.	15.1	Fatty acid	295.2274	277, 233			C ₁₈ H ₃₂ O ₃	1.3	Hydroxyoctadecadienoic acid*	(Napolitano, Cerulli, Pizza, & Piacente, 2018)
68.	15.1	Fatty acid	271.2278	253			C ₁₆ H ₃₂ O ₃	1.2	Hydroxyhexadecanoic acid*	(Feussner & Feussner, 2019)
69.	15.4	Fatty acid	269.2108	251		225	C ₁₆ H ₃₀ O ₃	5.1	Oxohexadecanoic acid*	(Nam et al., 2019)
70.	16.1	Fatty acid	295.2271	277, 233			C ₁₈ H ₃₂ O ₃	-3.1	Hydroxyoctadecadienoic acid*	(Liang et al., 2018)
71.	16.1	Fatty acid	297.2423				C ₁₈ H ₃₄ O ₃		Hydroxy- octedecaenoic acid*	(Napolitano et al., 2018)
72.	16.3	Fatty acid	311.2225	293, 275, 223			C ₁₈ H ₃₂ O ₄	1.1	Dihydroxy- octadecadienoic acid*	(Liang et al., 2018)
73.	16.6	Fatty acid	277.2171	259			C ₁₈ H ₃₀ O ₂	1.8	Octadecatrienoic acid*	(Osman, 2004)
74.	16.93	Fatty acid	355.3216	309			C ₂₂ H ₄₄ O ₃	0.22	Hydroxydocosanoic acid*	(Feussner & Feussner, 2019)
75.	17.1	Fatty acid	279.2331	261			C ₁₈ H ₃₂ O ₂	-0.6	Octadecadienoic acid (linoleic acid)	(Osman, 2004)
76.	17.5	Fatty acid	279.2331	261			C ₁₈ H ₃₂ O ₂	-1.8	Octadecadienoic acid (linoleic acid)	(Osman, 2004)
77.	17.6	Fatty acid	255.2333				C ₁₆ H ₃₂ O ₂	-0.8	Hexadecanoic acid (palmitic acid)	(Razafimamonjison et al., 2017)

* Metabolites detected for the first time from *Adansonia digitate* L. fruit.

3.1.4. Biochemical Assessment

Fasting blood glucose (FBG) was assessed to determine the effect of the *Adansonia digitata* L. fruit extract on the glucose level of diabetic rats. Untreated diabetic rats showed the highest FBG level. The 150 mg/kg of the fruit extract twice/ week was able to keep the blood glucose levels of the diabetic rats within the normal range (**Figure 14**). Unexpectedly, the high dose treatment (300 mg/kg) did not reduce the blood glucose levels, and most of the rats were above 250 mg/dL. Diabetic rats in all experimental groups gained weight throughout the experiment compared to normal healthy rats and normal rats receiving the extract.

Liver functions (ALT, AST, and Alkaline phosphatase), renal functions (Creatinine, Blood Urea Nitrogen, and uric acid), and lipid profile (Cholesterol, Triglycerides, and HDL) were tested to examine the effect of the *Adansonia digitata* L. fruit extract (**Figure 15 and Table 3**).

Liver functions and Lipid Profile

The alkaline phosphatase (ALK) was reduced to near normal (166 ± 50 U/L) in treated groups, compared to the untreated diabetic rats (559 ± 328 U/L). The ALT liver enzyme was slightly higher in the tested groups compared to the normal rats (28 ± 12.02 U/L). Untreated diabetic, metformin-treated, and *Adansonia digitata* L. extract treated rats showed high levels of ALT, compared to normal. In other words, *Adansonia digitata* L. extract did not have an effect on the ALT enzyme in the tested rats. AST liver enzyme was elevated in the untreated diabetic group compared to normal. Metformin and *Adansonia digitata* L. treated groups showed a decrease in the AST levels. *Adansonia digitata* L. administration in normal rats did not alter the ALK, ALT or AST levels.

The lipid profile was assessed by the total cholesterol, triglycerides, and high-density lipoprotein (HDL) levels. Cholesterol levels increased in diabetic rats compared to normal rats, which had cholesterol levels in range (49.5 ± 7.77 mg/dL). *Adansonia digitata* L. fruit extract significantly (p -value < 0.05) reduced the cholesterol levels in both low and high doses treatment groups (Cholesterol was 46.2 ± 8.31 mg/dL in 150 mg/kg treatments and in 300 mg/kg treatment it was 79.33 ± 9.86 mg/dL).

Triglyceride levels were higher in metformin-treated rats compared to normal and untreated diabetic rats. *Adansonia digitata* L. fruit extract reduced TG levels in both high and low treatment groups compared to metformin-treated. HDL increased in all experimental groups compared to the normal rats. However, *Adansonia digitata* L. fruit extract administration in normal rats caused an elevation in Cholesterol, TG, and HDL.

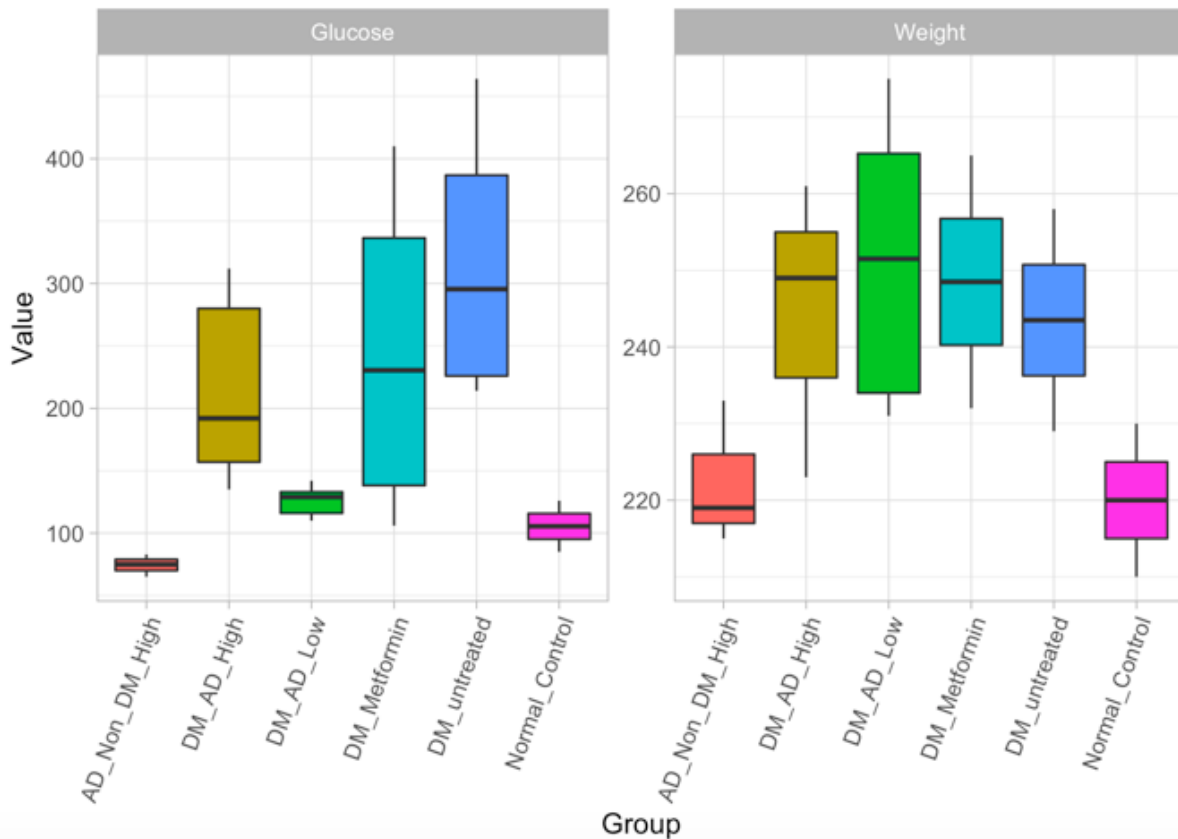


Figure 14: Fasting blood glucose (FBG) and body weight of rats.

*The fasting blood glucose result was statistically significant p-value (≤ 0.05)

Renal Functions

The results of the kidney function tests revealed that Creatinine levels were high in the metformin-treated diabetic group, as well as in the *Adansonia digitata* L. high dose (300 mg/kg) treated group. However, in the low dose of *A. digitata* (150 mg/kg) and the untreated diabetic groups, Creatinine levels were within the normal range.

Urea level was elevated in the untreated diabetic rats compared to normal control. Metformin treatment caused an insignificant reduction in the urea levels. Urea levels in the *Adansonia digitata* L. treated groups were 36.8 ± 7.49 mg/dL and 50.33 ± 11.15 mg/dL, in the low and high dose groups, respectively.

Uric acid levels increased in diabetic rats. Metformin treatment caused a significant reduction in uric acid levels. All rats receiving *Adansonia digitata* L. extract showed high levels of uric acid compared to normal healthy rats.

Table 3: Comparison of the six animal groups' blood biochemical analysis results compared to the normal healthy rats.

	Normal Healthy	DM untreated	DM + Metformin	DM + 150 mg/kg AD	Healthy + 300 mg/kg AD	DM + 300 mg/kg AD
ALT (U/L)	28 ± 12.02	37.5 ± 5.19	34 ± 15.87	32.6±11.2	34.66 ± 9.8	35.2 ±8.40
AST (U/L)	99 ± 16.97	115.75 ± 30.20	82.75 ± 33.52	83.2±13.66	94.66±19.5	69.8±12.47
Alkaline phosphatase (U/L)	166 ± 50	559 ± 328	200 ± 76	215 ± 71	364 ± 35	242 ± 56
Creatinine (mg/dL)	0.585 ± 0.049	0.49 ± 0.027	0.69 ± 0.11	0.54±0.099	0.55 ± 0.125	0.63 ± 0.11
Urea (mg/dL)	37 ± 1.414	71.75 ± 13.94	64.75 ± 29.41	36.8 ± 7.49	50.33±11.15	34.2 ± 7.19
Cholesterol (mg/dL)	49.5 ± 7.77	87.5 ± 22.98	73.5 ± 24.90	46.2 ± 8.31	79.33±9.86	55 ± 15.16
TG (mg/dL)	53.5 ± 3.53	84.75 ± 26.11	134.5 ± 73.66	87.8±25.03	116.33±66.5	110.6±26.8
HDL (mg/dL)	32 ± 12	41 ± 13	42 ± 17	46 ± 7.3	43 ± 11	47 ± 8.1
Uric Acid (mg/dL)	0.95 ± 0.77	2.22 ± 0.35	0.74 ± 0.30	2.32 ± 1.47	2.16 ± 0.61	2.30 ±1.02

*The results are expressed as the mean ± standard error (n=6).

The ALK, cholesterol, TG and urea biochemical parameters were statistically significant, p-value (≤ 0.05), the test was performed using R software version 3.5.2

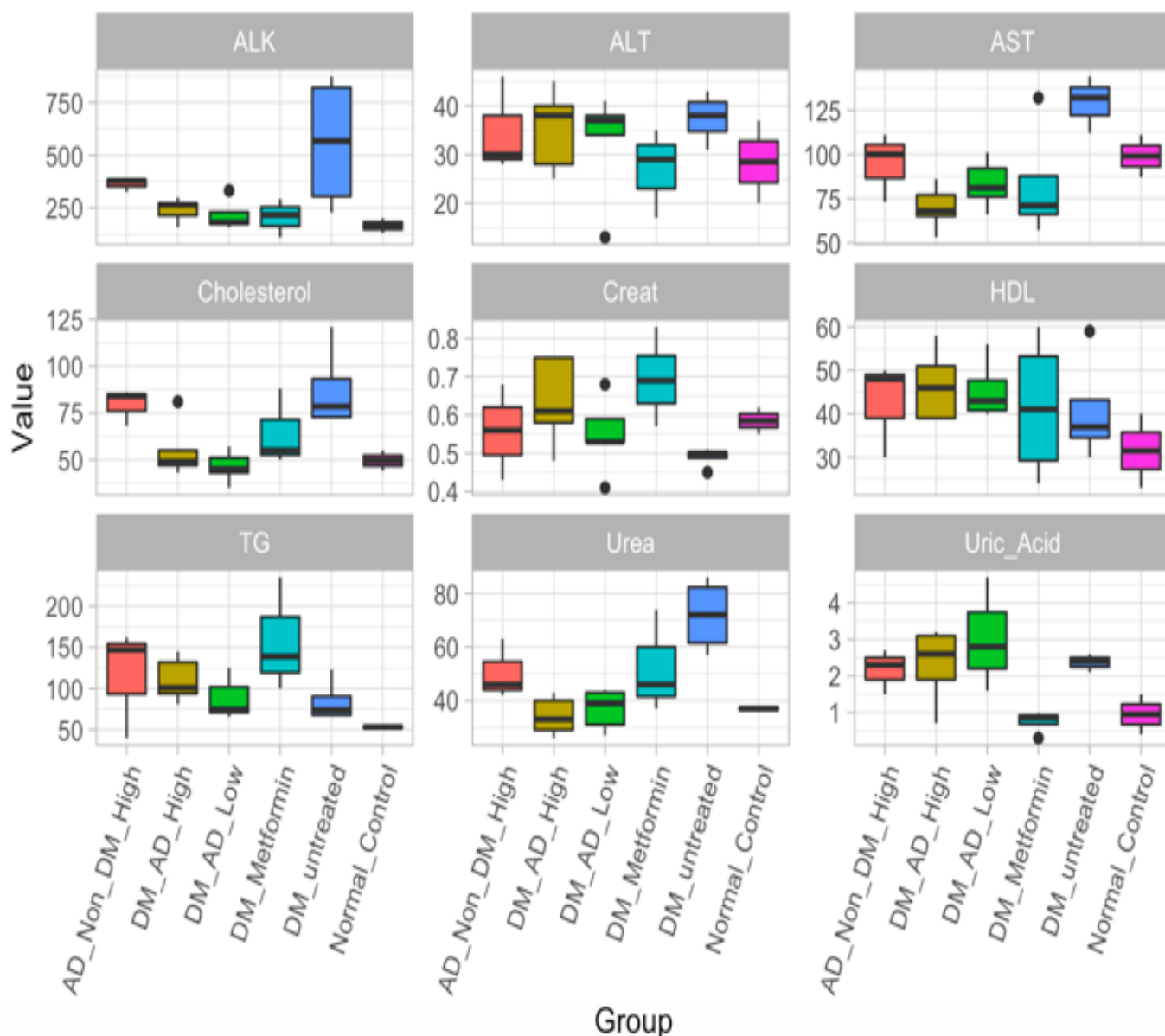


Figure 15: Biochemical analyses of the lipid profile, liver, and kidney functions.

Results of biochemical analyses in the experimental groups. The groups are normal healthy rats (Normal_Control), diabetic rats without any treatment (DM_untreated), diabetic rats administrated metformin orally (DM_Metformin), diabetic rats i.p. injection with a low dose, which is 150 mg/kg, of *Adansonia digitata* L. (DM_AD_Low), diabetic rats i.p. injection with high dose, which is 300 mg/kg, of *Adansonia digitata* L. (DM_AD_High), and healthy rats i.p. injection with high dose of *Adansonia digitata* L. (AD_Non_DM_High). Figure generated with R software version 3.5.2.

3.1.5. Histopathological Analysis

Pancreas, kidney, and liver tissues from each group (n = 9 slides/rat) were stained with hematoxylin and eosin (H&E) and examined on Olympus BX43 light microscope attached with a digital camera (DP27) and CellSens dimensions software.

Histopathology of the Pancreas

Pancreatic tissue from the normal control and normal rats that received a high dose of *Adansonia digitata* L. revealed a normal histological structure. The untreated diabetic group showed shrinkage of pancreatic islets, severe necrosis of β -cells, and reduction in islets cell number. Similar changes were observed in the metformin-treated diabetic group. However, rats from *Adansonia digitata* L. treated diabetic groups (both low dose, 150 mg/kg and high dose, 300 mg/kg) showed an increase in islets cell numbers with normal appearance of most cells (**Figure 16**).

Histological scoring of the pancreatic islets diameter (**Figure 17**), revealed that the largest diameter was recorded in normal controls and non-diabetic rats that received a high dose of *Adansonia digitata* L. While the smallest diameter was recorded in untreated diabetic rats and diabetic rats treated with metformin, respectively.

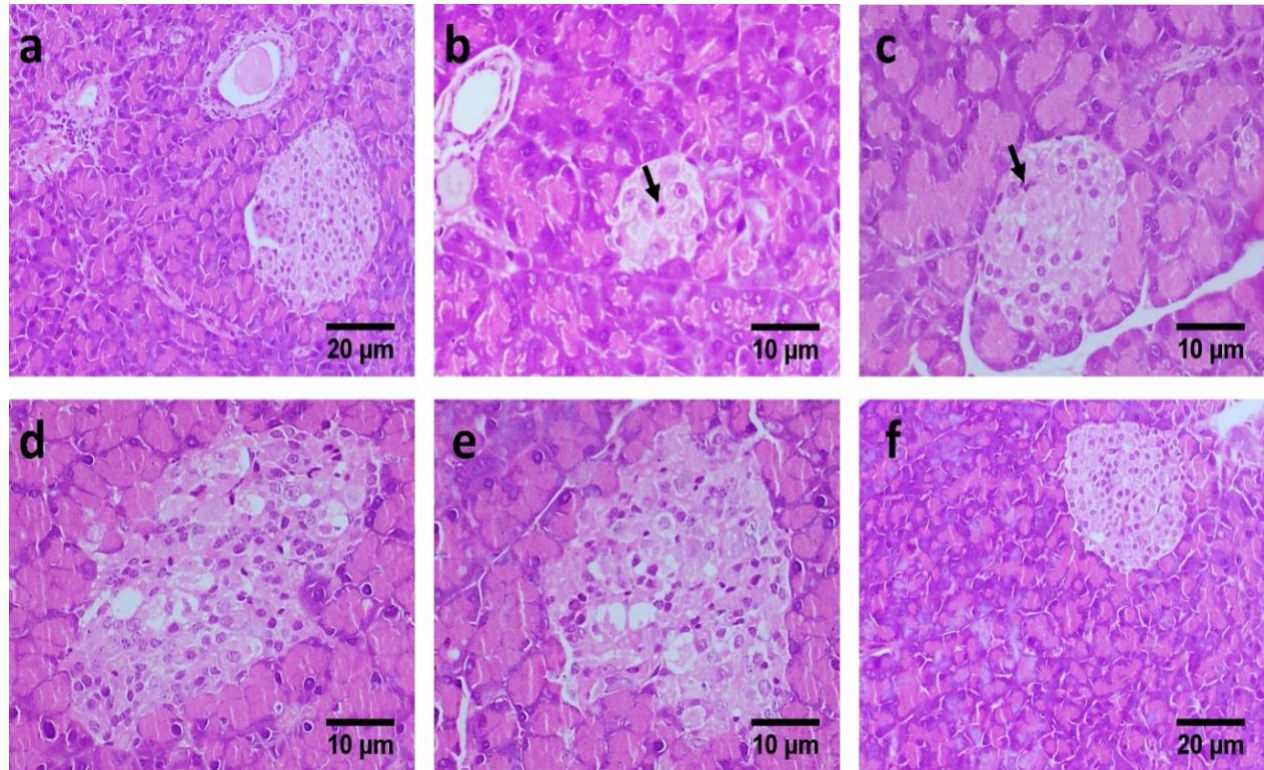


Figure 16: Histopathology of the Pancreas.

Photomicrographs of the pancreas stained by (H&E). (a) Healthy control group showing a normal histological structure of the pancreas with a regular outline of pancreatic islets. (b) DM_Untreated group showing a reduction in both the size of pancreatic islets and the number of the islets cells with necrobiotic changes (arrow). (c) DM_metformin group showing a similar pathological alteration detected in the untreated diabetic group. (d) DM_AD_Low group showing a normal appearance of islets with an increase in both islet diameter and number of islet cells. (e) DM_AD_High group showing improvement in the histological structure and number of the cells of the islets compared to the control group. (f) AD_Non_DM_High group showing a normal histological structure of both endocrine and exocrine portion.

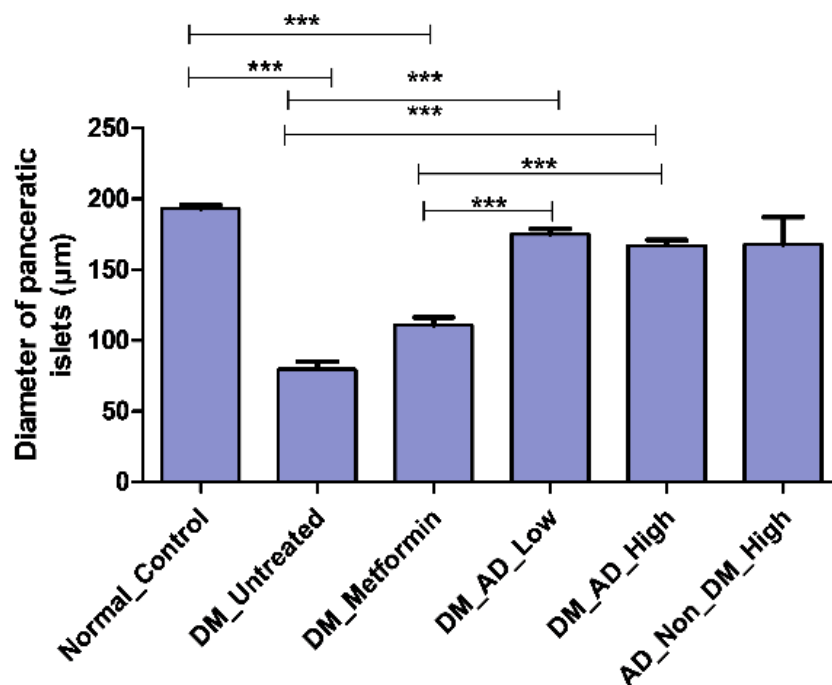


Figure 17: Histological scoring of the diameter of the pancreatic islets (µm).

Data expressed as mean \pm SD. ***Significant difference between groups, $P < 0.001$

Histopathology of the Liver

Histological examination of liver sections of the healthy control rats (**Figure 18**) and non-diabetic rats receiving the high dose of *Adansonia digitata* L showed normal histological structure. The untreated diabetic group showed necrosis of hepatocytes adjacent to the central vein associated with inflammatory cell infiltration. In untreated diabetic rats, inflammation, necrosis, and vacuolar degeneration were detected. Liver sections from metformin-treated diabetic rats showed similar histopathological changes to the untreated diabetic group. Treatment with *Adansonia digitata* L. in both low (150 mg/kg) and high (300 mg/kg) doses, improved the histological structure of the liver. Histological score of liver lesions (**Figure 19**), showed a significant increase in tissue damage in both untreated diabetic and the metformin-treated diabetic groups (** $P < 0.01$, *** $P < 0.001$).

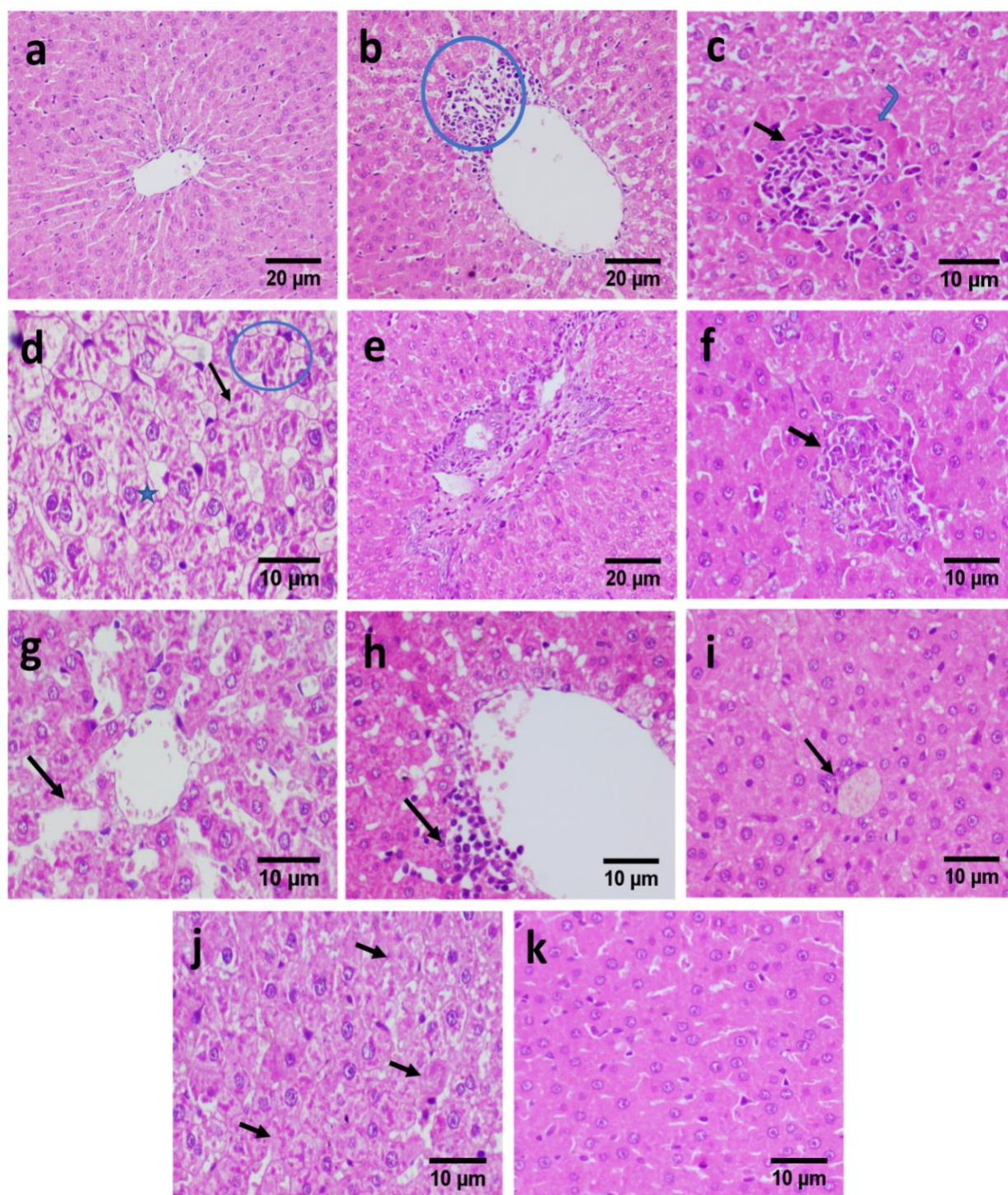


Figure 18: Histopathology of the Liver.

Photomicrographs of liver sections stained by (H&E). (a) Control group showing normal histological structure. (b) DM- untreated group showing necrosis of hepatocytes adjacent to central

vein associated with infiltration of inflammatory cells (**circle**), (c) focal area of necrosis associated with inflammatory cell infiltration (black arrow) as well as coagulative necrosis in the hepatocytes (**blue curved arrow**), (d) vacuolar degeneration (**asterisk**) and karyorrhexis (fragmentation of nucleus) in some hepatocyte (**circle**) and (e) inflammation of the portal area. (f) DM_Metformin group showing a focal area of necrosis associated with inflammatory cells infiltration (arrow), (g) vacuolar degeneration with dilation in sinusoid (arrow), and (h) infiltration of the inflammatory cell adjacent to the central vein (arrow). (i) DM_AD_Low group showing the restoration of cellular structure with minimal infiltration of inflammatory cells (arrow). (j) DM_AD_High group showing slight granulation of hepatic cytoplasm with necrosis of minute cluster of hepatocytes (spotty necrosis), (arrows). (k) AD_Non_DM_High group showing a normal cellular structure of hepatocytes.

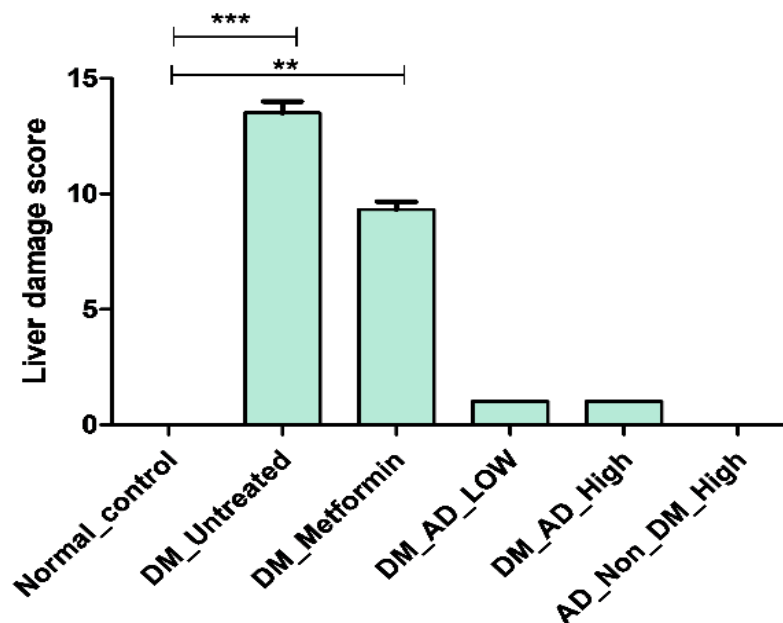


Figure 19: Histological damage score in liver tissue.

Data presented as median \pm SD. **P < 0.01, ***P < 0.001.

Histopathology of the Kidney

Normal histology of the kidney was seen in the healthy control group (**Figure 20**). Vacuolar degeneration, necrobiotic changes, and interstitial nephritis were observed in the untreated diabetic group. Kidney sections from the metformin-treated group showed degenerative and necrobiotic changes in renal tubules. Treatment of diabetic rats with low (150 mg/kg) or high (300 mg/kg) dose of *Adansonia digitata* L. improved the histological structure of the kidney, and no inflammation or necrosis was observed. No histological changes were detected in normal rats receiving a high dose of *Adansonia digitata* L. fruit methanolic extract.

Scoring of kidney damage is shown in (**Fig. 21**). A significant increase in renal damage score was detected in the untreated diabetic group and metformin-treated diabetic rats when compared with the control group. There was no significant difference between diabetic rats treated with either low or high doses of *Adansonia digitata* L. fruit methanolic extract (** $P < 0.01$, *** $P < 0.001$).

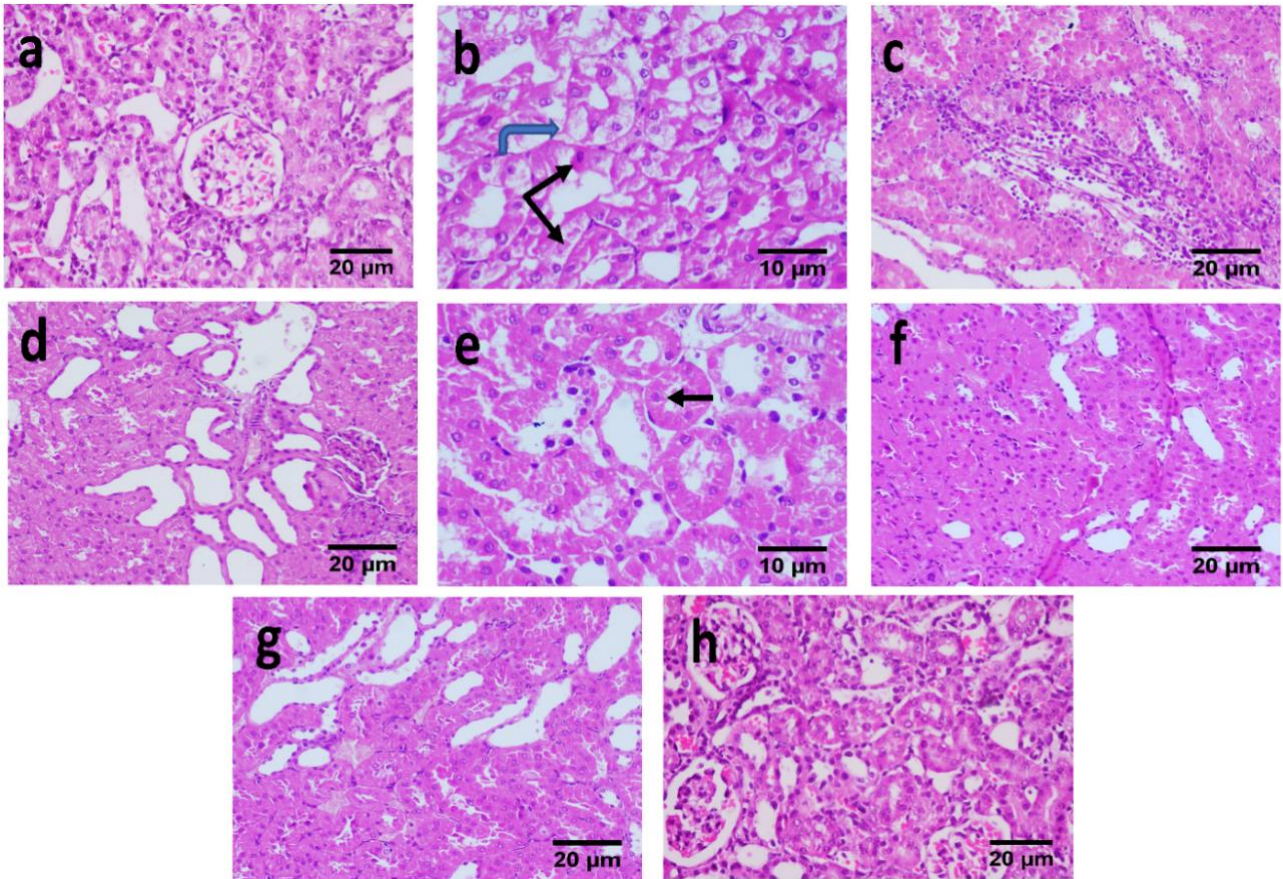


Figure 20: Histopathology of the Kidney.

Photomicrographs of **kidney** sections stained by (H&E). (a) Control group showing normal histological structure. (b) DM_Untrated group showing vacuolar degeneration of tubular epithelium (blue curved arrow) with necrobiotic changes (black arrows) and (c) Interstitial nephritis which characterized by infiltration of inflammatory cells between renal tubules. (d) DM_Metformin group showing cloudy swelling of tubular epithelium associated with dilation of some renal tubules and (e) necrobiotic changes in the epithelium lining renal tubules (arrow). (f) DM_AD_Low group showing cloudy swelling. (g) DM_AD_High group showing a similar change in the rats treated with a low dose of *Adansonia digitata* L. h) AD_Non_DM_High group showing normal cellular architecture.

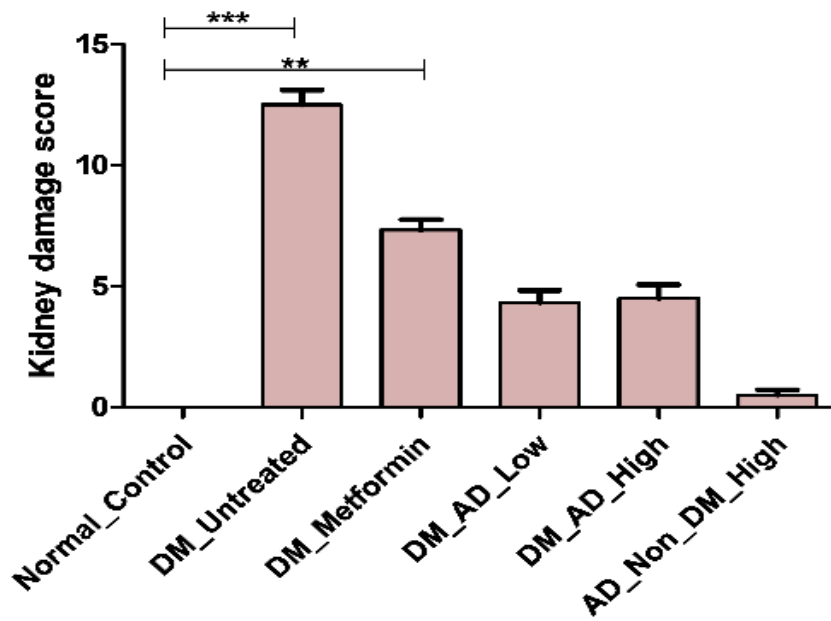


Figure 21: Histological damage score in kidney tissue.

Data presented as median \pm SD. ** $P < 0.01$, *** $P < 0.001$.

Chapter Four: Discussion

Diabetes Mellitus (DM) is considered one of the fast growing diseases around the world (Soltesz, Patterson, & Dahlquist, 2007). The search for an effective anti-diabetic medication is never stopping. In this study, we assessed the anti-diabetic potential of *Adansonia digitata* L. methanolic fruit extract. Baobab or *Adansonia digitata* L. is a traditional folk medicinal plant common to Africa. In 2008, the Baobabs plant obtained EU regulatory approval as a food additive (Agency, 2002).

One of the main challenges when using a new drug or plant extract is to assess its toxicity. The MTT assay is a quantitative, sensitive, and reliable colorimetric assay (Riss et al., 2016). The *Adansonia digitata* L. plant extract showed an IC₅₀ equals 105.7 µg/mL, which indicates a substantial safety margin for the tested plant extract.

The toxicity of The *Adansonia digitata* L. plant extract was also assessed *in vivo* by administering 300 mg/kg (the highest dose in the current study) to normal healthy rats. *Adansonia digitata* L. administration in normal rats did not alter the body weight, serum glucose, or liver functions (ALK, ALT, or AST levels). Urea and Creatinine were not affected in normal rats. However, all rats receiving *Adansonia digitata* L. extract showed high levels of uric acid compared to normal healthy rats, which indicates a possible side effect of using the extract in the long term. No histological changes were detected in normal rats receiving high doses of *Adansonia digitata* L. fruit extract. The liver, kidneys, and pancreas appeared normal with healthy cellular architecture, confirming the biochemical results that show the safety of the extract.

One of the advantages of using natural plant extracts is their multi-potential. In addition to its anti-diabetic activity, *Adansonia* possesses natural antioxidant properties. These natural antioxidants are essential because of their health maintenance effects, as well as their anti-aging effect (J. F. Weiss & Landauer, 2003). The observed DPPH IC₅₀ for *Adansonia digitata* L. methanolic fruit extract in this study was 114.8 µg/mL. The low IC₅₀ value indicates higher antioxidant activity. The current finding is consistent with previous findings that examined the methanolic extract of *Adansonia digitata* L. Stem, leaf, fruit pulp, and seed (S. Vertuani et al., 2011).

The *Adansonia digitata* L. fruit methanolic extract was tested *in-vivo* for its anti-diabetic potential. At low doses, the extract reduced fasting blood glucose (FBG) levels, and surprisingly, higher doses were not as effective. The current findings are similar to those reported by Gwarzo and Bako (2013). They found that *Adansonia digitata* L. fruit pulp (300 mg/kg) administered orally reduced blood glucose levels in Alloxan diabetic Wistar rats.

Histological examination of the pancreatic tissue confirmed the diabetic state of the rats. Untreated diabetic rats showed shrinkage of pancreatic islets and severe necrosis of β -cells. Treatment with *Adansonia digitata* L. (both low, 150 mg/kg, and high dose, 300 mg/kg) improved the islet cell numbers with the healthy appearance of most cells. Metformin, on the other hand, did not cause a histological improvement of the pancreatic tissue.

The *Adansonia digitata* L. fruit extract did not affect body weight. All diabetic rats gained weight compared to healthy rats. To the best of our knowledge, no reported studies are discussing the effect of the fruit pulp extract on the weight in animal models of diabetes.

Liver enzymes are biochemical markers used to indicate hepatocyte necrosis and inflammation (Reichling & Kaplan, 1988). The alkaline phosphatase (ALK) was reduced to normal levels in treated groups compared to the untreated diabetic rats. The ALT liver enzyme was slightly higher in the tested groups compared to the normal rats. *Adansonia digitata* L. extract did not affect the ALT enzyme in the tested rats. *Adansonia digitata* L. and Metformin decreased the AST levels. *Adansonia digitata* L. administration in normal rats did not alter the ALK, ALT or AST levels.

The current results are consistent with those previously reported that stem bark extract of the *Adansonia digitata* L. reduced the ALK enzyme following oral administration of 200 mg/kg and 400 mg/kg (Adeoye & Bewaji, 2018).

The biochemical results are consistent with the histological findings. Both revealed that the injection of *Adansonia digitata* L. in both low (150 mg/kg) and high (300 mg/kg) doses improved the histological structure of the liver in diabetic rats.

The lipid profile was assessed by the total cholesterol, triglycerides, and high-density lipoprotein (HDL) levels. Cholesterol levels increased in diabetic rats compared to normal rats. *Adansonia digitata* L. extract significantly reduced the cholesterol and TG levels in both low and high dose

treatment groups. HDL increased in all experimental groups compared to the normal rats. However, *Adansonia digitata L.* fruit extract administration in normal rats caused an elevation in Cholesterol, TG, and HDL.

These results are contradictory with previous reports regarding the used doses Bako et al. (2014) that the fruit extract of *Adansonia digitata L.* reduced the cholesterol, triglycerides, and low-density lipoprotein LDL in alloxan diabetic Wistar rat model with the high dose 300 mg/kg. However, we are reporting that the low dose which is 150 mg/kg was more effective than the reported 300 mg/kg. On the other hand, the fruit crude extract in the current study and in previously reported findings was able to enhance the high-density lipoprotein (HDL), which is considered good cholesterol (Jomard & Osto, 2020), which is associated with a decrease in coronary risk (Bako et al., 2014; Luka & Tijjani, 2013).

Three biochemical parameters assessed kidney functions; Creatinine, Urea, and uric acid (Suliman, Osman, Abdoon, Saad, & Khalid, 2020). Creatinine levels were high in the metformin-treated diabetic group, and in the *Adansonia digitata L.* high dose (300 mg/kg) treated group indicating impairment of the renal function (Amartey, Nsiah, & Mensah, 2015). However, following the low dose of *Adansonia* (150 mg/kg), Creatinine levels were within the normal range.

Urea levels increased in the untreated diabetic rats compared to normal control. Metformin treatment and *Adansonia digitata L.* treatment reduced Urea levels indicating recovery of the renal function after treatment.

Uric acid levels increased in diabetic rats. Metformin treatment caused a significant reduction in uric acid levels. However, all rats receiving *Adansonia digitata L.* extract showed high levels of uric acid compared to normal healthy rats. This indicates a possible side effect of the use of *Adansonia digitata L.* extract.

The biochemical results were consistent with the histological examination, which showed that the treatment of diabetic rats with *Adansonia digitata L.* reversed the histological damage of the kidneys in diabetic rats.

Metabolites characterization

Compared to targeted analysis, metabolomics targets a large subset of metabolites within the biological system (Cevallos-Cevallos, Reyes-De-Corcuera, Etxeberria, Danyluk, & Rodrick, 2009). Liquid chromatography-mass spectrometry (LC-MS) is an advanced technique for the assessment of residues. LC has a universal separation mechanism which can be used for profiling secondary metabolites (Masia, Suarez-Varela, Llopis-Gonzalez, & Pico, 2016)

Recently, the chromatographic performance has been shifted to the ultra-performance liquid chromatography (UPLC), owing to its better separation efficiency (Jansson, Pihlström, Österdahl, & Markides, 2004; Klein & Alder, 2003).

Recent studies showed that the phytochemical analyses for the methanolic extract showed the presence of glycosides, flavonoids, tannins, saponins, terpenoids, and steroids. In addition, this methanolic extract from the fruit pulp showed anti-diabetic effects in Alloxan induced diabetes models (Gwarzo & Bako, 2013).

To characterize metabolites in the fruit extract, UHPLC-MS was employed for metabolites profiling. The UHPLC/MS results in **Table 2** showed several chemical classes.

Organic acids such as Citric acid and its isomer were detected in peaks 1 & 5 with [M-H]⁻ at m/z 191.0198, C₆H₈O₇, and characteristic fragment at m/z 111. The negative molecular network revealed a directly correlated ion at peak 6 [M-H]⁻ 205.0354, C₇H₁₀O₇, with a mass difference of 14 Da (CH₂). It was annotated as citric acid methyl ester (L. Makila et al., 2016). Other organic acids included peak 2 at m/z [M-H]⁻ 133.0142, C₄H₆O₅ annotated as malic acid and peak 13 at m/z with [M-H]⁻ 175.0615, C₇H₁₂O₅ and daughter ions at m/z 157 and 131 for the loss of water and CO₂ annotated as hydroxyglutaric acid (P. Lorenz et al., 2012).

Saccharides and their derivatives from which peaks 3 & 4 corresponded to two disaccharides, where 3 exhibited molecular ion [M-H]⁻ at m/z 341.1107, C₁₂H₂₂O₁₁ and were assigned as sucrose

previously identified in *Adansonia* fruits (A. J. T. Sokeng et al., 2019) while, **4**, unknown disaccharide had [M-H]⁻ 371.1196, C₁₃H₂₄O₁₂.

Butanediol-pentosyl-hexoside, peak **7** in the positive mode, showed a molecular ion at m/z 385.1699 [M+H]⁺, C₁₅H₂₈O₁₁ (H. Sun et al., 2016). Similarly, peak **14** showed a molecular ion at m/z 369.1753 [M+H]⁺, C₁₅H₂₈O₁₀, and a fragment at m/z 295 for the loss of C₄H₉O was assigned as butanol-pentosyl-hexoside (Fujimatu et al., 2003).

Additionally, the positive molecular network revealed the presence of 4 isomers of acylated hexoses; peaks **15**, **25**, **29** & **32**, which shared a molecular ion at m/z 295.1018 [M+H]⁺ and corresponded to the elemental composition of C₁₁H₁₈O₉. Manual inspection of their MS₂ spectra showed the occurrence of a common fragment ion at m/z 133 resulting from the cleavage of a hexose moiety. Thus they were assigned as dihydroxy-methylenebutanoic acid-*O*-hexoside isomers (I. M. Abu-Reidah et al., 2019), which are well recognized for their antibacterial properties (Shigetomi, Shoji, Mitsuhashi, & Ubukata, 2010). While peak **31**, with [M+H]⁺ at m/z 383.1905, C₁₆H₃₀O₁₀ was found within the same cluster and was annotated as methylbutyl -*O*-pentosyl-hexoside (I. M. Abu-Reidah et al., 2019).

Several aromatic metabolites were detected sparsely in both ionization modes and revealed themselves mostly as single nodes within both spectral networks. The first eluted at peak **8** [M+H]⁺ at m/z 127.0393, C₆H₆O₃ with a fragment at m/z 109 accounting for the loss of H₂O and annotated as pyrogallol (H. Tohma et al., 2017). Following at peak **12** was dihydroxybenzaldehyde with [M-H]⁻ at m/z 137.0243, C₇H₆O₃, and a characteristic fragment at m/z 109 for the cleavage of CO (Lutter et al., 2007). Lately eluted, peak **36** m/z 461.1653 [M-H]⁻ C₂₀H₃₀O₁₂ and showing a daughter ion at m/z 149 [M-H-162-132]⁻ suggesting for the loss of a hexoside and pentoside and further dehydration (Y. Zhang et al., 2012). Hence, it was annotated as homovanillyl alcohol-*O*-pentosyl -*O*-hexoside.

Following, methoxy benzaldehyde, eluted at peak **60** with a molecular ion at m/z 137.0600 [M+H]⁺, C₈H₈O₂, and fragment ion at m/z 122 and 109 for the consecutive cleavage of CH₃ and CO (Razanamaro et al., 2015). Similarly, peaks **61** & **62** with [M-H]⁺ at m/z 133.0651, C₉H₈O, &

165.0907, C₁₀H₁₂O₂ and fragments at m/z 105 & 137 accounting for the loss of CO were annotated as cinnamaldehyde (Seukep et al., 2013) and eugenol (Razanamaro et al., 2015), respectively.

Several reports have demonstrated the beneficial effects of cinnamaldehyde in diabetes management. For instance, it improves glucose uptake and increases insulin sensitivity, increases glycogenesis, and reduces gastric emptying rate (Zhu et al., 2017). Additionally, eugenol is reported to exert antidiabetic activity via suppressing intestinal α -glucosidase and inhibiting sugar binding to serum albumin (P. Singh et al., 2016).

The presence of benzoic acids in *A. digitata* fruits was previously reported (Ismail et al., 2019) and were scarcely detected in peak **9** assigned as vanillic acid from its [M-H]⁻ at m/z 167.0353, C₈H₈O₄ and fragment at m/z 123 [M-H-44]⁻ for the loss of CO₂ (Gruz et al., 2008). While peak **10** at m/z 153.0193 [M-H]⁻, C₇H₆O₄ and fragment at m/z 109 for the loss of CO₂ was annotated as dihydroxy benzoic acid. Benzoic acids, specifically vanillic acid, proved to protect against hyperglycemia and hyperlipidemia (Chang et al., 2015).

Cinnamates were detected at peaks **27** & **37**, and were annotated as hydroxycinnamic acid, coumaric acid isomers as concluded from their molecular ions at m/z 163.0401 [M-H]⁻, 165.0544 [M+H]⁺ and formula of C₉H₈O₃ (Ismail et al., 2019). Coumaric acid showed beneficial effects in the management of metabolic disorders *via* reducing serum glucose, cholesterol, and triglycerides levels (Amalan, Vijayakumar, Indumathi, & Ramakrishnan, 2016).

Quinic acid esters were detected in peak **17** annotated as *O*-caffeoyl quinic acid ester (chlorogenic acid) and **35** & **40** as 2 positional isomers of *O*-feruloyl quinic acid. Peak **17** exhibited a molecular ion at m/z 353.0891 [M-H]⁻, C₁₆H₁₈O₉ and a base peak at m/z 191 suggesting *O*-caffeoyl quinic acid ester (chlorogenic acid) (Ismail et al., 2019). While peaks **35** & **40** showed m/z 367.1024 [M-H]⁻, C₁₇H₂₀O₉ and a base peak at m/z 191 confirming their assignment as *O*-feruloyl quinic acid esters. Chlorogenic acids showed the capability to reduce fasting plasma glucose and glycosylated hemoglobin (Jin et al., 2015).

Other cinnamic esters included peak **19** assigned as *O*-feruloyl lactic acid at m/z 281.0660 [M-H]⁻, C₁₃H₁₄O₇ and yielding daughter ions at m/z 193 for the loss of lactic acid moiety (C₃H₆O₃) (S. Wu et al., 2019). Peak **50** with a molecular ion at m/z 293.0671 [M-H]⁻, C₁₄H₁₄O₇ and fragments at m/z 163 [M-H- 130]⁻ for the cleavage of C₅H₆O₄ (methyl malic acid) and m/z 145 & 119 for dehydration and decarboxylation was tentatively annotated as *O*-coumaroyl malic acid methyl ester (M. K. Lee et al., 2007). Another coumaroyl tyrosine conjugate was detected in peak **52** at m/z 326.1032 [M-H]⁻ and yielding fragments at m/z 206 for tyrosine and subsequent fragments at m/z 163 and 145 for deamidation and dehydration (Clifford & Knight, 2004).

Five cinnamoyl glycosides were identified in peaks **21**, **24**, **30**, **43**, **51**, to show fragment ions [M-H-162]⁻ indicative of a hexoside moiety. Peak **21** was annotated as dihydrocoumaroyl *O*-hexoside (dihydromelilotoside) at m/z 327.1082 [M-H]⁻, C₁₅H₂₀O₈ (Yang et al., 2007). While peak **24** as caffeoyl acid *O*-hexoside showing a molecular ion at m/z 341.0893 [M-H]⁻, C₁₅H₁₈O₉ (X.-N. Li et al., 2017). Similarly, two isomers for coumaroyl *O*-hexoside were detected in peaks **30** and **43** at m/z 325.0925 [M-H]⁻, C₁₅H₁₈O₈ (Ibrahim M Abu-Reidah et al., 2013), whereas ferulic acid *O*-hexoside was observed in peak **51** at m/z 355.1022 [M-H]⁻, C₁₆H₂₀O₉ (Ibrahim M Abu-Reidah et al., 2013). Ferulic acid was detected in peak **26** as feruloyl *O*-pentosyl hexoside at m/z 487.1457 [M-H]⁻, C₂₁H₂₈O₁₃ and fragment at m/z 193 corresponding to the loss of a hexoside and pentoside moieties (Ibrahim M Abu-Reidah et al., 2013).

Monomeric proanthocyanidin for (epi)catechin was detected with high abundance in peaks **16** & **28** at m/z 289.0716 [M-H]⁻, C₁₅H₁₄O₆ (Ismail et al., 2019). Dimeric procyanidins were seen in peaks **20**, **22** & **45** at m/z 577.1346, C₃₀H₂₆O₁₂ with fragment at m/z 407, and 289 which are characteristic of type B proanthocyanidin previously reported in *A. digitata* fruits (Ismail et al., 2019). Isomers of proanthocyanidin trimers were detected in peaks **23**, **33** & **34** exhibiting molecular ions at m/z 865.1992, C₄₅H₃₈O₁₈, and distinctive fragment ions at m/z 577, 407, 289 (Ismail et al., 2019). Procyanidin tetramer was recognized amply at peak **38** with a molecular ion at m/z 1153.2595 [M-H]⁻, following the same fragmentation pattern as the dimers and trimers (Ismail et al., 2019). All detected proanthocyanidins were grouped in the negative molecular network as group **c** suggestive for their close biosynthetic relationship, except for the tetramer which appeared as a self-looped node.

Interestingly, procyanidins as a major metabolite class in *Adansonia digitata* L. are reported to exert antidiabetic activity *via* various mechanisms of action (Wang, Wang, Ren, & Hu, 2017), including inhibition of α -amylase (Kato, Kushibiki, Inagaki, Kurokawa, & Kawabata, 2017), improvement of insulin resistance (Ogura et al., 2016), reduction of glucogenesis and increase glucose uptake (Li, Sui, Li, Xie, & Sun, 2016).

Flavonols and their glycosides were depicted in peaks **42**, **44**, **46**, **47**, **48**, **49**, & **54**. For example, quercetin-*O*-deoxyhexosyl hexoside isomers were detected in peaks **42** & **44** at m/z 609.1457 [M-H]⁻, C₂₇H₃₀O₁₆ showing the loss of a hexoside and deoxyhexoside moieties to reveal quercetin aglycone moiety at m/z 301 (Ismail et al., 2019). Likewise, kaempferol-*O*-deoxyhexosyl hexoside isomers in peaks **47** & **49** at m/z 593.1505 [M-H]⁻, C₂₇H₃₀O₁₅ and fragments at m/z 447 and 285 (Ferrerres et al., 2017) were identified along with kaempferol-*O*-hexoside isomers in peaks **46** & **54** at m/z 447.09314, C₂₁H₂₀O₁₁ and fragment ion at m/z 285. The identification of kaempferol was confirmed from its peak **48**, observed in the positive ionization mode at m/z 287.0543, C₁₅H₁₀O₆. Flavonols being antioxidant, anti-inflammatory, are known to suppress the activity of α -amylase, and glucose transporters, and to enhance pancreatic β -cell function (Zamora-Ros et al., 2014).

Furanocoumarins are reported to exhibit numerous biological activities, including anti-inflammatory (J.-K. Lee et al., 2010), vasodilatory (Duarte et al., 1995) and photosensitizing properties (De Leeuw, Assen, Van Der Beek, Bjerring, & Martino Neumann, 2011). They were observed for the first time in peaks **41**, **58** & **59**, as proved from their UV and MS spectra (Ganzera, Sturm, & Stuppner, 1997).

Peak **41** detected in the positive ionization mode exhibited a molecular ion at m/z 409.1111 [M+H]⁺, C₁₉H₂₀O₁₀, and a distinct fragment at m/z 247 for the loss of a hexoside moiety (Schuster, 1980). Thus, it was ascribed as khellol-*O*-hexoside. While, peaks **58** & **59** showing high abundance, exhibited molecular ions at m/z 261.0753 [M+H]⁺, C₁₄H₁₂O₅ and m/z 231.0648 [M+H]⁺, C₁₃H₁₀O₄ and were annotated as khellin and visnagin, respectively as concluded from their fragmentation pattern.

Likewise, pyranocoumarins were also detected for the first time in *Adonsonia digitata* fruits. An acylated derivative of the pyranocoumarin, khellactone was recognized at peak **65** as a Na adduct at m/z 411.1410 corresponding to $C_{21}H_{24}O_7$ as previously detected by (Olennikov et al., 2019). Its MS₂ spectrum showed fragments at m/z 351 $[M+Na-C_2H_3O_2]^+$ for the cleavage of an acetic acid moiety, m/z 245 for the consecutive loss of an isovaleryl moiety (C_5H_8O) followed by m/z 227 and 199 for the elimination of H_2O and CO . Consequently it was ascribed as isovaleryl-acetyl khellactone (Olennikov et al., 2019). *O*-substituted esters of khellactone are reported to have numerous biological activities to include antiobesity (Taira et al., 2017), anti-HIV (Xie, Takeuchi, Cosentino, & Lee, 1999) and anti-cancer (J. Y.-C. Wu et al., 2003) activities.

Similarly, peak **66** found in the positive ionization mode, with molecular ion at m/z 329.1377, $C_{19}H_{20}O_5$ was annotated as decursin following its fragmentation pattern with fragment ions at m/z 245 $[M+H-C_5H_8O]^+$ (H. J. Lee et al., 2009). The antibacterial (S. Lee, Shin, Kim, Oh, & Kang, 2003), anticancer (Ahn, Sim, & Kim, 1995; H. J. Lee et al., 2009) and neuroprotective (Kang & Kim, 2007) properties of decursin have been previously documented.

Fatty acids composition of *Adonsonia* fruits comprised saturated fatty acids, monounsaturated fatty acids, and oxygenated fatty acids. Fatty acids were eluted at peaks **67-77** and were annotated based on their molecular formula and fragmentation pattern and in comparison with previously reported literature.

A megastigmane hexoside was observed at peak **55** with a molecular ion at m/z 405.2123 $[M-H]^-$ and fragment at m/z 225 for the loss of a hexoside moiety and dehydration (K. Singh et al., 2014).

4.1 Conclusion and future perspectives

The current study shows the potential of *Adansonia digitata* L. methanolic fruit pulp crude extract as an antidiabetic plant. The extract has low toxicity both *in vitro* and *in vivo* and a strong antioxidant activity.

The UHPLC/MS showed that the fruit extract contains several chemical groups, such as procyanidins. This chemical class is known to have antidiabetic activity via various mechanisms of action, such as inhibition of α -amylase, improvement of insulin resistance, reduction of glucogenesis, and increase glucose uptake. Moreover, a total of 77 compounds were dereplicated belonging to different classes, including organic acids, sugars, alcohols, phenolics, coumarins, and fatty acids. Almost 50% of the identified metabolites (40 compounds) are reported for the first time in *Adonsonia digitata* fruits.

The biochemical parameters supported the hypothesis of the antidiabetic effect of the *Adansonia digitata* L. However, further, characterization is needed to identify the optimal dose and mode of administration. The histopathological investigation supported the biochemical findings and revealed that the fruit extract has hepatoprotective, and renoprotective effects *in-vivo*.

Further studies are needed for the isolation and investigation of the anti-diabetic effect of the main identified ingredients, such as the procyanidins. Furthermore, a study of the molecular pathway of such active chemicals is necessary to understand the antidiabetic mechanisms better.

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Appendix

Table 4: Histopathological Scoring of tissue lesions. (Klopfleisch, 2013; Knodell et al., 1981).

Table 3: Score Description					
Liver			Kidney		
Lesion	Description	score	Lesion	Description	Score
Degeneration`	Normal	0	Degeneration of tubular epithelium	Normal	0
	< 25% of the entire area	1		< 25% of the entire area	1
	25-50% of the entire area	2		25-50% of the entire area	2
	50-75% of the entire area	3		50-75% of the entire area	3
	>75% of the entire area	4		>75% of the entire area	4
Necrosis	Normal	0	Necrosis of tubular epithelium	Normal	0
	<25% of the area	4		< 25% of the entire area	2
	25-50% of the area	6		25-50% of the entire area	4
	>50% of the area	8		50-75% of the entire area	6
	25-50% of the area + bridging necrosis	10		>75% of the entire area	8
Inflammation	Normal	0	Interstitial infiltration	Normal	0
	<1/3 of the area	1		< 25% of the entire area	1
	1/3 – 2/3 of the area	2		25-50% of the entire area	2
	>2/3 of the area	3		50-75% of the entire area	3
				>75% of the entire area	4
The damage was obtained by summation the entire score of the above histopathological lesions					