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Perfection is not attainable, but if we chase perfection we can catch excellence.

Vince Lombardi

To My Mother "*Naseem Akhtar*"
May Allah keep her soul in peace and rest Aameen

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LIST OF ABBREVIATIONS

AAPH	2,2'-azobis(2-methylpropionamide) dihydrochloride
AGEs	Advanced glycation end products
AMG	Aminoguanidine
AP-1	Activator protein-1
BBI	Broadband inverse
BSA	Bovine serum albumin
BSTFA	N,O-Bis(trimethylsilyl) trifluoroacetamide
CDC	Centres for disease control and prevention
CEL	Carboxyethyl-lysine
CML	Carboxymethyl-lysine
COSY	Correlation spectroscopy
CPRG	Chlorophenol red- β -D-galactopyranoside
DAD	Diode array detector
DEPT	Distortionless Enhancement by Polarization Transfer
3-DG	3-deoxyglucosone
D.I.Khan	Dera Ismail Khan
DPPH	2,2-diphenyl-1-picrylhydrazyl
DOLD	Deoxyglucosone-derived lysine dimers
DMSO	Dimethyl sulphoxide
ECM	Extracellular matrix
ELSD	Evaporative light scattering detector
FAS	Fatty acid synthesis
FCS	Fetal calf serum

FFI	2, 2(furyl), 4-(5) 2 (furanyl)1H-imidazole
GAE	Gallic acid equivalents
GO	Glyoxal
GOLD	Glyoxal lysine dimer
GC	Gas chromatography
HMBC	Heteronuclear multiple bond correlation
HPLC	High performance liquid chromatography
HSQC	Heteronuclear single quantum coherence
KPK	Khyber pakhtoon khawa
15-LOX	15-lipoxygenase
MBC	Minimum Bactericidal Concentration
MEM	Minimum Essential Medium
MGO	Methyl glyoxal
MHB	Mueller Hinton Broth
MIC	Minimum Inhibitory Concentration
MOLD	Methylglyoxal lysine dimers
MRC-5	Medical Research Council -5 (human lung fibroblast)
MS	Mass spectroscopy
NMR	Nuclear magnetic resonance
NP	Normal phase
NBT	Nitro blue tetrazolium
NF-IL6	Nuclear factor interleukin-6
PBS	Phospahte buffer saline
PC-3	Human prostate cancer cell line

PES	Phenazine ethosulphate
Pnp	<i>p</i> -nitrophenyl- α -D-glucopyranoside
QTOF-MS	Quadrupole-time-of-flight mass spectrometer
RAGE	Receptor for AGEs
ROS	Reactive oxygen species
RPMI	Roswell Park Memorial Institute
RiOS	Reverse osmosis
RUE	Rutin equivalents
SP-1	Specificity protein-1
TCA	Trichloroacetic acid
TFC	Total flavonoid contents
TLC	Thin layer liquid chromatography
TPC	Total phenolic contents
TM	Traditional medicine
TMCS	Trimethylchlorosilane
TQD-MS	Triple quadrupole mass spectrometer
UV-VIS	Ultraviolet-visible light
WHO	World Health Organization

CHAPTER 1
INTRODUCTION
AND AIM OF THE WORK

1.1 Traditional medicine

Medicinal plants are intrinsic components of traditional medicinal systems in treating and preventing an array of ailments. Traditional medicine (TM) is a broad term that refers to “systems” such as Traditional Chinese Medicine (TCM), Indian Ayurveda and Arabic Unani medicine, or multiple forms of indigenous medicine. However, in some countries TM is termed as "alternative" or "complementary" medicine (WHO, 2002).

The history of traditional medicine is as old as human civilization. Earliest records confirm that herbal medicines have been used and documented in Roman, Greek, Egyptian, Chinese and Indian medicinal systems for about 5000 years. Traditional herbal medicine has also been practised from ancient times in American, Arabian countries and Japan. The records of traditional medicine systems in the Indian subcontinent (India, Pakistan, Bangladesh, Nepal, Bhutan, Sri Lanka and the Maldives) include Ayurveda, Rigveda, Siddha and Yunani. Here the correlation between TM and biodiversity is demonstrated by an old custom of “healing powers” associated with the “earth's natural systems” that involves medicinal plants and animal species, the ambient healthy air and spring water (Good, 1980; Gesler, 1992). For example, Ayurveda of India has a documented history dating back to 1500 BCE (Sharma, 1994; Dwarakanath, 1952) and its authenticity is proven most recently in a genome-wide SNP (single nucleotide polymorphism) analysis (Affymetrix, 6.0) suggesting that India’s traditional medicine has a genetic basis, and its Prakriti-based (constitutional types of every human) practice in vogue for many centuries resonates with personalized medicine (Govindaraj *et al.*, 2015).

In the Indian subcontinent, practice of TM is mainly accomplished by “herbal practitioners” that are mainly referred to as “Sadhu” or “Hakeems”. Their practice is mainly based on secret herbal recipes. These recipes were formulated through traditional knowledge transferred from generation to generation. The safety, accessibility, cheapness and sociological acceptance are major factors in the popularity of traditional medicine in developing nations (Mesia, 2009). This trend continues even in our modern era and about 4 billion people, 80% of the world's population, presently use plant-based medicine for some aspects of their primary health care (WHO, 2008).

More interestingly, in a review covering anticancer drugs for instance, it has been reported that of 175 small molecules, approximately 48.6% of drugs were actually being either natural products or directly derived therefrom (between 1981 to 2010)(Newman and Cragg, 2012). WHO has compiled a comprehensive list of 21, 000 medicinal plant species (including synonyms), that are used worldwide, based on the data collected from 91 countries (WHO, 2002a).

1.2 Traditional medicine and the 21st century medicine market.

Indeed, immense progress has been made in the discovery of synthetic medicines for the treatment or prevention of several life threatening diseases; however, excessive and prolonged use of such drugs may produce toxic side effects that are considered as a continuous hazard in modern health systems (Alves and Rosa, 2007; Dubey *et al.*, 2004). In the 21st century, medicinal herbs are gaining importance in the mainstream healthcare system in developed countries. A large number of people quest for relatively safe remedies and approaches to healthcare. The demand for herbal medicines, herbal pharmaceuticals, herbal health products, nutraceuticals, food supplements and herbal cosmetics, etc. is boosting globally. The main reason for this trend is thriving appreciation of herbal products as showing less side effects, showing improved compatibility with physiological flora, and being available at reasonable prices (Dubey *et al.*, 2004; Sharma *et al.*, 2008). Statistics demonstrate that traditional herbal medicine includes 30–50% of the total medicinal usage in China. In North America, Europe and other well-developed regions over 50% of the population have used traditional preparations at least once. In 2003, it was reported that the world market for herbal medicines raised to over \$60 billion(US) per year (WHO, 2003) and estimates suggest that global medicinal plant business will reach \$5 trillion (US) by 2050 (Shinwari, 2010).

1.3 Ethnomedicinal assessment and drug discovery

In modern times ethnomedicinal assessment of medicinal plant species is considered as a primary step in the discovery of herbal drugs (Vitalini *et al.*, 2013), followed by isolation of bioactive principles, their pharmacological screening and finally ending up in finding new therapeutically active drugs. However, the approval of such a plant based drug or formulation on the market in modern times needs a series of procedures and measures. A number of approaches can be adopted in this regard; however, the scheme proposed by Pieters and Vlietinck (Fig. 1.1) is of particular importance. The scheme proposes two possible operational modes i.e working with standardised extracts or the isolation of pure active components. In both approaches finally toxicity and safety studies, as well as clinical trials are needed for final drug approval (Pieters and Vlietinck, 2005).

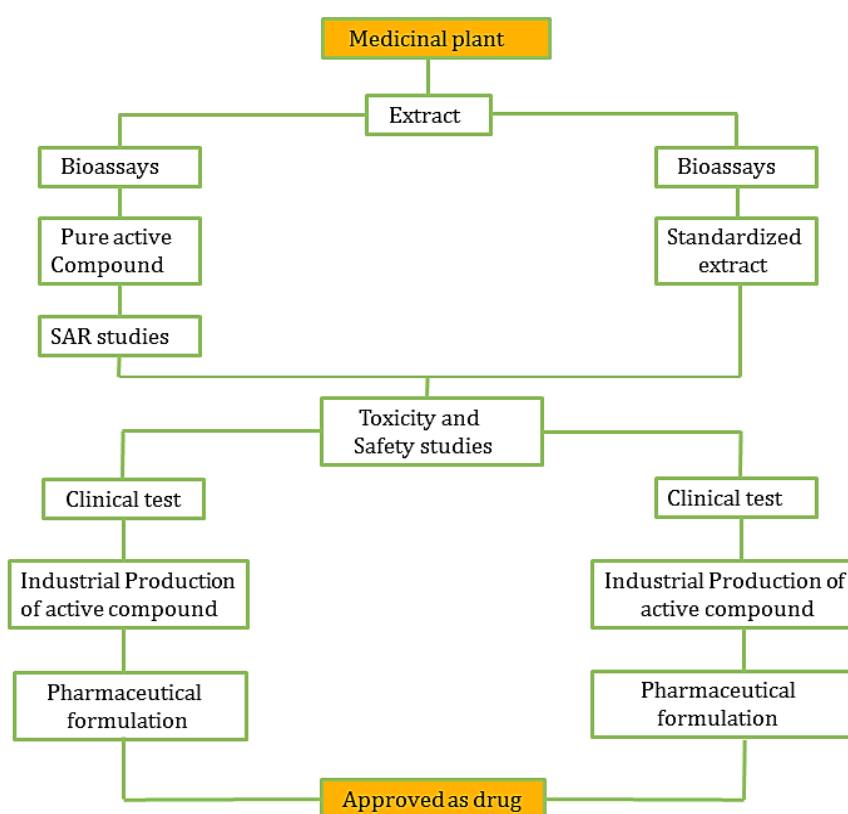


Figure 1.1. Scheme for the study of medicinal plants proposed by Pieters and Vlietinck, 2005.

1.4 Traditional medicine in Pakistan

Pakistan has an area of 80,943 km² and lies between 60°55' to 75° 30' E. longitude and 23° 45' to 36° 50' N. latitude. It has an altitude ranging from 0 to 8611 m, with nine major ecological zones and a unique biodiversity.

The country has a subtropical climate and a total of 1572 genera and 5521 species of flowering plants have been identified till now, out of which 400–600 are medicinally important (Bibi *et al.*, 2015). Pakistan has four phytogeographical regions including the Saharo-Sindian (9.5% of species), Sino-Himalayan (10%), Irano-Turanian (45%) and Indian element (6%) (Shinwari *et al.*, 2000). The plant “hotspots” of Pakistan are spread over 13 natural regions from alpine pastures to mangrove forest (Shinwari *et al.*, 2002).

The traditional healthcare system of Pakistan is mainly based on indigenous home-based herbal remedies and experienced local healers called “Hakeems” that use their centuries old traditional medicinal knowledge. Although the indigenous knowledge about medicinal uses of plants has been recorded widely in Pakistan in recent years (Kayani *et al.*, 2015; Rehman *et al.*, 2015; Bibi *et al.*, 2014; Abbasi *et al.*, 2013; Qureshi, 2012; Ahmad *et al.*, 2014; Shinwari *et al.*, 2000; Shinwari *et al.*, 2002) and efforts have been made in isolation of active chemical constituents from medicinal plants (Bibi *et al.*, 2010; Bibi *et al.*, 2011; Choudhary *et al.*, 2011; Irshad *et al.*, 2011; Amin and Khan, 2011; Jamil *et al.*, 2012; Ejaz *et al.*, 2014), such efforts need to be broadened especially to less explored plants.

Therefore, the current project was designed to isolate bioactive phytochemicals from some less explored but commonly used medicinal plants from Pakistan, as a part of running programmes in the host laboratory on antimicrobial (especially antiprotozoal) and antidiabetic properties (inhibition of α -glucosidase activity; antioxidant activity; inhibition of Advanced Glycation Endproducts-AGEs) of medicinal plants and natural products (Upadhyay *et al.*, 2014a, Upadhyay *et al.*, 2014b; Mesia *et al.*, 2008).

1.5 Diabetes mellitus

Diabetes mellitus is a multifarious metabolic disorder (involving carbohydrate, fat and protein metabolism) that results from either insufficiency or dysfunction of insulin, ending up in elevated fasting and post-prandial blood sugar levels. It is generally categorized as Type I and II diabetes (primary forms) (Fig. 1.3). The Type I (also referred to as insulin dependent) or juvenile onset diabetes is characterized by cell mediated, autoimmune destruction of the pancreatic cells, whereas in Type II (also referred to as non-insulin dependent), the patients are unable to respond to insulin (insulin resistance). Type II is the most common form and constitutes up to 90% of diabetes worldwide, whereas Type I accounts for 5–10% of patients. According to recent estimations, 387 million people worldwide are affected by the disease (IDF (International Diabetes Federation), 2014). According to WHO the major burden is in developing countries. Pakistan is one of the 19 countries and territories of the IDF MENA (Middle-East and North-Africa) region. It has been estimated that 415 million people have diabetes in the world and more than 35.4 million people in the MENA region; by 2040 this will rise to 72.1 million. There were over 7 million cases of diabetes in Pakistan in 2015 (IDF, 2015).

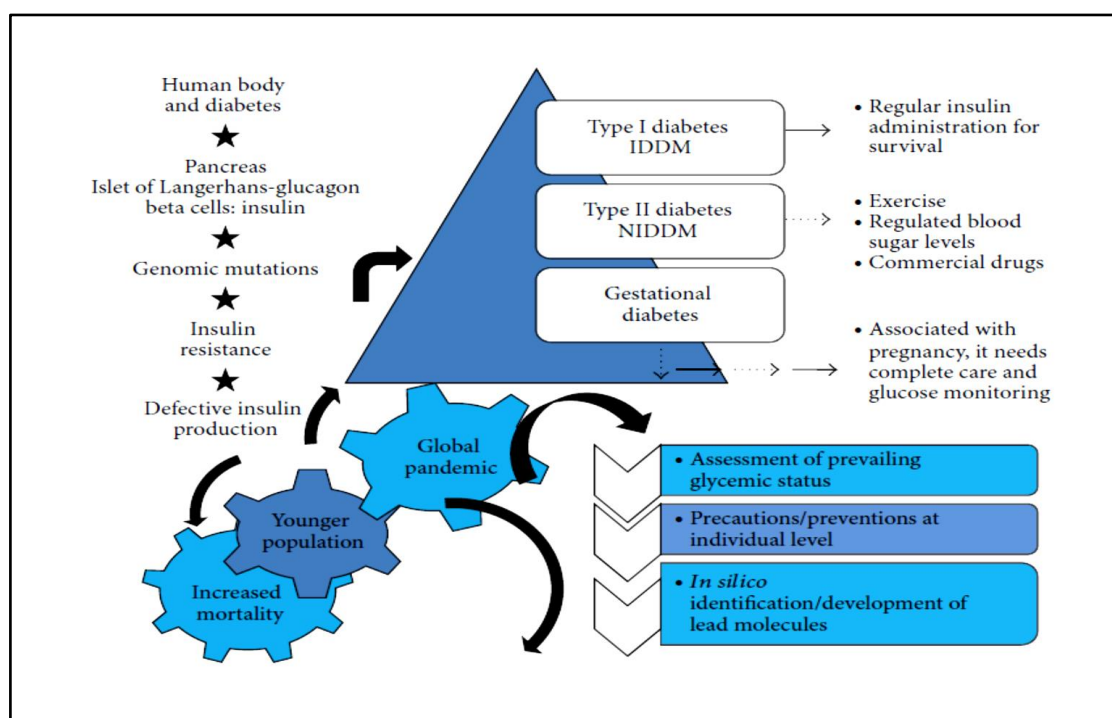


Figure 1.3. Occurrence and management of diabetes (Adopted from Tiwari, 2015).

The development of diabetes can be prevented or delayed in people with impaired glucose tolerance by implementing lifestyle changes or the use of therapeutic agents. Indeed, the pathophysiology of diabetes needs more investigations. However, it has been proven that free radicals and oxidative stress are associated with the development and pathogenesis of diabetes (Matteucci and Giampietro, 2000; Lipinski, 2001). In diabetic patients, extra-cellular and long living proteins such as laminin, collagen, elastin, several plasma proteins, including haemoglobin, serum albumin and transferrin are the major targets of free radicals. As a result of continuous and long term exposure to glucose, plasma proteins endure glycation and chemical modifications, which results in the formation of glycoproteins (Austin *et al.*, 1987; Glugliano *et al.*, 1996) that finally form “advanced glycation endproducts” (AGEs) after a series of rearrangements.

1.6 Advanced glycation end product (AGEs)

Glycation is a non-enzymatic reaction between the carbonyl group of reducing sugars and a free amino group of proteins, leading to the formation of a Schiff's base. The Schiff's base is the first product of the glycation reaction that is fast and highly reversible (Ulrich and Cerami, 2001). Subsequently thermodynamically unstable Schiff's bases are converted to a stable and reversible Amadori product. Proteins bearing an Amadori product are then referred to as glycated proteins or Maillard reaction products. Finally the Amadori product undergoes a series of dehydration and fragmentation reactions, resulting in a variety of carbonyl compounds including methylglyoxal (MGO), glyoxal (GO), glucosones, 3-deoxyglucosone (3-DG) and so on (Thornalley *et al.*, 1999). These carbonyl compounds are more reactive than the original sugar and act as reaction propagators, leading to the formation of advanced glycation end-products (AGEs) (Fig.1.4).

The AGEs formed are known to be quite heterogeneous, depending on the glycating agent. For example, glyoxal-derived AGEs include carboxymethyl-lysine (CML) and glyoxal lysine dimer (GOLD), while methylglyoxal-derived AGEs are carboxyethyl-lysine (CEL), argpyrimidine and methylglyoxal lysine dimers (MOLD), and 3-deoxyglucosone-derived AGEs are pyrroline and deoxyglucosone-derived lysine dimers (DOLD) (Thornalley *et al.*, 1999).

The glycation reaction mainly occurs at the terminal amino groups of lysine, arginine and histidine, whereas glycation-mediated protein crosslinking is associated with arginine residues (Munch *et al.*, 1999). The dicarbonyl compounds, like glyoxal, deoxyglucosone or methylglyoxal, favour the formation of intracellular AGEs (Wells-Knecht, 1995). Various proteins, including collagen, interact with AGEs or reactive dicarbonyl compounds and form protein crosslinks (Reddy, 2004). Quantitative analysis of AGEs have shown that intracellular proteins are more heavily AGEs-modified than extracellular proteins (Thornalley *et al.*, 2003)(Fig.1.4).

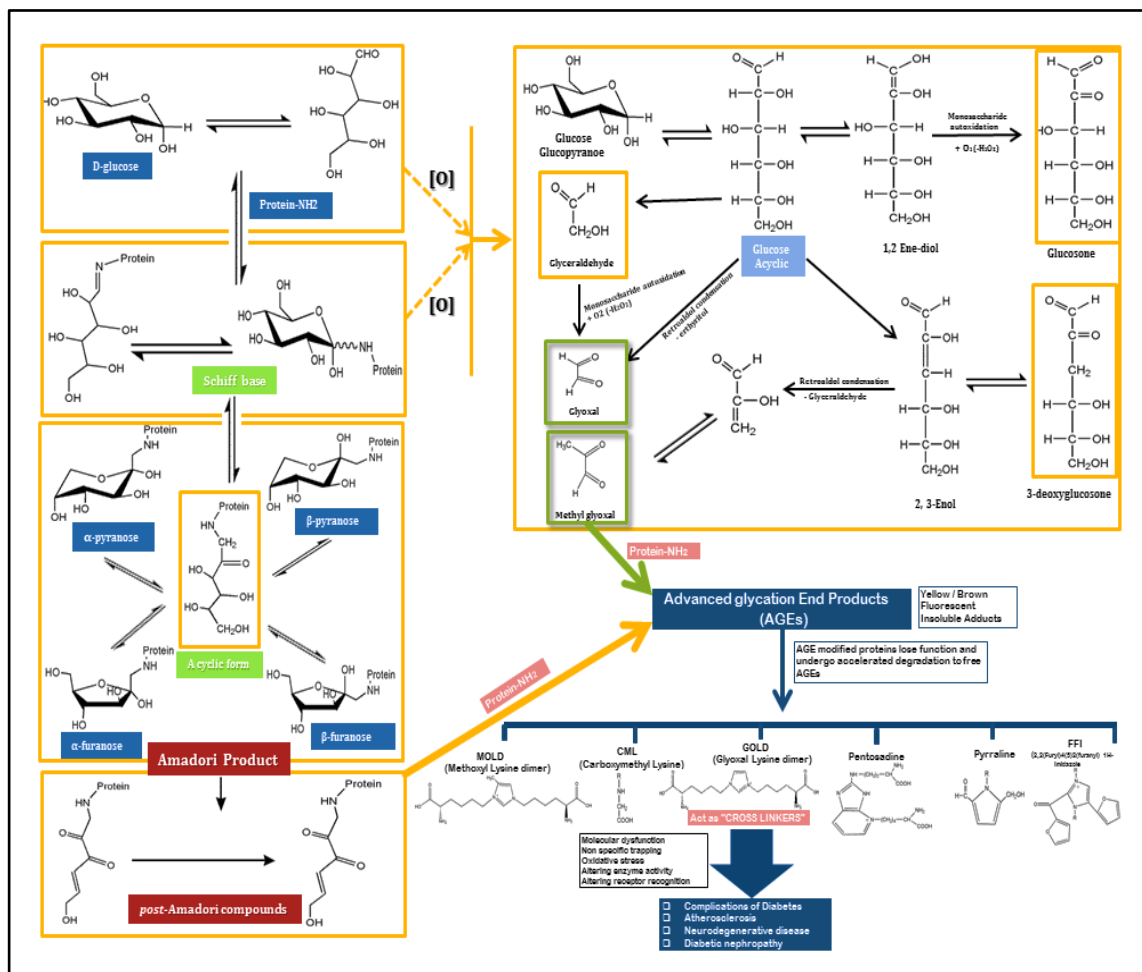
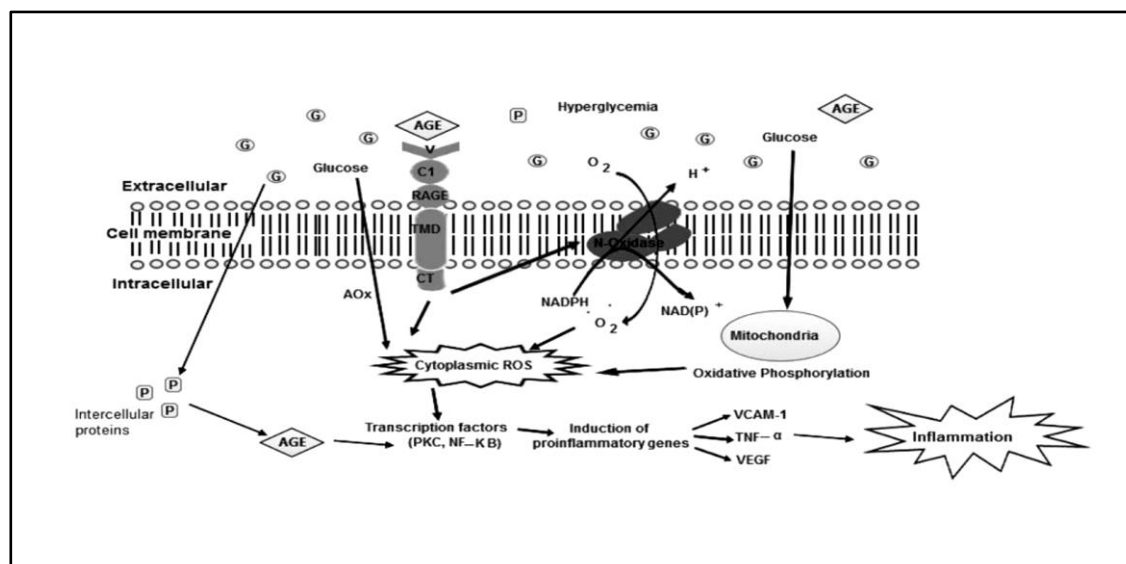


Figure 1.4. AGEs formation and complications (adopted from Adrover *et al.*, 2008 and Thornalley *et al.*, 1999).

1.6.1 AGEs and pathogenesis of diabetic complications

AGEs are mainly involved in pathogenesis of numerous diseases depending on their mode of interaction with target cells. In general AGEs cause modification in the structure and function of intracellular proteins, extracellular matrix (ECM) and cell receptors, and initiate a series of altered cellular processes. In addition the AGE modified protein activates a downstream signalling cascade by binding to receptor for AGEs (RAGE) leading to nuclear translocation of transcription factors such as NF- κ B, specificity protein-1 (SP-1), activator protein-1 (AP-1) and nuclear factor interleukin-6 (NF-IL6). All these then contribute to various cellular processes including pro-inflammatory responses, overexpression of RAGE and generation of reactive oxygen species (ROS), etc. (Brownlee, 2001). Other complications of AGEs include atherosclerosis, nephropathy, uremia, poor wound healing, diabetic retinopathy, neurodegenerative diseases, cancer, and the normal ageing processes (Schmidt *et al.*, 1994; Basta *et al.*, 2004; Heijst *et al.*, 2006) (Fig.1.5)



RAGE; receptors for AGEs, V, C1 and C2(extracellular domains for RAGE), TMD; Transmembrane domain, CT; cytoplasmic tail, ROS; reactive oxygen species, AOx; Autooxidation of glucose, PKC; protein kinase C, NF- κ B; nuclear factor kappa light chain enhancer of activated B cells, VCAM-1; Vascular cell adhesion protein 1, TNF- α ; tumour necrosis factor α , VEGF; Vascular endothelial growth factor, NADPH nicotinamide adenine dinucleotide phosphate.

Figure 1.5. Simplified mechanisms of the pathogenesis of AGEs in hyperglycemic condition (Adopted from Khangholi *et al.*, 2016)

1.6.2 Natural products as AGEs-inhibiting compounds

A number of plant species and plant products have been shown to exhibit antiglycation activity. Some biologically important phytochemicals, particularly flavonoids and polyphenols, terpenoids, alkaloids and iridoids have been studied extensively for their antioxidant and anti-glycation activities (Fig. 1.6).

1.6.3. Polyphenols

The polyphenolic compounds are main constituents of plants and are known for a variety of biological activities. A number of polyphenolics have been reported for their antiglycation potential in literature; for instance: vescalagin (**1**), a hydrolyzable tannin recently isolated from the fruit of *Syzygium* sp. It showed a protective effect against methylglyoxal-induced inflammation and carbohydrate metabolic disorder in rats. The effect was considered to be caused by an increase in concentration of D-lactate that inhibited AGE formation and a decrease in cytokine release to prevent damage to β -cells of the pancreas (Chang *et al.*, 2013).

Similarly several natural stilbenoids such as resveratrol and its derivatives have been reported to exhibit a broad-spectrum of biological activities, including anti-glycation activity. For example, oxyresveratrol (**2**) isolated from the heartwood of *Artocarpus lakoocha* showed anti-glycation activity with IC_{50} 8.1 μ M (Povichit *et al.*, 2010).

Nevertheless, a number of different types of polyphenolic compounds were isolated from *Rhus verniciflua* and tested for their antiglycation effect. It was reported that butein (**3**) (IC_{50} 0.7 μ M) and sulfuretin (**4**) (IC_{50} 1.3 μ M) showed a remarkable inhibitory effect on recombinant human aldose reductase and the accumulation of AGEs (IC_{50} 200 μ M) and (IC_{50} 1240 μ M) respectively (Lee *et al.*, 2008).

1.6.4. Flavonoids

The flavonoids are generally known for their antiglycation effects. Flavones are more active than the corresponding flavanones, isoflavones, and flavonols. In general, some structural features of flavonoids are considered as essential for the inhibition of AGEs formation, such as hydroxyl groups at C-3, C-5 and C-7. Methylation or glucosylation of the hydroxyl groups alters the activity of flavones, flavonols and flavanones (Matsuda *et al.*, 2003).

A series of natural flavonoids has been investigated for their potential as an inhibitor of protein glycation. Quercetin (**5**) is an important flavonol and possesses a broad spectrum of biological activities. It is known as a potent antiglycation agent. *Psidium guajava* leaf extract was found to inhibit glycation processes in an albumin/glucose test system and quercetin was identified as one of the active principles. It has been found to exhibit a strong inhibitory effect on glycation of albumin in various assays (Wu *et al.*, 2009a).

Interestingly, the natural flavonol glycosides are reported as effective anti-glycation agents. The Korean medicinal plant *Artemisia capillaris* was explored for its potential antiglycation activities. A series of compounds were isolated. Some flavonol glycosides i.e isorhamnetin 3-robinobioside (**6**), hyperoside (**7**) and isorhamnetin 3-galactoside-7-rhamnoside (**8**) exhibited moderate antiglycation activity (IC₅₀ 48 μM, 82 μM and 155 μM, respectively) compared to the positive control aminoguanidine (IC₅₀ 920 μM) (Jung *et al.*, 2012). Although the exact mechanism for this inhibition is not known yet, it is generally accepted that strong antioxidant capacities, free radical scavenging and/or metal ion trapping activities are mainly involved (Peng *et al.*, 2008). Likewise, silymarin (**9**) is an example of a complex of flavonoids, which acts as a specific RAGE blocker and therefore could reduce the burden of AGEs in diabetic patients. Silymarin has also been identified as an antioxidant with antiglycation properties in both *in vitro* and *in vivo* systems (Wu *et al.*, 2009b).

1.6.5 Terpenoids

Terpenoids hold a wide structural diversity and reveal a wide spectrum of biological activities. The terpenoids are classified into a number of different types including sesquiterpenes, diterpenes, triterpenes, etc.

The sesquiterpenes are commonly found in many medicinal plants and are considered as anti-AGEs agents. β -Caryophyllene (**10**), a sesquiterpene lactone, is reported as having a protective role against plasma and tissue glycoprotein components in streptozotocin-induced hyperglycemic rats (Basha and Sankaranarayanan, 2015). Similarly, labdane diterpenes and labdadiene (**11**) were isolated from the rhizomes of *Alpinia zerumbet*. These labdadienes exhibited inhibitory activities on the formation of fructosamine adducts and dicarbonyl compounds with IC_{50} 168.0 μ M and this activity was comparable with the reference flavonoids quercetin and rutin (Chompoo *et al.*, 2011).

1.6.6 Alkaloids

Alkaloids are effective anti-AGEs compounds. A number of alkaloids from different classes have been tested for AGE inhibition over the past few years. Cyclopeptide alkaloids including nummularine-R (**12**), and hemsine-A (**13**) were isolated from *Ziziphus oxyphylla* Edgw. (Rhamnaceae) and tested for AGEs inhibition. Nummularine-R (IC_{50} 720.2 μ M) and hemsine-A (IC_{50} 277.7 μ M) presented moderate anti-glycation activity. All compounds were non-toxic to the PC-3 cell line (Choudhary *et al.*, 2011).

Wang *et al.* (2007) have isolated sinomenine (**14**) from the stem and root of *Sinomenium acutum*. In an attempt to know the molecular mechanism of inhibition of this compound, retinal microglial were activated by AGEs treatment. It was noticed that sinomenine weakened ROS production in a dose-dependent way and reduced the nuclear translocation of NF-kB p65 in AGEs-activated cultured retinal cells.

More recently in a review by Abbas *et al.* (2016) new potent inhibitors of protein glycation belonging to different classes such as flavonoids, alkaloids and terpenes, have been identified. Some interesting activities were highlighted with regard to small compounds such as carnosine (**15**) (IC₅₀ 16.3 μM) and nicotinamide (**16**) (IC₅₀ 17.8 μM) as potential AGE inhibitors.

1.6.7 Iridoids

Iridoids are monoterpenes based on the cyclopenta [c] pyranoid skeleton, which are known for a number of pharmacological activities including the inhibition of protein glycation (Ghisalberti *et al.*, 1998; West *et al.*, 2014).

Catalpol (**17**), a common iridoid, was isolated from *Rehmannia glutinosa*, and tested for anti-inflammatory effects on AGE-stimulated THP-1 cells. It was noticed that catalpol causes a considerable decrease in the expression of pro-inflammatory mediators. Moreover, the effective anti-AGEs potential was shown by the decrease in transcriptional activation of NF-κB (Choi *et al.*, 2013).

More recently iridoids obtained from dietary sources including *Morinda citrifolia*, Asian cornelian cherry (*Cornus officinalis*), European cornelian cherry (*Cornus mas*) and olive (*Olea europaea*) were tested for their potential antiglycation activities. The major iridoids from *Morinda citrifolia*, deacetylasperulosidic acid (**18**) and loganic acid (**19**), were tested for AGE inhibition *in vitro*. Both inhibited glycation in a concentration-dependent manner (West *et al.*, 2014).

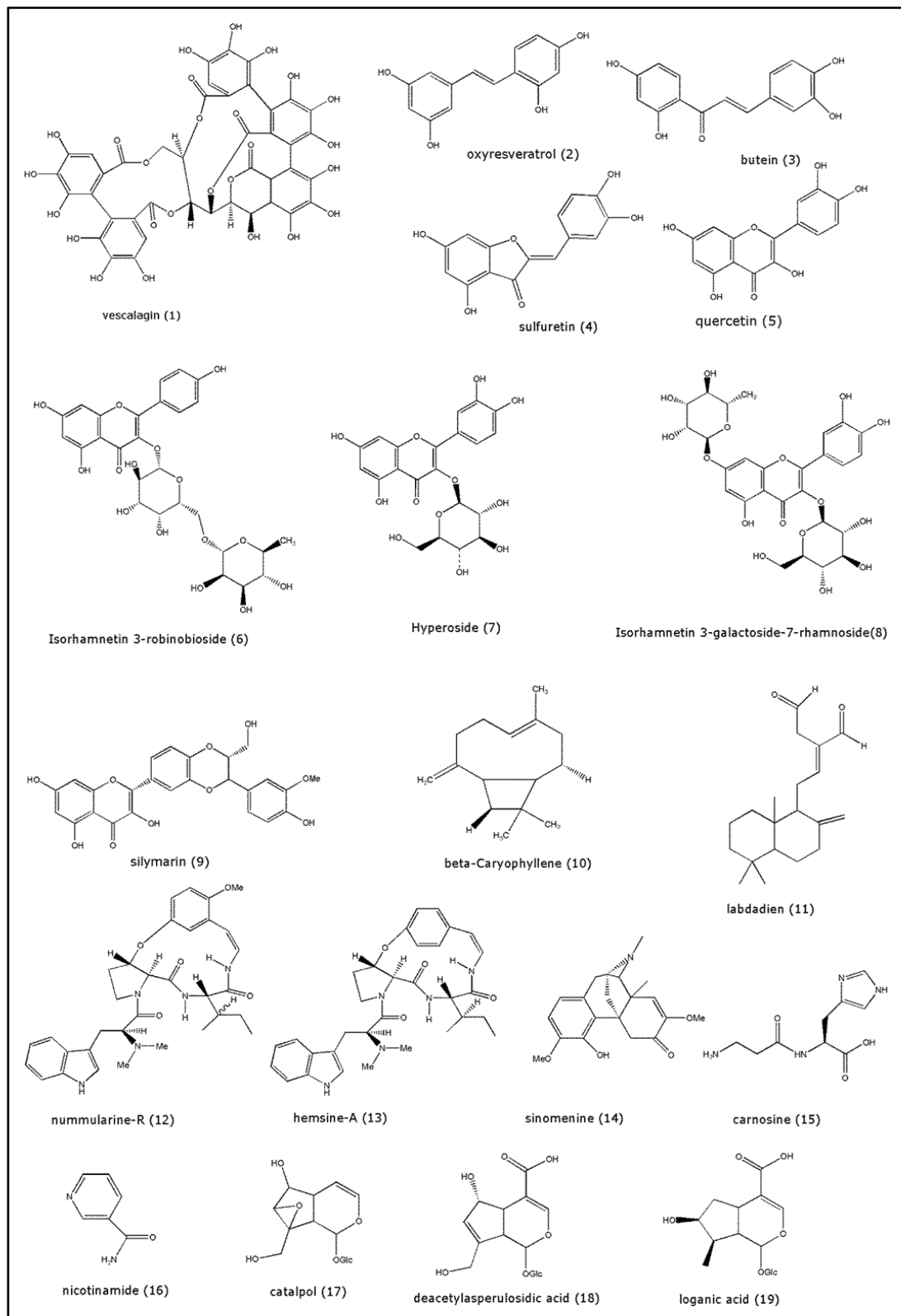


Figure 1.6. Antglycation compounds isolated from medicinal plants

1.7. Inhibition of α -glucosidase

Inhibition of alpha-glucosidase enzyme activity in the gastro-intestinal tract is a common therapeutic approach to reduce glucose-uptake in diabetic patients. Through a competitive inhibition of α -glucosidase, these drugs help to inhibit the fast breakdown of polysaccharides and thus control the blood sugar level.

Numerous α -glucosidase inhibitors (Fig 1.7) have been isolated from natural sources, for instance acarbose (**20**) and voglibose (**21**) that are clinically used as α -glucosidase inhibitors in the treatment of diabetes mellitus (Playford *et al.*, 2013). Commercially available α -glucosidase inhibitors (**20**, **21**, **22**), mainly contain nonsugar moieties, are associated with serious side effects (Yin *et al.*, 2014). Therefore, there is a need to look for promising molecules as an alternative for existing α -glucosidase inhibitors. In the literature a number of constituents, for instance terpenes, alkaloids, flavonoids and polyphenols have been reported as α -glucosidase inhibitors.

In an investigation by Mbaze *et al.*, (2007), a number of compounds were isolated from methanolic extracts of *Fagara tessmannii* (Rutaceae). Among these 3 β -acetoxy-16 β -hydroxybetulinic acid (**23**) presented a strong inhibition of α -glucosidase with an IC₅₀ of 7.6 μ M against α -D-glucosidase and IC₅₀ 397 μ M against β -D-glucosidase compared to acarbose (IC₅₀ of 780 μ M) (Mbaze *et al.*, 2007).

Similarly the crude bark extract of *Uncaria laevigata* was explored for potential α -glucosidase inhibitors. A number of triterpenes were isolated from the crude extract and tested for α -glucosidase inhibitory activity. Among all ursolic acid (**24**) and 3 β -hydroxy-30-methoxy-6-oxo-urs-12,19(20)-dien-28-oic acid (**25**) showed interesting inhibitory activity with IC₅₀ values of 49 μ M and 16 μ M, respectively, as compared with that of genistein (IC₅₀ 36 μ M), used as positive control (Wang *et al.*, 2013).

Alkaloids have been reported as interesting inhibitors of α -glucosidase. In a study aimed at the isolation of promising α -glucosidase inhibitors, methanolic extracts from the medicinal parts of 40 traditional Chinese herbs were assessed. From the methanol

extract of the leaves of *Adhatoda vasica* Nees (Acanthaceae), two alkaloids, vasicine (**26**) and vasicinol (**27**) were isolated. Both compounds inhibited rat intestinal α -glucosidase activity (IC_{50} = 125 and 250 μ M, respectively), and they were both found to be competitive inhibitors (K_i = 82 and 183 μ M, respectively) (Gao *et al.*, 2008).

Similarly in another investigation two new phenylpropanoyl amides, namely chaplupyrrolidones A (**28**) and B (**29**), were isolated from the leaf extract of *Piper sarmentosum*. Chaplupyrrolidone B (**29**) (IC_{50} 430.0 μ M) revealed α -glucosidase inhibitory activity and was 18-fold more active compared to its demethylated congener chaplupyrrolidone A (**28**) (IC_{50} 7800 μ M). Kinetic evaluation studies indicated that chaplupyrrolidones B inhibited α -glucosidase in a non-competitive manner (K_i = 1.04 mM) (Damsud *et al.*, 2013).

Flavonoids are widely distributed in plants and are known for a number of activities including α -glucosidase inhibition. In an investigation by Feng *et al.*, (2011), the ethanolic leaf extract of *Aquilaria sinensis* (Lour.) Gilg.b. was explored in a quest for α -glucosidase inhibitors. Four new compounds were isolated namely aquilarisinin (**30**), aquilarisin (**31**), hypolaetin 5-*O*- β -*D*-glucuronopyranoside (**32**), and aquilarixanthone (**33**) in addition to other (known) compounds. These were further tested for α -glucosidase inhibition, and interesting results were obtained for quilarisinin (**30**) (IC_{50} 273.7 μ M), aquilarisin (**31**) (IC_{50} 634.7 μ M), hypolaetin 5-*O*- β -*D*-glucuronopyranoside (**32**) (IC_{50} 298.9 μ M), and aquilarixanthone (**33**) (IC_{50} 678.1 μ M), that were comparable with acarbose (IC_{50} 576.2 μ M) (Feng *et al.*, 2011).

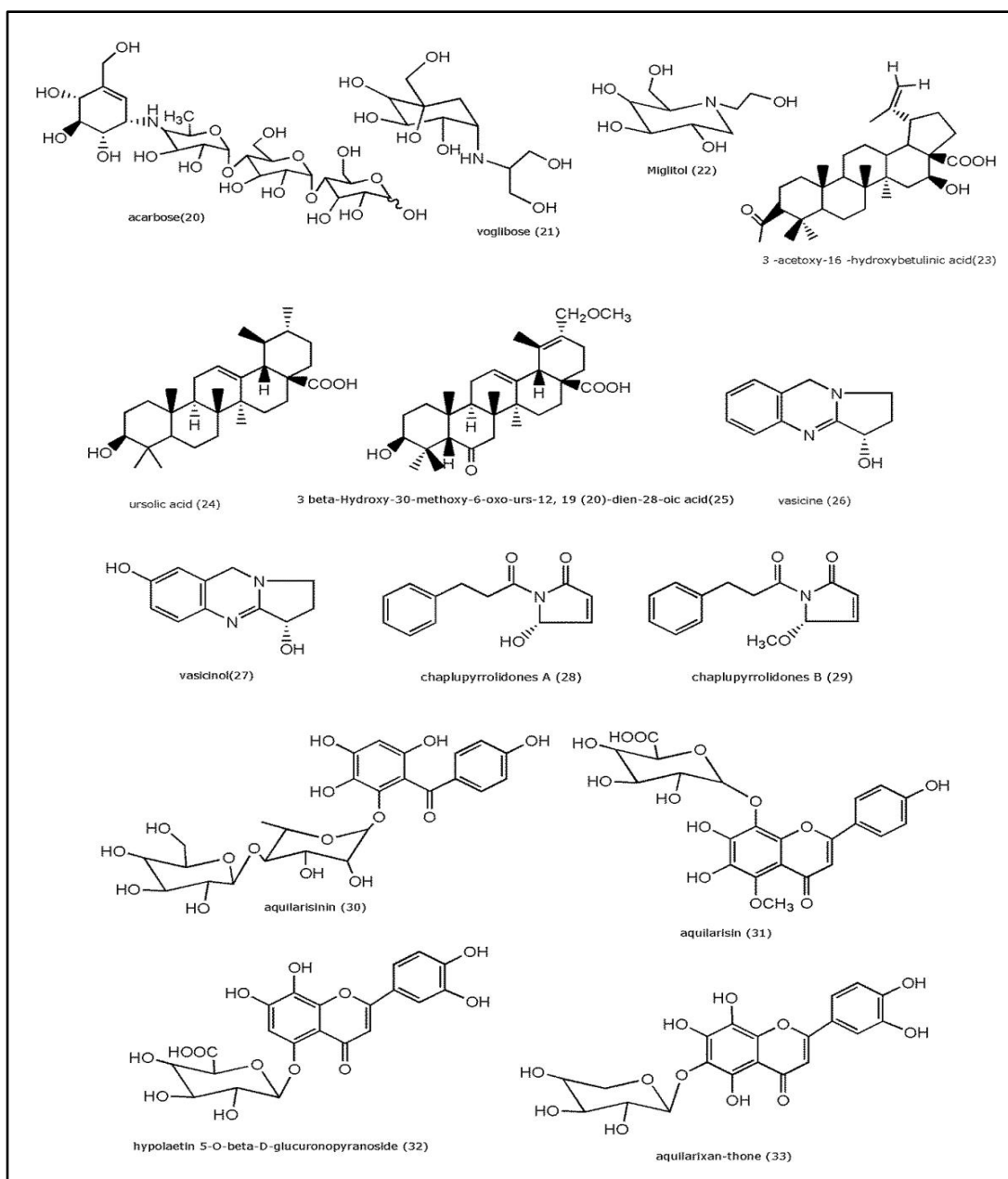


Figure 1.7. Alpha-glucosidase inhibitors isolated from medicinal plants

1.8 Microbial infections

Microbial infections are one of the important health hazards in developing countries including Pakistan, where they account for 12% of deaths (CDC, 2010). Both Gram-positive and Gram-negative bacterial strains including *Bacillus*, *Staphylococcus*, *Salmonella* and *Pseudomonas* are mainly involved in such infections (Ahameethunisa and Hoper, 2010). Poor hygiene, lack of proper health facilities and lack of knowledge are major reasons for this widespread health concern in developing countries. The antibiotics are mainly used as a remedy for bacterial infection; however, the cost of therapy, severe side effects like hypersensitivity, allergic reactions and immune suppression limit their usage. Furthermore prolonged and over-usage of antibiotics that generally leads towards development of resistance is of special concern these days (Alder, 2005; Walsh, 2004).

The increasing incidence of drug-resistant pathogens has drawn the attention of the pharmaceutical and scientific communities towards studies on the potential antimicrobial activity of plant-derived substances based on traditional use (Savoia, 2012). Literature surveys show that plant-based medicines play a promising role in the treatment of infectious diseases (Ahmad and Beg, 2001; Bibi *et al.*, 2010; Obeidat, 2011). Some antimicrobial compounds isolated from plants are shown in (Fig 1.8).

Also in Pakistan medicinal plants are considered as primary sources of medicines for the treatment of bacterial infections. A number of ethnopharmacological investigations conducted in various parts of the country have demonstrated the impact of traditional plants as sources of antimicrobial agents (Mahmood *et al.*, 2012_b; Ullah *et al.*, 2013; Ahmad *et al.*, 2015; Barkatullah *et al.*, 2015; Yaseen *et al.*, 2015). It has been reported that simple phenols, phenolic acids, quinones, flavonoids, tannins, alkaloids, coumarins, terpenoids, lectins, polypeptides and polyacetylenes are the major classes of antimicrobial compounds from plants (Cowan, 1999; Bibi *et al.*, 2011; Jamil *et al.*, 2012; Ejaz *et al.*, 2104; Ashraf *et al.*, 2015).

In recent years a number of interesting antibacterial metabolites have been isolated from different plant sources that exhibited significant activities. A naturally occurring 2,2': 5',2'' terthiophene (**34**) presented promising inhibitory activity against *S. aureus* (MIC = 0.08 μ M) and this activity mainly occurs in the presence of UV light (Ciofalo *et al.*, 1996).

The flavonoids and flavonones are well known for their antimicrobial potential. Some compounds including sophoraflavanone G (a geranylated flavonone) (**35**) also act synergistically with other molecules like vancomycin. In Thailand sophoraflavanone G was isolated from *Sophora exigua* Criab (Leguminosae), and presented promising activity against *S. aureus* (MIC = 7.3–14.7 μ M) (Tsuchiya *et al.*, 1996).

In a recent study methanolic extracts of *Oncoba spinosa* Forssk. (Salicaceae) were investigated and five flavonoids were isolated. Out of all, quercetin 3-O- α -L-rhamnopyranosyl- β -D-glucopyranoside (**36**) and quercetin (**37**) were the most active compounds against bacteria (MIC = 10.5- 21.1 μ M) and fungi (MIC = 212 – 423 μ M), respectively (Djouossi *et al.*, 2015).

In an investigation by Nissanka *et al.*,(2001) two alkaloids i.e 8-acetonyl-dihydranitidine (**38**) and 8-acetonyl-dihydroavicine (**39**) were isolated from the stem bark of *Zanthoxylum tetraspermum*. In the antimicrobial assays, 8- acetonyl-dihydranitidine (**38**) and 8- acetonyl-dihydroavicine (**39**) showed significant inhibition of bacterial growth, i.e MIC 3.8 μ M and 8.0 μ M, respectively, and MBC 61.6 μ M and 32.1 μ M, respectively (Nissanka *et al.*, 2001).

Coumarins are important secondary metabolites that possess a broad spectrum of pharmacological properties including antimicrobial activities. The dichloromethane extract of the fruits of *Heracleum mantegazzianum* Sommier & Levier (Apiaceae) was investigated for its antimicrobial activities and the isolation of active compounds. From the most active fraction xanthotoxin (**40**) and bergapten (**41**) were isolated. Xanthotoxin (**40**) showed a very high activity against gram-positive bacteria with very low MIC values (13.8 μ M), followed by bergapten (**41**) (MIC = 57.8 μ M).

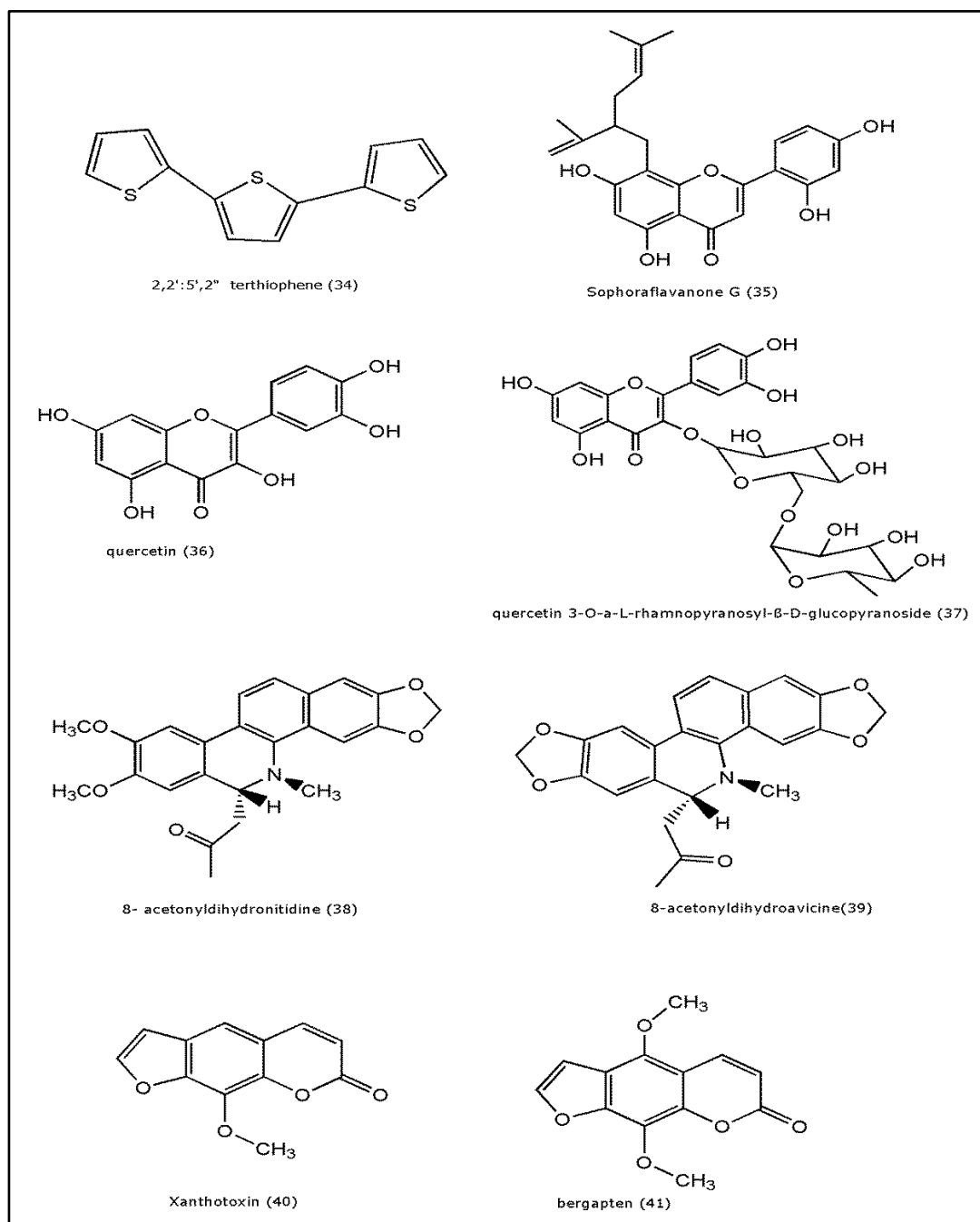


Figure 1.8. Antibacterial compounds isolated from medicinal plants

1.9 Leishmaniasis

Leishmaniasis, common in tropical and sub-tropical parts of the world, is a protozoal infection caused by single cellular, haemoflagellate protozoan parasites of the genus *Leishmania* (*Leishmania major*, *Leishmania tropica*, *Leishmania aethiopica*, *Leishmania mexicana*, *Leishmania amazonensis* and *Leishmania braziliensis*), transmitted by the bite of an infected female sandfly *Lutzomyia* or *Phlebotomus* (Handman and Bullen, 2002) (Fig. 1.9). Leishmaniasis is endemic in over 98 countries with more than 350 million people at risk. It is estimated that 1.3 million new cases of leishmaniasis (0.3 million visceral leishmania and 1.0 million cutaneous leishmania) occur every year (WHO, 2014a). Also in Pakistan leishmaniasis is reported as an emerging dermal disorder (Iqbal *et al.*, 2016).

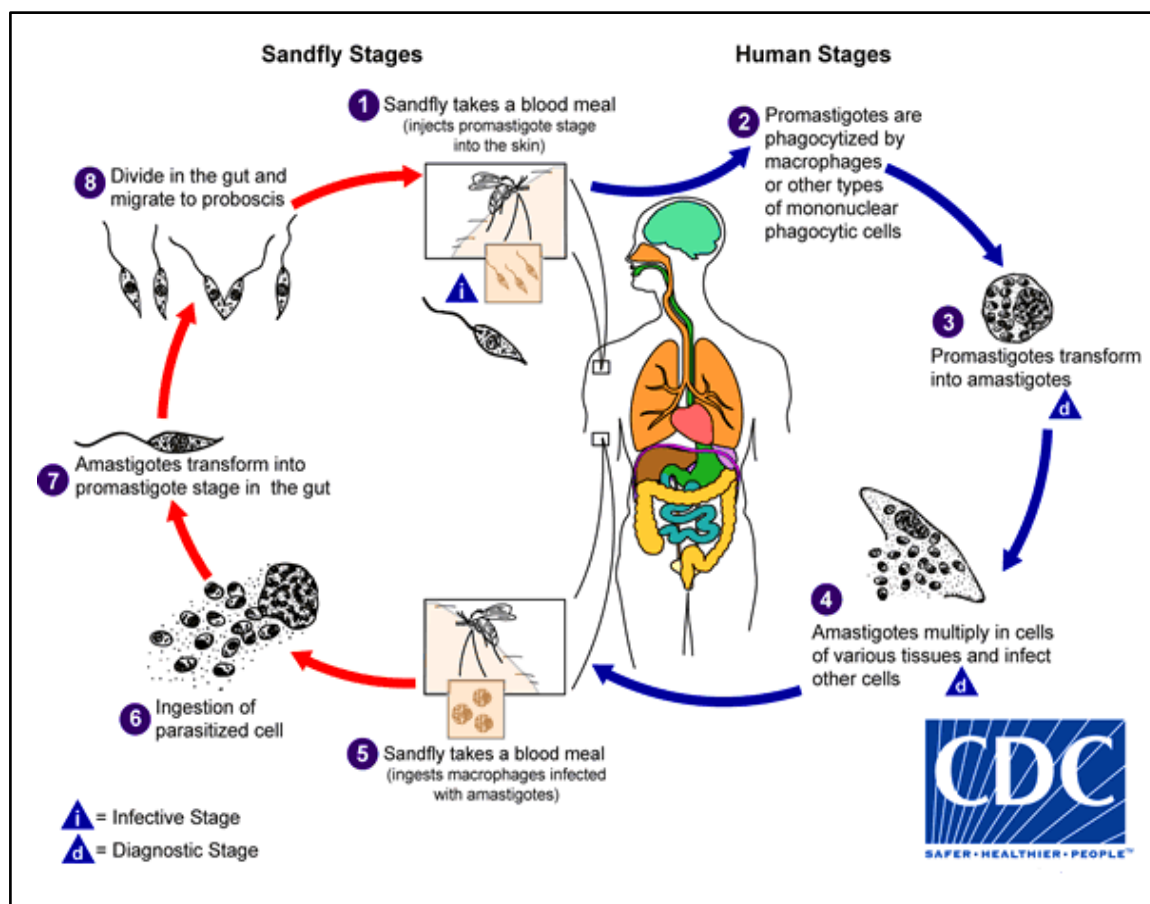


Figure.1.9 Life cycle of *Leishmania* sp. (Adopted from CDC, 2015)

In general the first-line treatment against leishmaniasis is still based on pentavalent antimonial drugs, such as sodium stibogluconate and meglumine antimoniate. However, these are associated with severe undesirable effects (Choudhary *et al.*, 2010). Plants are promising sources for new drug candidates. More than hundred plant species have been reported to possess antileishmanial activity (Kvist *et al.*, 2006).

The quest for new antileishmanial compounds is still continuing. For instance, the sesquiterpene lactone dehydrozaluzanin C (**42**) was isolated from the petroleum ether extract of leaves of *Munnozia maronii* (Asteraceae) and it inhibited the growth of eleven species of *Leishmania* promastigotes at concentrations between 2.5 and 10.0 µg/mL (Fournet *et al.*, 1993). More recently two new sesquiterpenes fnarthexol (**43**) and fnarthexone (**44**) were isolated from *Ferula narthex* in addition to some known compounds. Conferol (**45**) was found to be the most potent compound with IC₅₀ value of 30.1 µM, whereas newly isolated compounds i.e fnarthexol (**43**) (IC₅₀ 114.2µM) and fnarthexone (**44**) (IC₅₀ 123.4 µM) presented moderate activities (Bashir *et al.*, 2014) (Fig 1.10).

Flavonoids have a broad spectrum of activities including antileishmanial properties. Quercitrin (quercetin 3-O-α-L-rhamnopyranoside) (**46**) was isolated from *Kalanchoe pinnata* and tested for its antileishmanial activity. The compound presented a potent activity (IC₅₀ approximately 2.23 µM) with a low toxicity profile. This was the first demonstration of antileishmanial activity for a flavonoid glycoside (Muzitano *et al.*, 2006). In another investigation by Tasdemir *et al.* (2006) more than 100 compounds were screened for antileishmanial potential. It was reported that fisetin (**47**), 3-hydroxyflavone (**48**), luteolin (**49**) and quercetin (**50**) were the most potent compounds with IC₅₀ values of 2.2, 2.9, 2.7, and 3.3 µM, respectively.

In a recent study the antileishmanial activity of extracts and phyto-constituents of *Moringa oleifera* Lam. was investigated. The ethyl acetate fraction was found to be the most active with an IC₅₀ value of 27.5 µg/mL. From the ethyl acetate fraction, among other isolated compounds, niazinin (**51**), a thiocarbamate glycoside, was reported as highly active (IC₅₀ 5.25 µM) (Kaur *et al.*, 2014).

Quinones are well known for their antileishmanial activity. In a report by Mori *et al.* (2008) antileishmanial activity of cordiachromes (quinone derivatives) isolated from *Cordia fragrantissima* was determined. Among all the tested samples, cordiachrom A (52) and cordiachrom B (53) presented interesting leishmanicidal activity with IC_{50} 17.0 μ M and 10.2 μ M respectively.

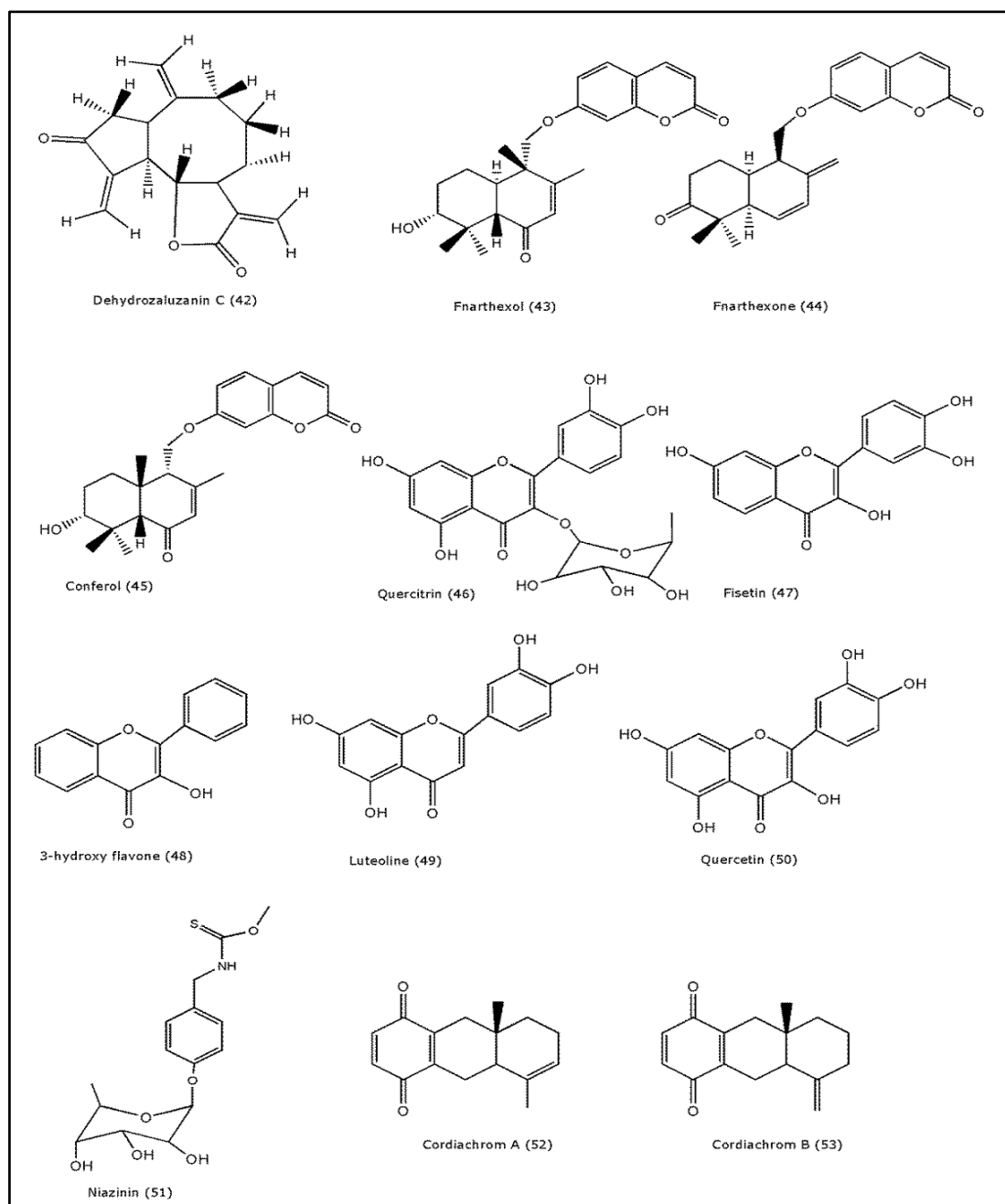


Figure.1.10 Antileishmanial compounds isolated from medicinal plants

1.10 Malaria

Malaria is caused by protozoan parasites from the genus *Plasmodium*, including *P. falciparum*, *P. vivax*, *P. malariae*, *P. ovale* and more recently the hitherto monkey malaria parasite *P. knowlesi* (White, 2008). Amongst all, *P. falciparum* is considered as the most lethal form. All *Plasmodium* parasites are transmitted to humans through female *Anopheles* mosquitoes (Diou *et al.*, 2009) (Fig. 1.11).

In Pakistan mainly *P. vivax* (responsible for approximately 64% of infections) and *P. falciparum* (causing 36% of infections) are involved in spreading malaria (Asif, 2008; WHO, 2012). Indeed, malaria accounts for 16% of the total disease burden of the country and therefore has become a serious health concern these days in both in rural and urban areas (Shah *et al.*, 2014). Regional statistics of WHO are even more alarming. It is estimated that the annual incidence of malaria is 1.5 million cases and Pakistan is placed in the “group 3 countries” of the WHO “Eastern Mediterranean” region, based on prevalence of malaria (WHO, 2014_b). Keeping in mind the severity of the issue, WHO has initiated a number of malaria control programmes in various parts of Pakistan especially the Khyber Pakhtunkhwa (KPK) and Balochistan Provinces.

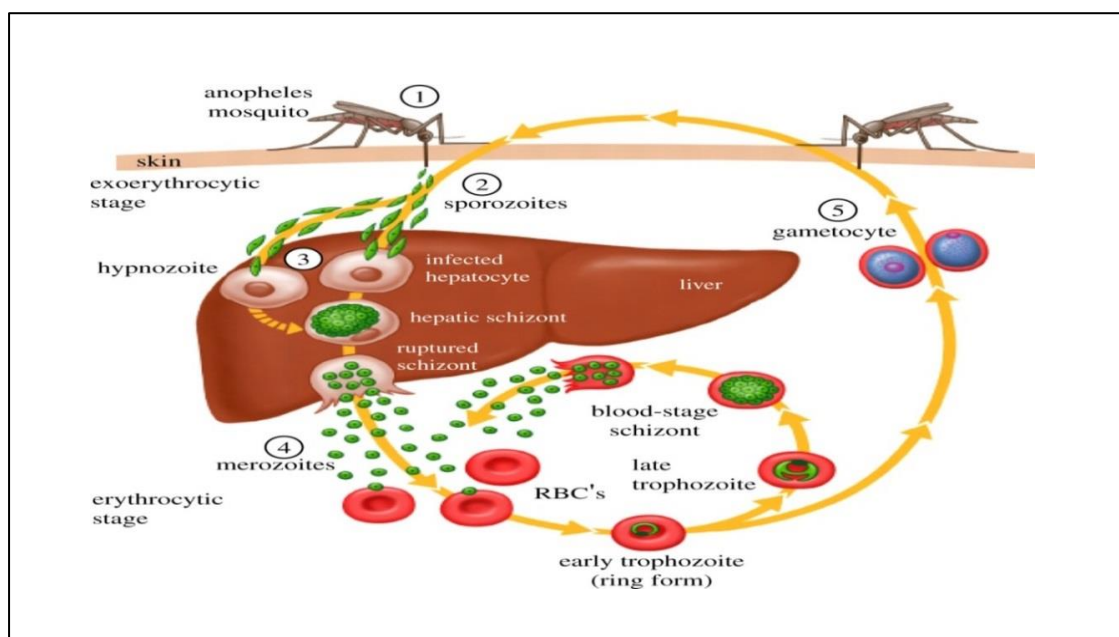


Figure. 1.11 Life cycle of malarial parasite (Adopted from Hill, 2011)

In general, a large number of anti-malarial drugs are commercially available to treat malaria all over the world. However, they are associated with dose-related side effects such as vomiting, nausea, fatigue, pulmonary toxicity, neuropsychiatric effects, blindness and neurotoxicity. In addition, drug resistance against the malarial protozoan is a serious concern these days, which is responsible for failure of therapy in many cases (Ruiz-Irastorza *et al.*, 2010, Gravani *et al.*, 2013).

Medicinal plants and their constituents have a long history of effective use in prevention and treatment of malaria in various parts of the world, and are still considered as promising sources for new antimalarials (Chiyaka *et al.*, 2009). Also a wide majority of rural communities in Pakistan still rely on medicinal plants and herbal formulations for the treatment of malaria (Irshad *et al.*, 2011; Marwat *et al.*, 2011a; Shah *et al.*, 2014; Sher *et al.*, 2015). From literature it is obvious that diterpenes, alkaloids, flavonoids, coumarins, steroids and phenolics are mainly responsible for antimalarial activity of many plant extracts (Fig. 1.12)

Abietane type diterpenes were isolated from an apolar *n*-hexane fraction of *Plectranthus barbatus* and tested against erythrocytic schizonts of *Plasmodium falciparum*, intracellular amastigotes of *Leishmania infantum* and *T. cruzi* and free trypomastigotes of *T. brucei*. The compound 5,6-didehydro-7-hydroxy-taxodone (**54**) showed promising activity with acceptable selectivity against *P. falciparum* (IC₅₀ 9.2 μM, SI 10.4) and *T. brucei* (IC₅₀ 1.9 μM, SI 50.5) supporting its effectiveness against important protozoans (Mothana *et al.*, 2014).

Also several plant-derived alkaloids have shown very interesting antimalarial activities and thus have a great potential for anti-malarial drug development. The indole alkaloids are famous for their antimalarial activities. Nkunya *et al.* (2004) explored *Monodora angolensis* and *Isolona cauliflora* (Annonaceae) for their antimalarial activities. They finally isolated a series of prenylated indole alkaloids from both plants. It was reported that 6-(3-methyl-but-2-enyl)-1,3-dihydro-indol-2-one(**55**) and annonidine F (**56**) showed promising antimalarial activities against the multidrug resistant strain K1 of *P. falciparum* (IC₅₀ 104 μM and IC₅₀ 57.0 μM for each compound).

Naphthoisoquinolines are characterised by a unique structure with a linkage between the naphthalene and the isoquinoline portions. It is hypothesised that this characteristic structure is mainly responsible for antimalarial activity. In Kenya naphthoisoquinoline alkaloids were isolated from the stems and the leaves of *Ancistrocladus robertsoniorum*. All isolated compounds were found active, however ancistrocladisine (**57**) (IC₅₀ 1.4 µM), cis-dihydro ancistrocladisine (**58**) (IC₅₀ 1.7 µM) and ancistrobrevine B (**59**) (IC₅₀ 2.0 µM) presented significant activities against the *P. falciparum* K-1 strains (Bringmann *et al.*, 1999).

The flavonoids are commonly found in medicinal plants and account for an array of biological activities including antimalarial. It is considered that flavonoids act by inhibiting the fatty acid biosynthesis (FAS II) of the parasite. Several antimalarial flavonoids have been isolated from the stem bark of *Erythrina abyssinica* by Yenesew *et al.*(2004). These include chalcones, prenylated and non-prenylated isoflavones and flavones, pterocarpenes and flavenes. All compounds exhibited moderate antimalarial activity against the D6 and W2 strains of *P. falciparum*. Abyssinone V (IC₅₀ 4.9 µM) (**60**) and sigmoidin A (**61**)(IC₅₀ 5.8 µM) presented significant antimalarial activities.

Bioassay-guided fractionation of the ethanolic extract of the leaves and twigs of *Piptocoma antillana* (Asteraceae) was performed and a number of compounds were isolated. All compounds were tested for their antiproliferative and antimalarial potential. Two new sesquiterpene lactones i.e 5-O-methyl-5-epiisogoyazensolide (**62**) and 15-O-methylgoyazensolide (**63**) displayed significant (IC₅₀ 6.2 µM) and (IC₅₀ 2.2 µM) activity against *Plasmodium falciparum*.

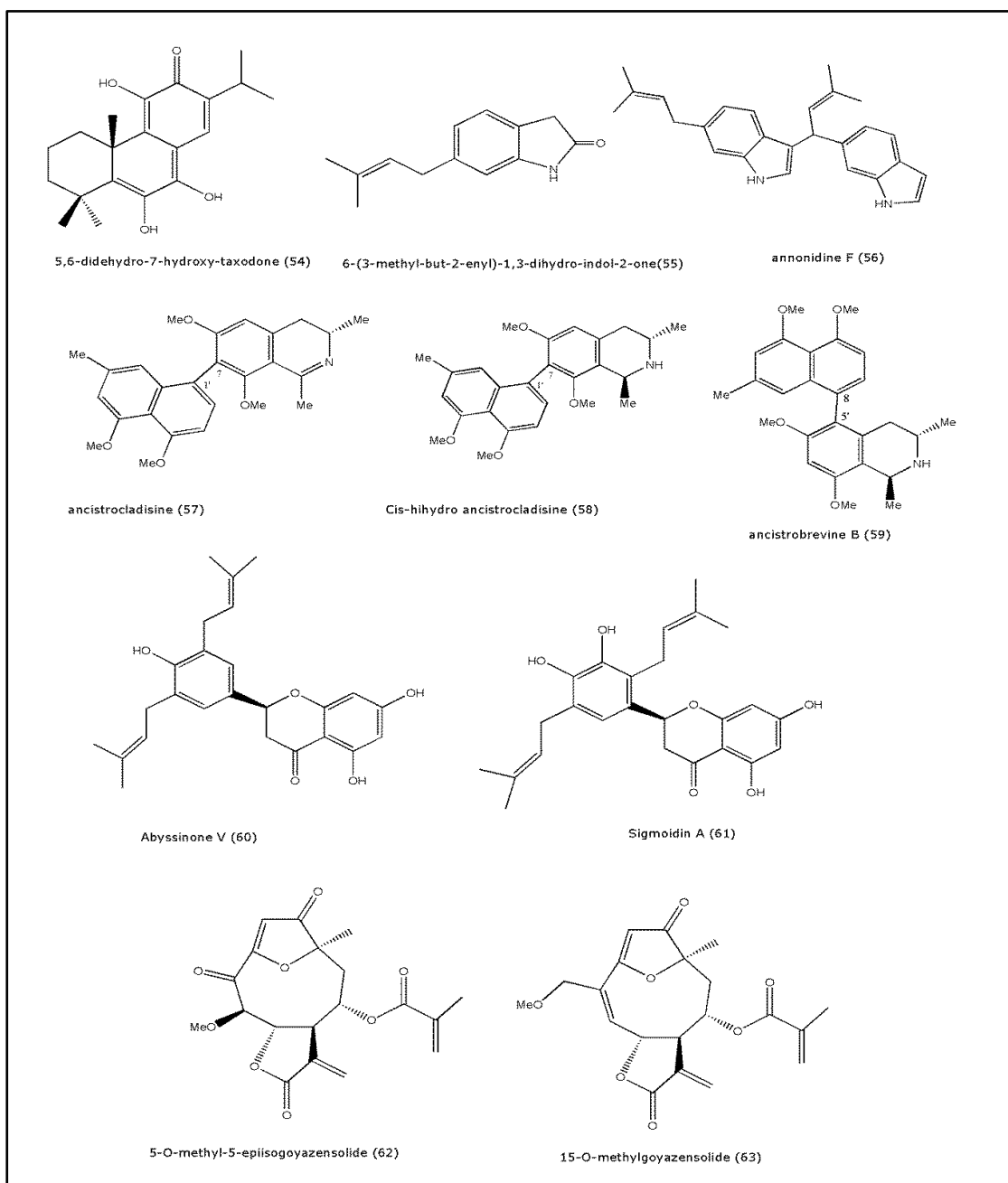


Figure 1.12 Antiprotozoal agents from plant origin

1.11 AIM OF THIS WORK

The aim of this thesis was to perform phytochemical and biological investigations on some selected medicinal plants from Pakistan.

Based on the importance of traditional medicine in Pakistan and various ethnopharmacological surveys, selected medicinal plants were collected from different phytogeographical regions of Khyber Pakhtunkhwa (KPK) province of Pakistan, including Dera Ismail Khan (wetlands), District Karak (hilly and sandy area) and Chitral (Himalaya region) (Fig 1.13).

D.I.Khan has an area of about 7326 km² and is situated between 70°11' and 71°20' E. longitude and between 31°15' and 32°32' N. latitude. D.I.Khan has mostly flat dry plains, commonly called Daman, about 80% of the total area, where a large number of streams and hill torrents discharge water (Chaudhry, 1998). The city D.I.Khan is located on the river Indus. Aquatic and xerophytic vegetation are commonly found in this study area, which is part of the country's richest biodiversity centre and a source of ethnobotanical knowledge. A number of ethnomedicinal surveys have reported the occurrence of important medicinal plants in this area (Marwat *et al.*, 2008; Marwat *et al.*, 2011a; Marwat *et al.*, 2011b; Mussarat *et al.*, 2014).

The district **Karak** covers an area of 3372 km² and is located at 33° 7'12N. latitude and 71° 5'41E. longitude. It is a semi-arid area that mainly consist of xerophytes and mesophytes. The study area is comprised of mountainous and plain area with mostly sandy soils. A number of ethnomedicinal surveys have been carried out in this area regarding the occurrence of medicinal plants (Khan *et al.*, 2013; Murad *et al.*, 2013; Khattak *et al.*, 2015, Adnan *et al.*, 2015; Rehman *et al.*, 2015).

The **Chitral** is the largest district (14850 km²) covering 20% of the total KPK province. It lies within 35°15'06" to 36°55'32"N. latitude and 71°11'32" to 73° 51' 34"E. longitude. The elevation of the area varies from about 1,070 m (about 3500 ft) in the extreme south in Arandu to 7,690m (25,230 ft) at the summit of Tirich Mir in the Hindu

Kush. The local community of Chitral has a centuries old tradition of using plants for the ailments of various medical problems and therefore people have typical methods of preparation of herbal medicines (Ali and Qaiser, 2009). Despite the ethnomedicinal importance of the area, only limited information is available (Awan *et al.*, 2001; Hussain *et al.*, 2007; Ali and Qaiser 2009; Khan *et al.*, 2011; Shah and Hussain, 2012; Abbasi *et al* 2013; Hadi *et al.*, 2013).

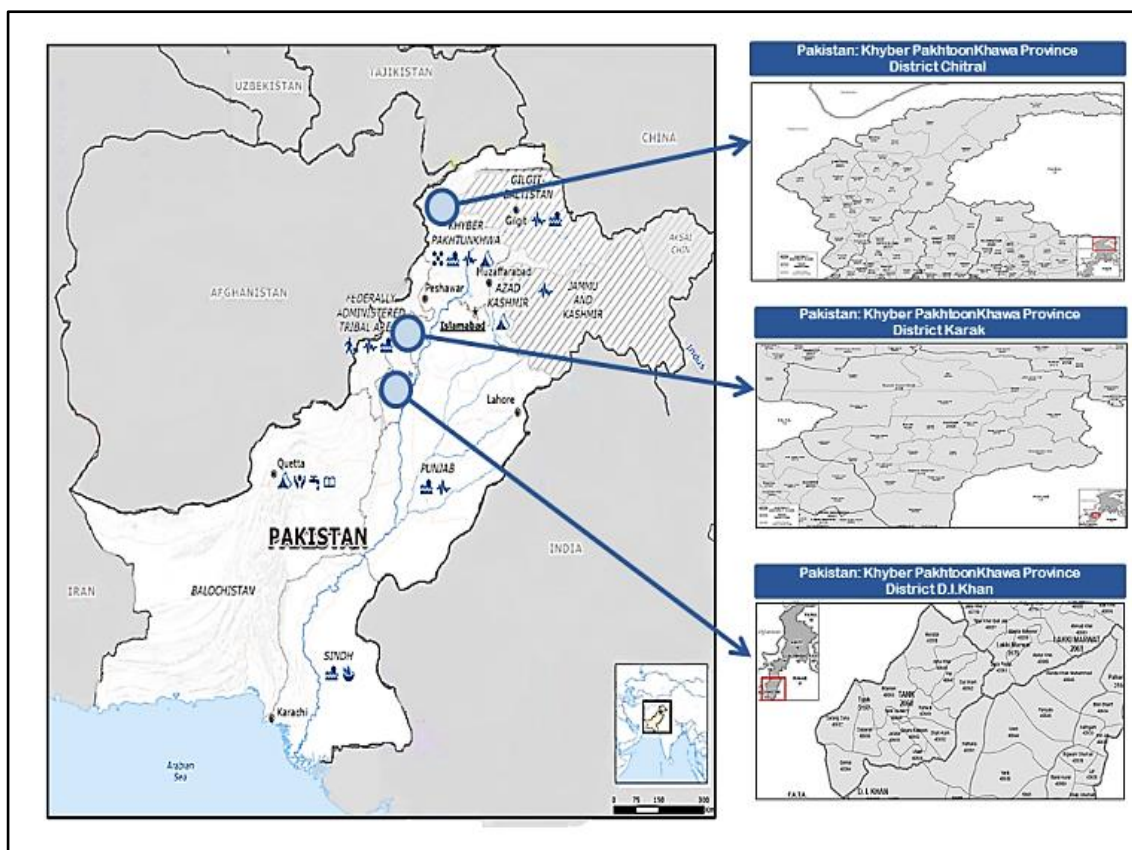


Figure 1.13. Map of Pakistan showing the sampling areas (Adopted from OCHA, Pakistan, <http://www.unocha.org/pakistan>)

In the present work some less evaluated medicinal plants from Pakistan have been selected in order to investigate their phytochemical constituents and biological properties. This selection was based on their extensive traditional use and on most recent ethnopharmacological surveys. The pharmacological focus was on their potential therapeutic usefulness in diabetic conditions, and on their antimicrobial properties

(antibacterial, antifungal, antileishmanial and antimalarial), according to running programmes in the host laboratory. In this way the following plant species were selected: *Nymphoides indica*, *Kickxia ramosissima* and *Ferula narthex*.

***Nymphoides indica* (L.) Kuntze** (Menyanthaceae) (**Chapter 3**) is a free floating leaved aquatic herb. It is commonly used as vegetable in local communities. A number of ethnomedicinal surveys reported its medicinal importance in various communities. In general it is used as one of the important ingredients of some traditional formulations. Most commonly *N. indica* is used for treatment of dysentery, scabies, jaundice and as antiproliferative, indicating the potential presence of antimicrobial and liver-protective constituents (Kensa, 2011; Panda and Misra, 2011; Kitdamrongtham *et al.*, 2013). Previously flavonoids (Bohm *et al.*, 1986) were reported from *N. indica*; however, no comprehensive phytochemical and pharmacological data was available in view of its medicinal importance.

***Kickxia ramosissima* (wall.) Janch**(Scrophulariaceae) (**Chapter 4**) is a small perennial herb, mainly found on rocky areas with shady places. It is commonly used for a number of ailments; for instance, as diuretic, to remove kidney stones, against fever, in rheumatism and in the management of snake and scorpion bites (Jain *et al.*, 2008; Bole and Pathak, 1988; Pandya *et al.*, 2012). In addition, ethnopharmacological surveys suggest that traditionally the plant has been used as an effective remedy for diabetes mellitus (Ahmad *et al.*, 2009; Qureshi *et al.*, 2010; Patel and Sachdeva, 2014). Limited phytochemical and pharmacological work has been published in the past and a few flavonoids, iridoids and mannitol were isolated (Khan *et al.*, 1991; Khan and Aqil, 1993). In view of its use against fever and in diabetic conditions, the potential presence of antimicrobial, alpha-glucosidase and AGEs inhibitors, respectively, can be hypothesised.

***Ferula narthex* Boiss.** (Umbelliferae) (**Chapter 5**) mainly grows on high altitude areas. The oleo-gum resin obtained from this plant species is used as traditional medicine. It is a very important part of Ayurveda formulations like Hingvaday and Ghrita. It has been used as a spice and as well as folk phytomedicine for centuries for the treatment of gastro-intestinal disorders, nervous disorders, respiratory problems, as anti-diabetic and to treat impotence (Kapoor, 2001; Abuzaitoon, 2010). A number of compounds

have been previously isolated from *Ferula* species including coumarins, various types of terpenes (sesquiterpenes and other terpenoids) and sulfur-containing compounds (Lee *et al.*, 2009). However, in the case of *Ferula narthex* limited work is available. More recently antileishmanial compounds have been isolated from this plant (Bashir *et al.*, 2014). Therefore in the current project *Ferula narthex* was selected to isolate potentially antidiabetic and antimicrobial compounds.

The investigations accomplished in this project on different medicinal plants of Pakistan embraced:

- a) Phytochemical characterization, i.e isolation and identification of constituents by using various chromatographic and spectroscopic techniques;
- b) *In vitro* antidiabetic (inhibition of alpha-glucosidase, antioxidant activity, inhibition of AGEs) studies of extracts and isolated compounds;
- c) *In vitro* antimicrobial studies of extracts and isolated compounds.

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CHAPTER 2

GENERAL EXPERIMENTAL METHODS

2.1 CHROMATOGRAPHIC METHODS

2.1.1 Solvents and reagents

All solvents including *n*-hexane, ethyl acetate, *n*-butanol, methanol, absolute ethanol (99.99%), isopropyl alcohol and methylene chloride were analytical grade and obtained from Fisher Scientific (Leicestershire, UK) and Acros Organics (Geel, Belgium). All reagents such as acetic acid glacial ($\geq 97\%$), sulphuric acid ($\geq 95\%$), hydrochloric acid (HCL) ($\geq 95\%$), potassium hydrogen phosphate ($\geq 98\%$), potassium dihydrogen orthophosphate ($\geq 95\%$), disodium hydrogen phosphate ($\geq 99\%$), sodium hydrogen phosphate ($\geq 95\%$), sodium hydroxide ($\geq 98\%$), sodium carbonate (99%), formic acid ($\geq 98\%$), dimethyl sulfoxide (DMSO) ($\geq 95\%$) and trichloroacetic acid (TCA) ($\geq 99\%$) were purchased from Acros Organics or Sigma-Aldrich (St. Louis, MO, USA). The solvents for HPLC i.e methanol (MeOH) and acetonitrile were HPLC grade and purchased from Fisher Scientific (Leicestershire, UK). RiOS water was prepared by reverse osmosis and water for HPLC was dispensed by a Milli-Q system, both from Millipore (Bedford, MA, USA). Water was passed through a 0.22 μm membrane filter before usage. Solvents used for GC-MS were purchased from Sigma-Aldrich including BSTFA + TMCS, 99:1 (N,O-bis (trimethylsilyl) trifluoroacetamide and trimethylchloro silane) (Supelco, USA), pyridine (dry, 99.8%) (Seccosolv).

The reference materials such as aminoguanidine ($\geq 99\%$), quercetin ($\geq 98\%$), glucose ($\geq 99\%$), 2,2'-azobis(2-methylpropionamide) dihydrochloride (AAPH) ($\geq 97\%$), Trolox ($\geq 97\%$), linoleic acid (99%), bovine serum albumin (BSA) ($\geq 98\%$), methylglyoxal (MGO) ($\geq 99\%$), 2,2-diphenyl-1-picrylhydrazyl (DPPH) ($\geq 99\%$), α -glucosidase (≥ 100 units/mg protein), Pnp (*p*-nitrophenyl- α -D-glucopyranoside ($\geq 99\%$), acarbose ($\geq 95\%$), sodium azide (99%) and 15-lipoxygenase from *Glycine max* (soybean) ($\geq 50,000$ units/mg solid) were purchased from Sigma-Aldrich (St. Louis, MO, USA).

2.1.2 General extraction scheme

The standard extraction scheme used during the project is shown below (Fig. 2.1).

2.1.3 Thin layer chromatography

Analytical plates for thin layer chromatography (TLC) were silica gel 60 F₂₅₄ plates (20 x 20 cm) for normal phase (Merck, Darmstadt, Germany). The spraying reagent *p*-anisaldehyde was prepared by mixing 0.5 mL *p*-anisaldehyde (Sigma-Aldrich) with 10 mL glacial acetic acid, 85 mL methanol and 5 mL sulphuric acid.

Dragendorff's reagent was prepared by mixing solution A and B. Solution A was prepared by adding of 1.7 g bismuth subnitrate in 100 mL water/acetic acid (4:1). Solution B was prepared by dissolving 40 g potassium iodide in 100 mL water. Solution A and B were combined, as follows: 5 mL A + 5 mL B + 20 mL acetic acid + 70 mL water.

The cerium sulphate reagent was prepared by mixing 1g CeSO₄ and 10g of trichloroacetic acid (TCA) with 4 mL of 15% H₂SO₄ initially. This was mixed thoroughly until a slurry was formed. Then the slurry was stirred using a magnetic stirrer on a hot plate at 60 °C and 15% H₂SO₄ solution was slowly added . The process was continued until a clear solution was obtained.

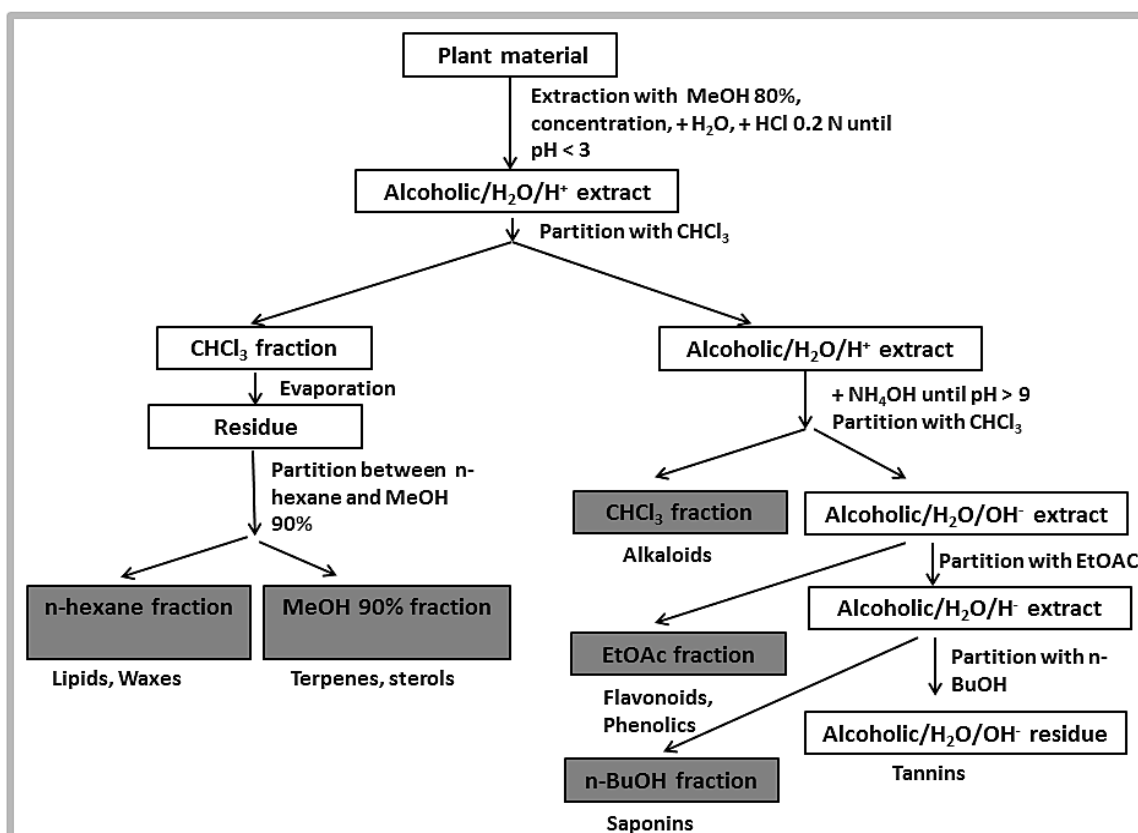


Figure 2.1 Schematic presentation of liquid-liquid fractionation

2.1.4. Flash Chromatography

Flash column chromatography was performed on a Reveleris iES system from Grace (Columbia, MD, USA) using the Reveleris® Navigator™ software. The system is equipped with a binary pump with four solvent selection, an ultraviolet (UV) and evaporating light scattering detector (ELSD) and a fraction collector. The column used was a pre-packed Flash Grace Reveleris silica cartridge (80 g) with a particle size of 40 µm. The ELSD carrier solvent was isopropyl alcohol.

2.1.5. High performance liquid chromatography

HPLC analysis was carried out on an Agilent® 1200 series system with degasser, quaternary pump, automatic injection, thermostatic column compartment and a diode array detector (DAD) (Agilent® Technologies, Santa Clara, California, USA). A silica based Gracesmart C₁₈ column (250 x 4.6 mm, 5 µm) (Grace Vydac, USA) and Phenomenex luna C₁₈ (250 x 4.6 mm, 5 µm) (Phenomenex, Torrance, CA, USA) was used together with a suitable precolumn to endure the lifetime of columns.

2.1.6. Semi-preparative high performance liquid chromatography

The isolation of compounds was carried out on a semi-preparative HPLC-DAD-MS system (Waters®) using a Luna 5µm (C₁₈) 100A 250 X 10.0 mm column (Phenomenex™) and Masslynx® 4.1 software. The system was equipped with HPLC pump 515 (Waters™ 2767), make-up pump (Waters™ 511), system fluid organiser (SFO), diode array detector (DAD) (Waters™ 2998), triple quadrupole mass spectrometer (TQD-MS) and automatic fraction collector was used to isolate compounds.

2.2. SPECTROSCOPIC METHODS

2.2.1 Nuclear magnetic resonance spectroscopy

NMR spectra were recorded on a Bruker DRX-400 instrument (Rheinstetten, Germany), operating at 400 MHz for ¹H and at 100 MHz for ¹³C, using a 3-mm broadband inverse (BBI) probe or a 5-mm dual ¹H/¹³C probe using standard Bruker pulse sequences. “Distortionless Enhancement by Polarization Transfer” spectra (DEPT-135, DEPT-90) were also recorded. The chemical shifts are given in ppm and coupling constants (*J*) in Hz. The multiplicity is indicated as s for singlet, d for doublet, t for triplet, q for quartet

and m for multiplet. Additionally two dimensional NMR experiments (2D-NMR) were performed where necessary. The 2D-NMR experiments included correlation spectroscopy (COSY) for ^1H - ^1H correlation, heteronuclear single quantum coherence (HSQC) for direct ^1H - ^{13}C correlation and heteronuclear multiple bond coherence (HMBC) for long-range ^1H - ^{13}C correlation. Structure elucidation was assisted by a ^{13}C -NMR database (NMR Predict version 4.8.57, Modgraph[®]). Deuterated solvents including CDCl_3 (99.8% D), DMSO-d_6 (99.9 % D) and CD_3OD (99.8% D) were purchased from Sigma-Aldrich.

2.2.2 Mass Spectrometry

High resolution mass spectra was obtained with an Agilent[™] 6530 quadrupole-time-of-flight mass spectrometer (QTOF-MS) equipped with an Agilent[™] Jetstream source that was used with the following parameters: gas temperature 325 °C, sheath gas flow 11L/min. Capillary, fragmentor and skimmer voltages were set to 3500 V, 150 V and 65 V, respectively, and OCT 1RF Vpp was set at 750 V. The mass spectrometer was operated in positive and negative ion mode at 20000 resolution. The instrument was calibrated and tuned with a tune mix (G1969-85000) and during acquisition the accuracy was monitored by using ES-TOF reference mass solution kit (G1969-85001) from Agilent[™]. Mass Hunter[®] (Agilent[™] Technologies) software was used for acquisition and processing.

2.2.3 GC-MS analysis

GC experiments were performed on a TRACE[®] 2000 Ultra GC (Thermo Scientific) on a column AT-5MS (Grace): length 30 m; ID: 0.25 mm; film thickness: 0.25 μm . Derivatisation was performed on dried samples using 20 μL of BSTFA + TMCS, 99 : 1 (Supelco, USA) in pyridine (dry) (Seccosolv) in a molar ratio 2-1, by heating for 1h at 70°C.

2.2.4 Optical rotation

The specific optical rotation was determined on a Jasco P-2000 polarimeter. The samples were dissolved in methanol and optical rotation was recorded at 589 nm with a path length of 50 mm.

2.2.5 Others

UV-Vis (ultraviolet-visible light) absorbance was measured on a Genesys-10UV (Thermoscientific) spectrophotometer. Fluorescence (excitation 335 nm, emission 385 nm) was measured on a Tecan™ Infinite M200 spectrofluorometer.

2.3 PHYTOCHEMICAL SCREENING

Phytochemical screening for various compound classes was carried out on every fraction using published methods with slight modifications (Sofowora, 1993, Trease and Evans, 1989, Herborne, 1973).

2.3.1 Chemical test for alkaloids

About 0.2 g of the extract was added to 2 mL of 2% H₂SO₄ and the mixture was slightly heated on a water bath for 2-3 min. The mixture was filtered and 4-5 drops of Dragendorff's reagent were added. Orange-red precipitate indicated the presence of alkaloids.

2.3.2 Chemical test for tannins

About 0.2 g of extract was mixed with 10 mL distilled water in a test tube and heated on a water bath. The mixture was filtered and 4-5 drops 0.1% ferric chloride (FeCl₃) were added to the filtrate. A dark green or blue-green coloration indicated the presence of tannins.

2.3.3 Chemical test for saponins

About 0.2 g of the extract was shaken with 5 mL of distilled water and then heated to boiling. Appearance of foam indicated the presence of saponins.

2.3.4 Chemical test for flavonoids

About 0.2 g of extract was dissolved in 2 mL of 10% NaOH. A change from yellow to a colourless solution observed after addition of 0.5 mL HCl, indicated the presence of flavonoids.

2.3.5 Chemical test for steroids

About 0.2 g of the extract was mixed with 2 mL of acetic anhydride. To this mixture, 1 mL of H₂SO₄ conc. was added. The colour change from violet to blue or green indicated the presence of steroids.

2.3.6 Chemical test for terpenoids (salkowski test)

About 0.2 g of extract was mixed with 2mL of chloroform and 3mL H₂SO₄ conc. was carefully added to form a layer. A reddish-brown colouration of the interface indicated the presence of terpenoids.

2.3.7 Total phenolic contents (TPC)

The total phenolic content (TPC) of all fractions was estimated using Folin-Ciocalteu's reagent according to Bursal and Gulcin (2011) with slight modifications. Briefly, 10 µL of each sample (0.2 mg/mL) or reference compound (gallic acid 0.0156–0.5 mg/mL) was mixed with 100 µL of Folin-Ciocalteu's reagent, and incubated at room temperature and after 5 min, 90 µL of 10% Na₂CO₃ was added to the above mixture. These mixtures were then incubated at room temperature for 40 min. The absorbance was measured at 765 nm. TPC was expressed as gallic acid equivalents (GAE). Based on findings of Dudonne *et al.* (2009), TPC were categorized as very high (> 300 mg GAE/g), high (200 - 300 mg GAE/g), moderate (50-200 mg GAE/g), low (15-50 mg GAE/g), very low (<15 mg GAE/g).

2.3.8 Total flavonoid contents (TFC)

The total flavonoid content (TFC) of all fractions was assessed according to Sun *et al.* (2011). Briefly, 20 µL of each sample (1 mg/mL) or reference compound (rutin) was mixed with 30 µL of 5% sodium nitrite. After 6 min, 50 µL of a 10% AlCl₃ solution was added, and the mixture was kept untouched for 5 min. Next, 100 µL of a 10% NaOH solution was added, the mixture was incubated at 25°C for 15-20 min, and the absorbance was measured at 510 nm. TFC was expressed as rutin equivalents (RUE). TFC were categorized as very high (>300 mg RUE/g), high (200-300 mg RUE/g), moderate(50-200 mg RUE/g), low (15-50 mg RUE/g), very low (<15 mg RUE/g).

2.4. BIOLOGICAL ACTIVITIES

2.4.1 Antiglycation assay

Glucose-mediated protein glycation models are generally used to determine AGE inhibition. However, AGE-protein adducts can be formed both under oxidative and non-oxidative conditions, and therefore AGEs inhibitors should be differentiated from common antioxidants (Rahbar, 2007). Evaluation of the inhibition of AGEs formation starting from glucose and BSA involves all possible mechanisms. However, Reactive Carbonyl Substances (RCS) such as glyoxal and methyl glyoxal that are intermediate products of the Maillard reaction, have already undergone the oxidation processes starting from glucose. Inhibition of their reaction with BSA is representative of non-oxidative glycation reactions.

2.4.1.1 Antiglycation assay (BSA-Glucose assay)

The antiglycation assay was performed according to the method developed by Matsuura *et al.* (2002) with minor modifications. The reaction mixture (300 μ L) contained bovine serum albumin (BSA) (10 mg/mL, 135 μ L), D-glucose (500 mM, 135 μ L) dissolved in phosphate buffer (50 mM, pH 7.4, containing sodium azide (0.02%, added to prevent bacterial growth) and test compounds (30 μ L), at different final concentrations (1.5 – 0.023 mM for isolated constituents or 100–1.17 μ g/mL for extracts) dissolved in 100 % DMSO. The mixtures were incubated at 60°C for 48 h. After incubation, the reaction mixture was allowed to cool down at room temperature. Then 100 μ L reaction mixture was transferred to a new plastic tube (1.5 mL) and the reaction was stopped by adding 10 μ L of 100% (w/v) trichloroacetic acid (TCA) and to precipitate proteins. The TCA-added mixture was kept at 4°C for 10 min, followed by centrifugation (14000 rpm, 4 °C, 4 min). The supernatant containing unreacted D-glucose, test sample and interfering substances was discarded, whereas the precipitate containing AGEs-BSA was redissolved with 0.8 mL alkaline PBS (137 mM NaCl, 8.1 mM Na₂HPO₄, 2.68 mM KCl, 1.47 mM KH₂PO₄, pH 10). The change in fluorescence intensity (excitation 335 nm, emission 385 nm; excitation 370, emission 440) due to formation of AGEs was monitored by spectrofluorometry (Tecan™ Infinite M200, Giessen, The Netherlands). In order to eliminate interference by autofluorescence of the test compounds, a parallel incubation of test substance with BSA at 60°C without D-glucose was performed for all

samples. Quercetin (0.5- 0.0312 mM final concentration) and aminoguanidine (3 - 0.046 mM final concentration) were used as reference compounds.

The AGEs inhibition was calculated as

$$\% \text{ inhibition} = \{1 - (F_{\text{BSA} + \text{glucose} + \text{test substance}} - F_{\text{BSA} + \text{test substance}}) / (F_{\text{BSA} + \text{glucose}} - F_{\text{BSA}})\} \times 100$$

where F is the fluorescence intensity. The concentration required for 50% inhibition (IC₅₀) was calculated using Sigma plot® 13.0. The AGEs inhibition was categorized as mild (1-25%), moderate (26-50%), high (50-70%), and very high (70-100%).

2.4.1.2 Protein-glyoxal interaction (BSA-MGO assay)

The antiglycation assay using methylglyoxal (MGO) was performed according to the method developed by Peng *et al.* (2007) with slight modifications. Briefly, methylglyoxal (135 µL, 5.75 mM) was incubated with BSA (135 µL, 10 mg/mL) dissolved in phosphate buffer (50 mM, pH 7.4, containing sodium azide (0.02%), added to prevent bacterial growth) and test compounds (30 µL) at different final concentrations (1.5 – 0.023 mM) in 100% DMSO. The reaction mixture was incubated at 37 °C for one week. Control solutions only contained methylglyoxal (135 µL, 5.75 mM) BSA (135 µL, 10mg / mL) and DMSO (100%) only. The blank samples of similar composition were also prepared simultaneously and kept at 4 °C for one week. The change in fluorescence intensity (excitation 335 nm, emission 385 nm; excitation 370, emission 440) due to formation of AGEs was monitored by spectrofluorometry (Tecan® Infinite M200, Giessen, The Netherlands). Aminoguanidine (3-0.046 mM final concentration) was used as reference compound ,

The AGEs inhibition was calculated as

$$\% \text{ AGEs inhibition} = [1 - (S - S_b) / (C - C_b)] \times 100$$

where S and C were fluorescence of test samples (in DMSO) and control (test mixtures containing only DMSO) incubated at 37 °C, and where S_b and C_b were fluorescence for

samples incubated at 4 °C. The concentration required for 50 % inhibition (IC₅₀) was calculated using Sigma plot® 13.0.

2.4.2 Inhibition of α-glucosidase.

α-Glucosidase inhibitory activity was assayed according to method adopted by Choudhary *et al.* (2010) with some modifications. Briefly, 50 µL of a solution of α-glucosidase from *Saccharomyces cerevisiae* (0.2 U/mL in 0.1 M phosphate buffer at pH 6.8, Sigma-Aldrich, St. Louis, MO, USA) was incubated with 20 µL of test compounds or extracts at different concentrations (ranging from 6 mM – 0.372 mM in 25% DMSO for isolated constituents or from 5 mg to 0.039 mg/mL for extracts) at 37 °C for 10 min. Then 50 µL of the substrate, *p*-nitrophenyl-α-D-glucopyranoside (0.7 mM stock, final concentration 0.29 mM) was added to the reaction mixture and incubated again for 30 min at 37 °C. The final concentrations of test substances in the reaction mixture ranged from 1 – 0.062 mM for isolated constituents, and from 834 µg to 6.5 µg/mL for extracts. The reaction was stopped by adding Na₂CO₃ (100 µL, 200 mM stock) solution and the absorbance was measured at 400 nm. Final DMSO concentration remained below 7.5% during the assay. Acarbose (1 – 0.031 mM final concentration) was used as reference compound.

The % inhibition was calculated using the following formula:

$$\% \text{ Inhibition} = 100 - [\text{OD}_{(\text{test sample})} / \text{OD}_{(\text{control})}] \times 100$$

Alpha-glucosidase inhibition was categorised as mild (1-25%), moderate (26-50%), high (50-70%), and very high (70-100%).

2.4.3 DPPH radical scavenging activity.

The ability of the crude extract or fractions to scavenge DPPH radicals was evaluated according to Szabo *et al.* (2007) with some modifications. An aliquot (50 µL) of test sample (prepared in a concentration range (3.84 – 0.12 mg/mL in methanol) or quercetin (reference compound with stock concentration ranging from 1.08 – 0.0312 mg/mL in methanol) was mixed with PBS (450 µL, 10 mM, pH 7.4) and 1.0 mL of a methanolic solution of DPPH (0.1 mM), yielding a series of test samples in a final

concentration range of 128 – 4 µg/mL) and a series of quercetin samples in a final concentration of (1 – 36 µg/mL). After 30 min the absorbance was recorded at 517 nm and % inhibition was calculated using the following formula:

$$\text{DPPH radical scavenging activity (\%)} = [(OD_{\text{control}} - OD_{\text{sample}}) / OD_{\text{control}}] \times 100$$

Based on findings of Dudonne *et al.* (2009), DPPH inhibition was categorized as very low (1-10%), low/week (10%-30%), moderate (30-60%), high (61-69%) and very high (70-100%).

2.4.4. Inhibition of 15-lipoxygenase

The soybean lipoxygenase (LOX) assay is used as an indication of anti-inflammatory activity (Muller, 1994). LOX is a key enzyme in the inflammatory cascade, whose inhibition is correlated to the ability of the inhibitors to reduce Fe³⁺ at the active site to the catalytically inactive Fe²⁺. LOXs contain a “non-heme” iron per molecule in the enzyme active site as high-spin Fe²⁺ in the native state and the high-spin Fe³⁺ in the activated state. Several LOX inhibitors, such as phenolic derivatives, are excellent ligands for Fe³⁺. The enzyme 15-lipoxygenase has consistently been reported as involved in oxidation of low-density lipoproteins (LDL) that is responsible for expansion of atherosclerosis (Steinberg, 1999).

The iridoids are more particularly known for their anti-inflammatory activities (Park *et al.*, 2010; Gousiadou *et al.*, 2013). As the major constituents of *Kickxia* were iridoids, it was therefore considered interesting to test the isolated iridoids for their potential to inhibit 15-lipoxygenase. Also flavonoids are reported to possess lipoxygenase inhibitory (Lyckander and Malterud, 1996; Malterud and Rydland, 2000), it was therefore decided also to investigate the isolated flavonoids in the 15-LOX assay.

15-LOX from soybean was used for peroxidation of linoleic acid, and inhibition was carried out as described previously with slight modifications (Malterud and Rydland 2000). Briefly, 12.5 µL (2 - 0.062 mM) of sample, dissolved in DMSO, was added to 487.5 µL of enzyme solution (200 U/mL). The mixture was incubated at room temperature for

5 min. Absorbance was recorded immediately after the addition of 500 μL of substrate (250 μM linoleic acid in 0.2 M borate buffer, pH 9) every min up to 5 min at 234 nm using a Genesys-10UV spectrophotometer.

% inhibition of enzyme activity was calculated as:

$$\% \text{ inhibition of enzyme activity} = \left(\frac{[\Delta A1/\Delta t] - [\Delta A2/\Delta t]}{[\Delta A1/\Delta t]} \right) \times 100$$

where $\Delta A1/\Delta t$ and $\Delta A2/\Delta t$ are the increase rate in absorbance at 234 nm for sample without test substance and with test substance, respectively.

2.4.5 Inhibition of linoleic acid peroxidation

In general anti-inflammatory activity of compounds is believed to be the result of inhibition on the arachidonic acid (AA) pathway. AA is an unsaturated (C_{20}) fatty acid that is produced from membrane phospholipids. The derivatives of AA are potent mediators of inflammation. It is therefore considered that by inhibition of the biosynthesis of pro-inflammatory molecules, the inflammation can be reduced or ended by using antioxidants and anti-inflammatory drugs (Grantstrom 1984).

During the AAPH (2,2'-azobis(2-methylpropionamidine)-dihydrochloride) assay oxidative changes and the effect of the test compound are evaluated. AAPH (water soluble Azo compound) produces alkyl peroxy free radicals through spontaneous thermal decomposition and induces linoleic acid peroxidation. Oxygen species are known to play a key role in various pathophysiological situations and also directly affect lipid peroxidation in membranes and cellular components.

The inhibition of linoleic acid lipid peroxidation was studied by a previously described method (Liegeois *et al.*, 2000; Rajic *et al.*, 2010). Briefly, 10 μL of a 16 mM linoleic acid dispersion and 10 μL of test compound (8-1 μM final concentration) was added to the UV cuvette containing 0.93 mL of 0.05 M phosphate buffer, pH 7.4. The oxidation reaction was initiated under air by the addition of 50 μL (40 mM) of AAPH solution. The rate of oxidation at 37 $^{\circ}\text{C}$ was monitored by recording the increase of absorption at 234 nm caused by conjugated diene hydroperoxides. The results were compared to the standard inhibitor (Trolox). The antioxidant power (AOP) is defined as the slope of the

curve representing the inhibition time of oxidation (T_{inh}) vs. the concentration of antioxidant and is expressed in min/ μ M.

2.4.6. Antimicrobial assays

The antimicrobial tests were performed in the Laboratory for Microbiology, Parasitology and Hygiene (LMPH) of the University of Antwerp. The primary *in vitro* evaluation of the samples was performed against a broad panel of pathogens to allow proper evaluation of selectivity. This approach involved integrated and robotic logistics to attain a reasonably high throughput. Compound stock solutions were prepared in DMSO (100%) at 20 mM (pure compounds) or 20 mg/mL (extracts or fractions). The compounds were then serially diluted (2 or 4 fold) in DMSO followed by a further (intermediate) dilution in demineralized water to ensure a final in-test DMSO concentration less than 1%. Test plates were identical for all screens and were produced as a single batch.

2.4.6.1 Antileishmanial activity

Leishmania infantum MHOM/MA(BE)/67 was maintained in Golden Hamster and spleen amastigotes were collected for preparing infected inocula. The primary peritoneal mouse macrophages were used as host cells and collected two days after peritoneal stimulation with a 2% potato starch suspension. The assay was performed in 96-microwell plates. Each well contained 10 μ L of test compound, 190 μ L of macrophage / parasite inoculum (3×10^5 cells and 3×10^6 parasite per well) and further incubated for 5 days at 37 °C and 5% CO₂. After five days the parasite burden (mean number of amastigotes/macrophage) was microscopically assessed after Giemsa staining, and expressed as a percentage of the blank controls without test compound. Fungizone was used as reference drug. (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008).

2.4.6.2 Antitrypanosomal activity

Trypanosoma brucei Squib-427 strain (suramin-sensitive) was cultured at 37 °C and 5% CO₂ in Hirumi-9 medium (Hirumi and Hirumi, 1989) supplemented with 10% fetal calf serum (FCS). The assay was performed in 96-microwell plates. Each well contains 10 microliter of test compound, 190 μ L of trypomastogotes of an inoculation of 1.5×10^4

trypomastigotes/well . After an incubation of 72 h at 37°C under 5% CO₂. The parasite growth was assessed by adding 50 microliter/well resazurin (Raz *et al.*, 1997). After 24h incubation at 37°C the fluorescence was measured (λ exi 550 nm, λ emi 590 nm). The results were expressed as % reduction in parasite growth compared to the blank controls. Suramine was used as reference drug.

For the Chagas disease, *Trypanosoma cruzi* Tulahuen CL2, β -galactosidase strain (nifurtimox- sensitive) was maintained on MRC-5 (human lung fibroblast) cells in minimum essential medium (MEM), supplemented with 200 mM L glutamine and 5% FCSi. All cultures and assays were conducted at 37 °C under 5% CO₂. The assays were performed using sterilized 96 micro well plates. Each well contained 10 of test substance, 190 μ L MRC-5 cell/parasites inoculum (2×10^4 /mL cell and 2×10^5 parasite/mL). The parasite growth was compared with untreated infected control (100% growth) and non-infected controls (0% growth) after seven days of incubation. The parasite burden was assessed by adding substrate chlorophenol red- β -D-galactopyranoside (CPRG): 50 μ L/well of stock solution containing 15.2 mg of CPRG and 250 μ L Nonidet in 100 ml phosphate buffered saline (PBS) and the change in colour was measured spectrophotometrically at 540 nm after 4h of incubation at 37 °C. The results were expressed as % reduction in parasite burden compared to the control wells. The compounds were tested at five concentrations (ranging from 64 - 0.25 μ M). Benznidazole was used as reference drug (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008).

2.4.6.3 Antiplasmodial activity

The chloroquine-resistant *Plasmodium falciparum* strain K1 was used. Parasites were cultured in human erythrocytes at 37 °C under a low oxygen atmosphere (3% O₂, 4% CO₂ and 93% N₂) in a modular incubation chamber. The culture medium was RPMI-1640 supplemented with 0.5 m/v % albumax (Cos *et al.*, 2006). The assay was performed in 96 micro well plates. Each well containing 10 μ L of test compound, 190 μ L of malaria parasite inoculum (1% parasitaemia, 2% haematocrit) incubated for 72 h at 37°C in modular chamber under low oxygen atmosphere (3% O₂, 4% CO₂ and 93% N₂). After incubation, test plates were frozen at -20 °C. Parasite multiplication was measured by the Malstat method. This method is based on the ability of the lactate dehydrogenase

(LDH) enzyme of *P. falciparum* and the coenzyme 3-acetyl pyridine (APAD) to transform the lactate into pyruvate. One hundred microliters of Malstat reagent was transferred in a new plate and mixed with 20 μ L of the haemolysed parasite suspension for 15 min at room temperature. After addition of 20 μ L nitro blue tetrazolium (NBT)/phenazine ethosulphate (PES) solution (2 mg/mL of NBT and 0.1 mg /mL of PES) and 2h incubation in the dark, the optical density was spectrophotometrically read at 655 nm. Percentage of growth inhibition was compared to the negative blanks. Chloroquine was used as reference drug (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008).

2.4.6.4 Antibacterial activity

The bacterial strains *Staphylococcus aureus* representing gram-positive bacteria and *Escherichia coli* representing gram-negative bacteria were cultured in MHB (Mueller Hinton Broth) at 37 °C. Assays were performed in 96-well microtiter plates, each well containing 10 μ L of test compound together with 190 μ L of inoculum (5×10^3 cfu/mL). After 17 h incubation, bacterial viability was assessed fluorimetrically after addition of resazurin, as described above. The reference drugs used during analysis were erythromycin and trimethoprim (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008)

2.4.6.5 Antifungal activity

The antifungal activity of extracts was tested against *Candida albicans*, *Aspergillus fumigatus* and *Microsporum canis*. The fungal strains were cultured in RPMI-1640 medium supplemented with Mops buffer and glucose at 37 °C. Assays were performed in 96-well microtiter plates, each well containing 10 μ L of extract and 190 μ L of fungal inoculum (5×10^3 cfu/mL). After 24 h incubation, fungal viability was assessed fluorimetrically after addition of 10 μ L resazurin per well (λ exi 550 nm, λ emi 590 nm). The results are expressed as % reduction in fungal growth/viability compared to control wells and an IC₅₀ is determined. Reference drugs include miconazole and terbinafine (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008)

2.4.6.6 Cytotoxicity

MRC-5SV2 (human fetal lung fibroblasts) cells were cultivated in MEM supplemented with L-glutamine (20 mM), 16.5 mM sodium hydrogen carbonate, and 5% FCS. For the assay, MRC-5 cells (3×10^4 cells / well) were seeded onto the test plates containing 10

μL the pre diluted test substance and incubated at 37 °C and 5% CO₂ for 72 h. Cell viability was assessed fluorimetrically 4 h after addition of 50 microliter/well resazurin. Fluorescence was measured (λ exi 550 nm, λ emi 590 nm) and the results are expressed as % reduction in cell viability compared to the blank controls. Tamoxifen was used as reference drug (Cos *et al.*, 2006; Balde' *et al.*, 2010; Mesia *et al.*, 2008)

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CHAPTER 3

**PHYTOCHEMICAL AND BIOLOGICAL INVESTIGATIONS ON
NYMPHOIDES INDICA LEAF EXTRACTS**

Accepted for publication in Phytotherapy Research:

Phytochemical and pharmacological investigations on *Nymphoides indica* leaf extracts
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Sandra Apers and Luc Pieters.

3.1 INTRODUCTION

Wetlands deliver several ecosystem services that afford a variety of human activities (Bostian *et al.*, 2014) such as agriculture, food and water resources (Maya *et al.*, 2003; Grossmann 2012). Furthermore the origin of the majority of human advancements bear their roots around wetlands (Swapna *et al.*, 2011). Together with socioeconomic assistance wetlands also provide an imperative reservoir of medicinal plants worldwide (Cook, 1996; Marwat *et al.*, 2011; Panda and Mishra, 2011) and numerous reports from literature can be consulted that provide clear evidence about synthesis of many secondary metabolites through their natural pathways (Jain *et al.*, 2007, Das *et al.*, 2006). Some promising bioactivities offered by aquatic plants include antibacterial (Abu Ziada *et al.*, 2008; Fareed *et al.*, 2008; Haroon *et al.*, 2009; Sridevi *et al.*, 2010), antifungal (Bhosale *et al.*, 1999; Haroon, 2006) and antiviral (Verma *et al.*, 2008; Shin *et al.*, 2010) activities.

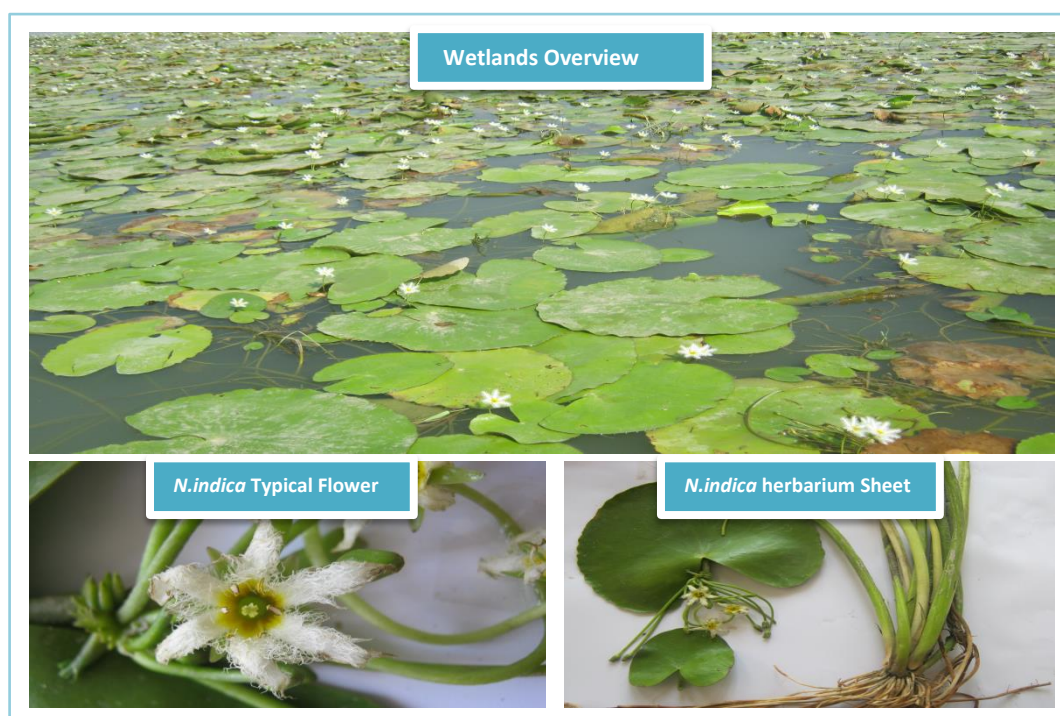


Figure 3.1 *Nymphoides indica* collection from natural habitat and herbarium sheet for identification.

Nymphoides indica (L.) Kuntze (Menyanthaceae) is a perennial rhizomatous free floating leaved aquatic herb (Schmidt, 2005). The genus *Nymphoides* is represented worldwide by about 50 species that are not only very diverse but also differ considerably in a number of biological features (Tippary *et al.*, 2009). *Nymphoides indica* is native to tropical America (Iverson, 2010), however it is reported from various parts of world including Thailand (Ito and Barfod, 2014), India (Madhavan *et al.*, 2009), Australia (Ohwi, 1984), Japan (Shibayama and Kadono, 2007). In Pakistan only three species occur, i.e *Nymphoides cristata* (Roxb.), *Nymphoides peltatum* (S.G. Gmel.) (Qaiser, 1977) and *Nymphoides indica* (Marwat *et al.*, 2009). These species are primarily reported in the districts of Attock, Thatta, Kashmir and D. I. Khan.

N. indica is consumed as a vegetable (Cruz-Garcia and Price, 2011; Swapna *et al.*, 2011) and has a consistent folklore usage in the Indian subcontinent for the management of numerous health indications by local healers (called hakeem or herbal doctors). A number of ethnomedicinal surveys can be accessed that principally emphasise on traditional use of this plant species for treatment of dysentery (Das *et al.*, 2006; Kensa, 2011), scabies (Panda and Misra, 2011), snake bites (Coea and Anderson 2005), jaundice (Jain *et al.*, 2007), anticonvulsant (Madhavan *et al.*, 2009), aphrodisiac (Yumnam and Tripathi, 2012), and as antiproliferative (Kitdamrongtham *et al.*, 2013).

Moreover it has been reported that it is a part of some important herbal formulations in India (Madhavan *et al.*, 2009), Nepal (Siwakoti, 2006), Thailand (Manosroi *et al.*, 2012) and Pakistan (Marwat *et al.*, 2011) that are used for various therapeutic reasons locally. In spite of the multimedical usage of *Nymphoides indica*, no comprehensive published data is available to date concerning its phytochemical constituents and pharmacological properties. Therefore the aim of the present work was to evaluate various medicinal and pharmacological properties of *N. indica* and its constituents, as a part of running programmes on antimicrobial, antioxidant and antidiabetic (inhibition of α -glucosidase activity; inhibition of AGEs (Advanced Glycation Endproducts) properties of medicinal plants and natural products (Upadhyay *et al.*, 2014_a, Upadhyay *et al.*, 2014_b; Mesia *et al.*, 2008).

3.2 EXTRACTION AND ISOLATION

3.2.1 Plant material

N. indica leaves were collected in June and July 2012 from wetlands of the river Indus in Dera Ismail Khan (KPK) Pakistan. The collected plant leaves were identified by the Islamabad Herbarium at the Taxonomy Department, Quaid I Azam University, Islamabad, Pakistan, where the voucher specimen was deposited (5298-GA, accession No. ISL-120019). The leaves were shade dried, powdered and sieved through a 20 mesh filter, and stored below 20°C till further use (Fig 3.2).



Figure 3.2 *Nymphoides indica* dried leaf powder.

3.2.2. Extraction and fractionation

Powdered leaves (1.92 kg) were extracted with 90% (v/v) methanol by double cold maceration. The extract was instantaneously filtered through Whatman No.1 filter paper using a vacuum pump. The filtrates were combined together and evaporated on a rotary evaporator under reduced pressure below 40°C. The resultant semisolid material was lyophilised with a final yield of 88.61 g, and stored below 20°C. Further liquid-liquid partitioning of the crude extract was performed according to a standard extraction scheme (Chapter 2). After partitioning with different solvents as shown in the scheme, *n*-hexane (1.8 g), methanol 90% (11.2 g), chloroform (2.03 g), ethyl acetate

(1.2 g) and *n*-butanol (19.58 g) fractions were obtained. The collected fractions were dried under reduced pressure at 40°C and stored below 20°C.

3.2.3. Phytochemical analysis

Phytochemical screening for various compound classes was carried out on every fraction using published methods (Sofowora, 1993, Trease and Evans, 1989, Harborne, 1973; Egwaikhide and Gimba, 2007; Farnsworth, 1966) with slight modifications for sterols, triterpenes, carbohydrates and saponins flavonoids, alkaloids, and tannins (chapter 2). Phytochemical screening of the different extracts and fractions revealed that flavonoids, tannins, saponins and triterpenes are present with varying concentrations in the different fractions of the leaves extract (Table 3.1). No alkaloids were detected. Fractions of medium polarity contained flavonoids and terpenes. The residual aqueous fraction was positive for tannins and saponins. These findings are consistent with previous reports (Madhavan *et al.*, 2009; Kitdamrongtham *et al.*, 2013).

Table 3.1 Phytochemical analysis of the *N.indica* leaves

Extract fraction	Phytochemicals				
	Alkaloids	Flavonoids	Tannins	Steroids/ triterpenoids	Saponins
MeOH 90%	-	+++	-	+++	-
Chloroform	-	++	-	++	-
Ethyl acetate	-	+++	-	+	-
n-Hexane	-	-	-	-	-
n-butanol	-	+	-	++	++
Aqueous	-	-	+++	-	+++

profoundly present: high degree of precipitation (dark coloration), moderately present: less degree of precipitation (medium colouration), slightly present: very low precipitation (very little colouration), absent: no change in colour (Senguttuvan *et al.*, 2014)

Phytochemicals: +++ profoundly present, ++ moderately present, + slightly present, - absent.

3.2.4. Thin layer chromatography

Each fraction was subjected to TLC (NP) using various solvent systems as mobile phase, including CHCl₃ / MeOH (70:30) with a few drops of NH₄OH for the methanolic fraction; CHCl₃ / MeOH (80:20) for the chloroform and ethyl acetate fractions; *n*-hexane / CHCl₃ (95:5) for the *n*-hexane fraction; and CHCl₃ / MeOH (25:75 or 37:63) the for *n*-butanol fraction. Developed TLC plates were examined under UV at 254 nm and 366 nm, after spraying with *p*-anisaldehyde reagent. For instance, the TLC profiling of the methanolic fraction (FR4) obtained after flash chromatography is shown below. In the highlighted area, four prominent spots are visible (Fig 3.3).

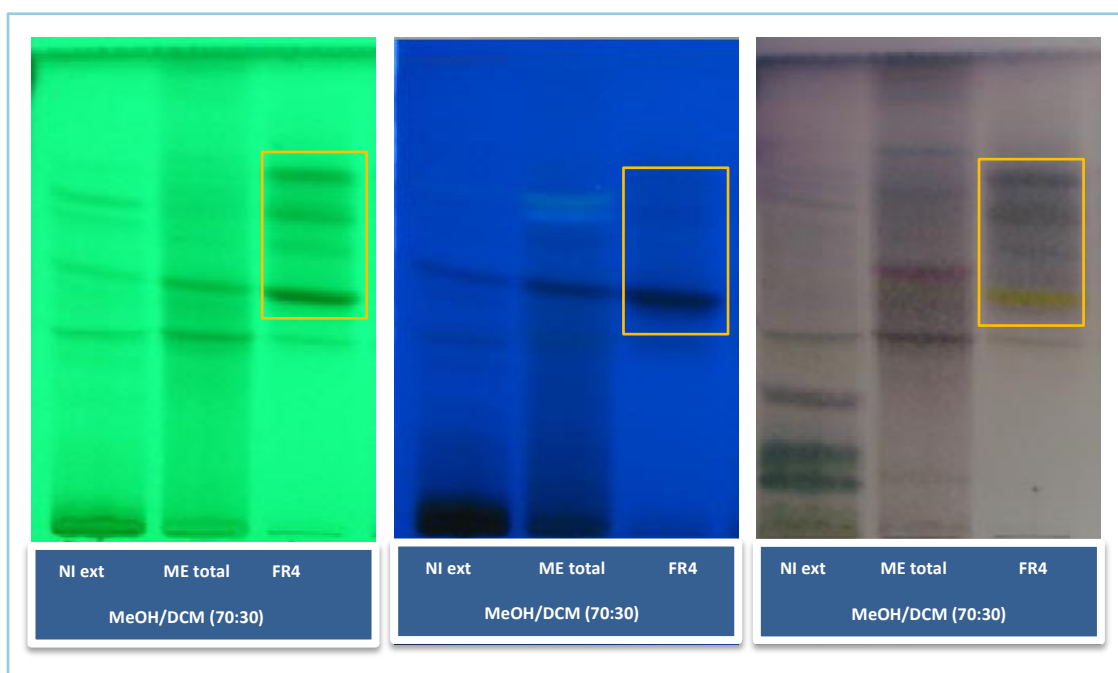


Figure 3.3 TLC profile of total plant extract (NI ext), methanolic extract (ME total) and fraction-4 (FR4) on silica TLC (NP) plate at 254 and 366 nm and after spraying anisaldehyde reagent.

3.2.5. Flash chromatography

An aliquot of 0.8 g from the *n*-hexane fraction was dissolved in 2 mL methanol, mixed with 1.1g silica and dried with nitrogen gas. The dried extract was subjected to flash chromatography with a gradient from *n*-hexane over methylene chloride to methanol. Based on UV and ELSD detection different subfractions were collected (Fig. 3.4). All fractions were analysed by TLC and similar fractions were combined. In this way **21** subfractions were obtained. Based on TLC profiling, subfractions NI4 (190 mg) and NI7 (150 mg) were selected for further HPLC profiling.

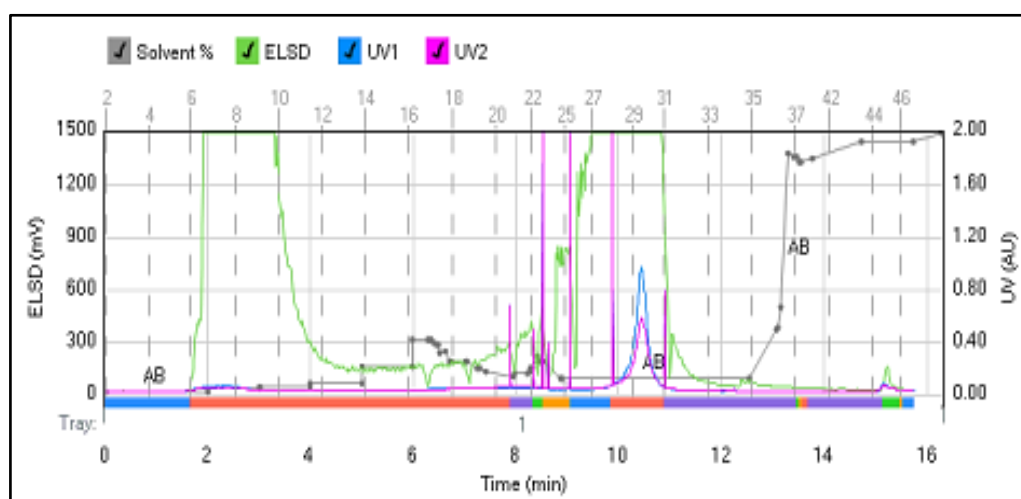


Figure 3.4 Flash chromatogram of *n*-hexane fraction.

An aliquot of 0.8 g from the methanolic fraction was subjected to flash chromatography with a gradient from ethyl acetate to methylene chloride over methanol. Based on UV and ELSD detection different subfractions were collected (Fig 3.5). All fractions were analysed by TLC and similar fractions were combined. A total of **19** subfractions were finally obtained. Finally subfractions NI4 (250 mg), NI6 (180 mg) and NI7 (150 mg) were selected for further HPLC profiling.

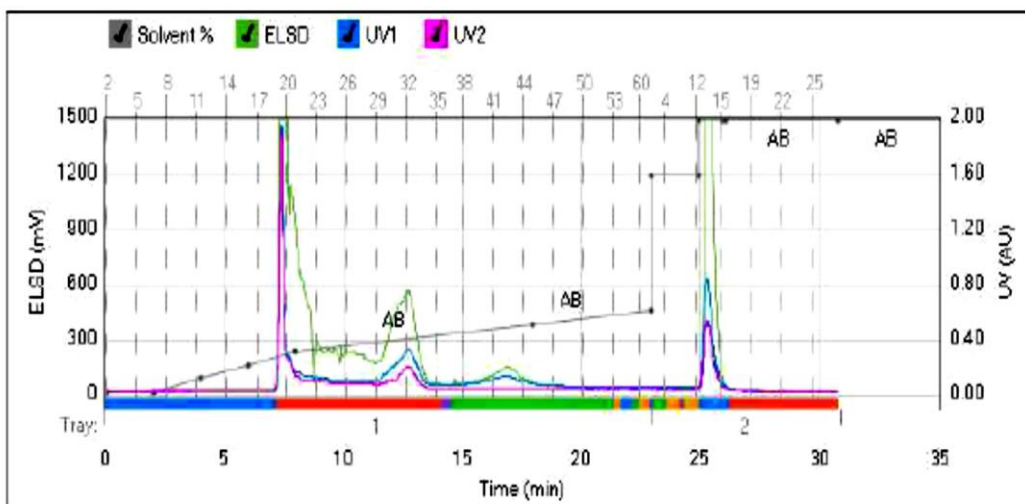


Figure 3.5 Flash chromatogram of methanolic fraction.

Similarly an aliquot of 0.9 g from the ethyl acetate fraction was subjected to flash chromatography with a gradient from ethyl acetate over methylene chloride to methanol. Based on UV and ELSD detection different subfractions were collected (Fig 3.6). All fractions were analysed by TLC and similar fractions were combined as described above; in this way **15** subfractions were obtained. Based on TLC profiling subfractions NI4 (126 mg) and NI7 (154 mg) were selected further HPLC analysis.

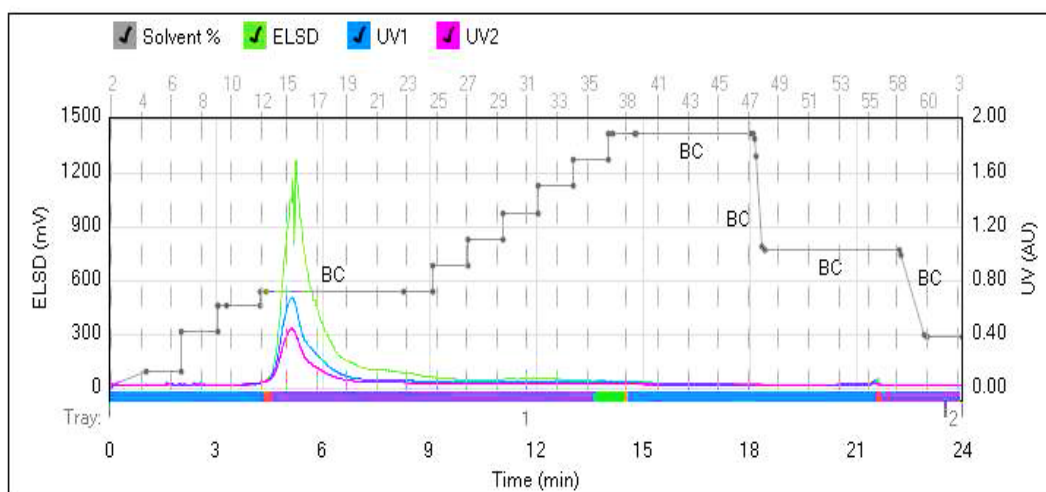


Figure 3.6 Flash chromatogram of ethyl acetate fraction.

Similarly an aliquot of 0.7 g from the chloroform fraction was subjected to flash chromatography with a gradient from ethyl acetate over methylene chloride to methanol as previously explained. Finally **10** subfractions were obtained (Fig 3.7). Based on TLC analysis subfraction NI2 (130 mg) was selected for HPLC analysis.

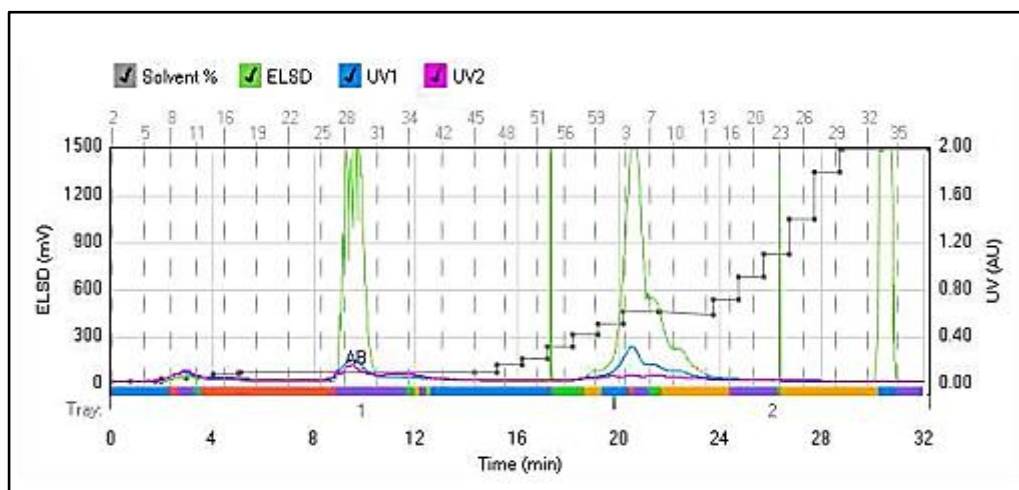


Figure 3.7 Flash chromatogram of chloroform fraction.

Finally an aliquot of 0.8 g from the *n*-butanol fraction was subjected to flash chromatography with a gradient from ethyl acetate over methylene chloride to methanol. Based on UV and ELSD detection different subfractions were collected (Fig 3.8). All fractions were analysed by TLC and similar fractions were combined. Finally **18** subfractions were obtained. Based on TLC analysis subfractions NI2 (126 mg) and NI5 (154 mg) were selected for further HPLC analysis.

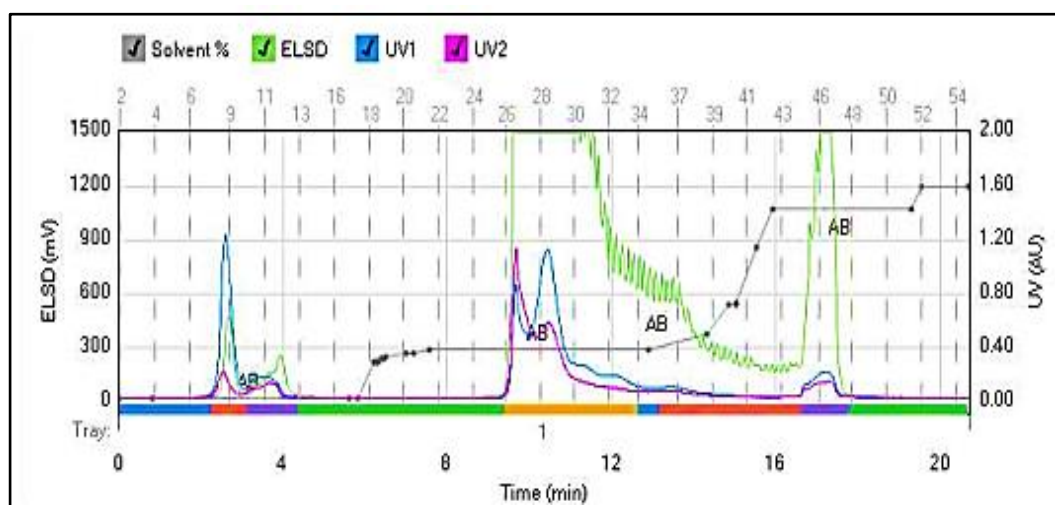


Figure 3.8 Flash chromatogram of *n*-butanol fraction.

3.2.6. HPLC analysis and isolation of compounds

The *n*-hexane fraction and all obtained subfractions NI4 and NI7 were analysed by HPLC using an optimized acetonitrile / H₂O + 0.1% formic acid gradient, ranging from 5% acetonitrile to 100% in 40 min at a flow rate of 1 mL/min. Samples were prepared in a concentration range from 1-10 mg/ mL in methanol (Fig 3.9). The isolation of pure compounds was performed by means of semi-preparative HPLC-DAD-MS using the same gradient at 3 mL/min, yielding compounds **1** (3.5 mg), **2** (4.2 mg), **3** (4.2 mg), **4** (3.6 mg) and **5** (5.2 mg).

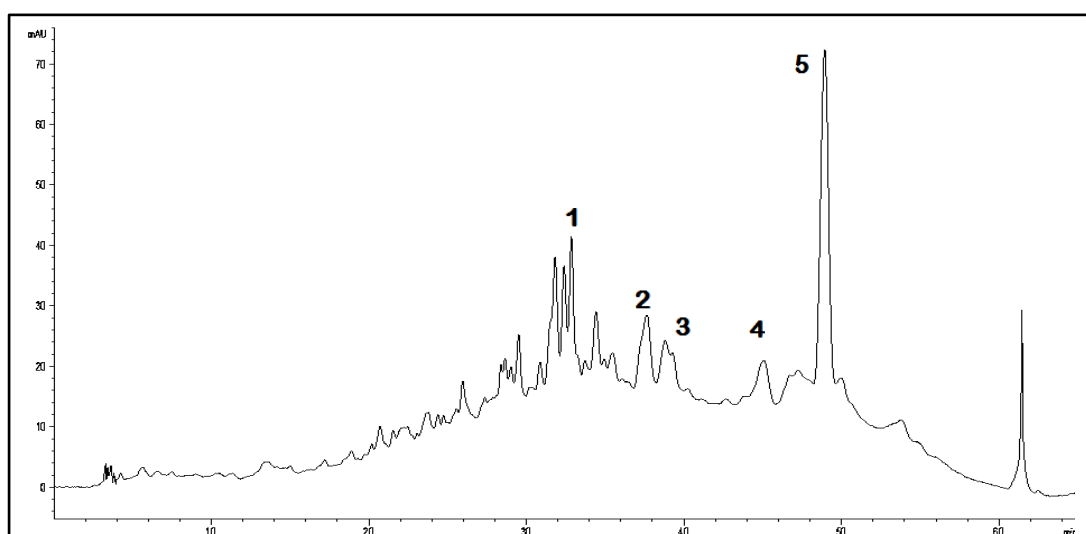


Figure 3.9 HPLC chromatogram of *n*-hexane fraction at 254 nm

The crude methanolic fraction and all subfractions NI4 (250 mg), NI6 (180 mg) and NI7 (150 mg) were analysed by HPLC using an optimized acetonitrile / H₂O + 0.1% formic acid gradient ranging from 15% acetonitrile to 100% in 40 min at a flow rate of 1 mL/min Fig. 3.10). The isolation of pure compounds was performed by means of semi-preparative HPLC-DAD-MS using the same gradient at 3 mL/min Compounds **6** (5.2 mg), **7** (5.7 mg), **8** (7.5 mg), **9** (4.2 mg) and **12** (3.3 mg) were isolated.

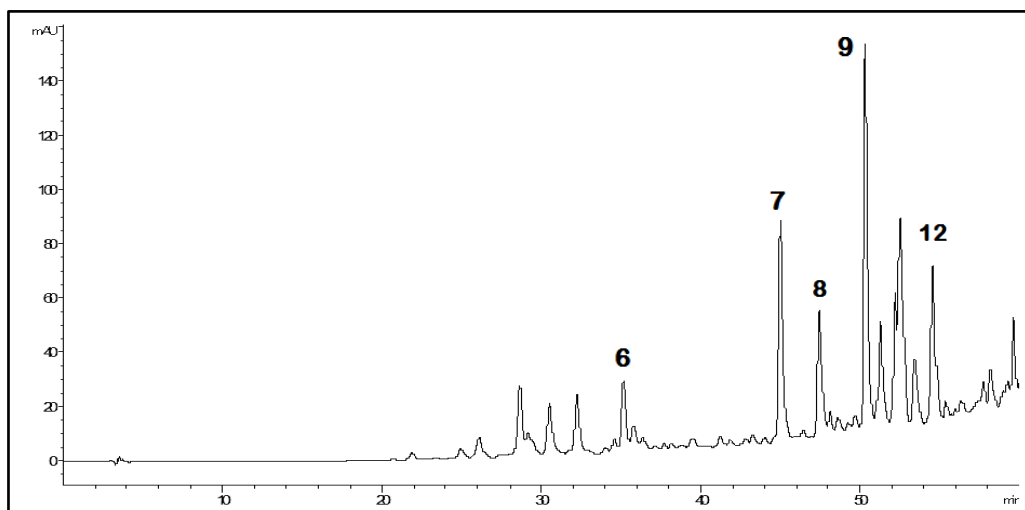


Figure 3.10 HPLC chromatogram of 90% methanol fraction at 254 nm

Similarly the crude ethyl acetate fraction and subfractions NI4 (126 mg) and NI7 (154 mg) were analysed by HPLC using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 35% acetonitrile to 70% in 45 min at a flow rate of 1 mL/min (Fig. 3.11). The compounds were isolated by means of semi-preparative HPLC-DAD-MS system using same gradient at a flow rate of 3 mL/min. Finally the compounds **9** (4.5 mg), **10** (3.2 mg) and **13** (2.0 mg) were isolated.

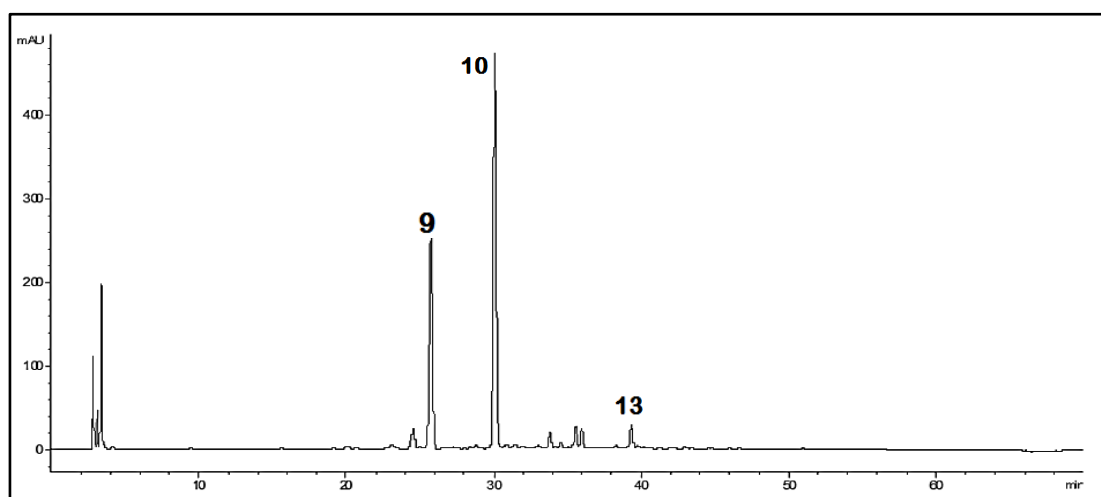


Figure 3.11 HPLC chromatogram of ethyl acetate fraction at 254 nm

The crude chloroform fraction and subfraction NI2 was analysed by HPLC using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 39% acetonitrile to 65% in 45 min at a flow rate of 1 mL/min (Fig. 3.12). Compound isolation was performed by means of semi-preparative HPLC-DAD-MS system using the same solvent gradient as HPLC at a flow rate of 3 mL/min. Finally compounds **10** (3.2 mg) and **11** (3.8 mg) were isolated.

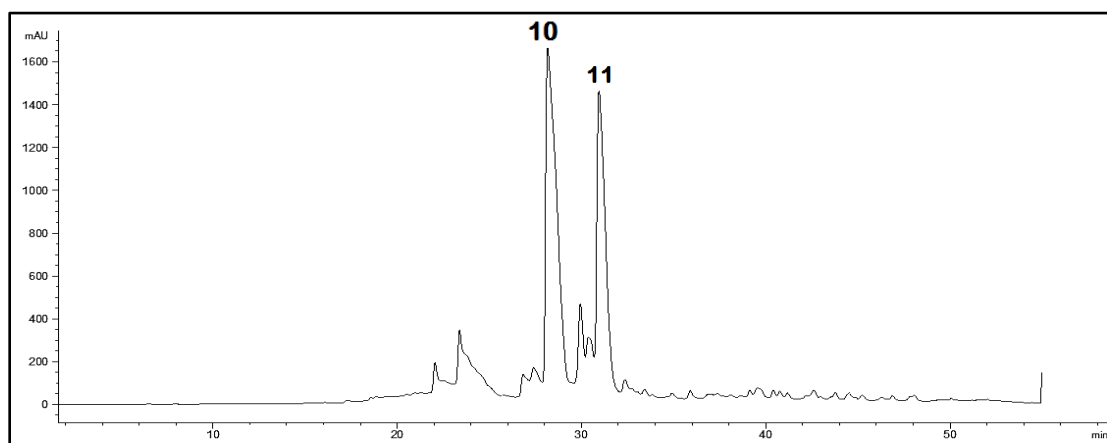


Figure 3.12 HPLC Chromatogram of ethyl acetate fraction at 254 nm

Finally the HPLC profiling of *n*-butanol fraction and subfractions NI2 (126 mg) and NI5 (154 mg) was performed using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 30% acetonitrile to 65% in 50 min at a flow rate of 1 mL/min (Fig 3.13). The compounds **9** (7.3 mg), **10** (4.6 mg), **13** (2.3 mg) and **14**(2.8 mg), were finally isolated by means of semi-preparative HPLC-DAD-MS. The solvent gradient was same as HPLC at a flow rate of 3 mL/min.

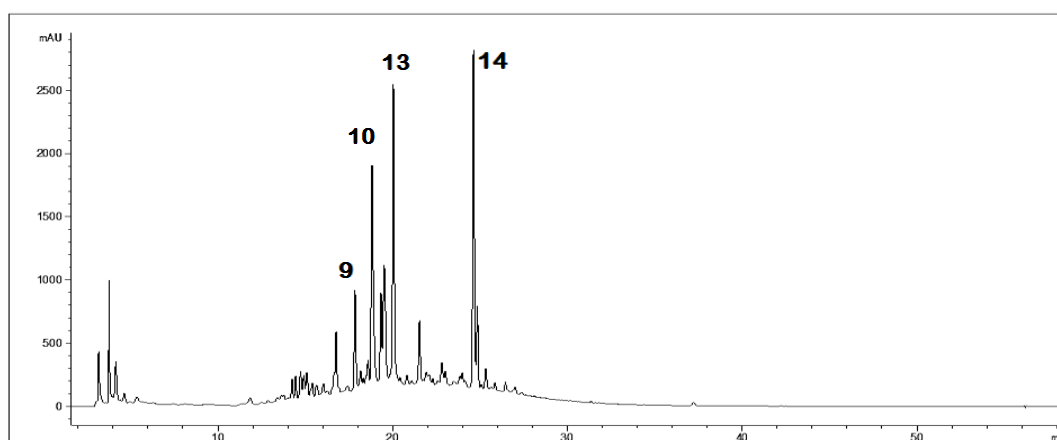


Figure 3.13 HPLC chromatogram of *n*-butanol fraction at 254 nm

3.3. STRUCTURE ELUCIDATION

3.3.1 Lipophilic compounds (1-5)

From the *n*-hexane fraction 5 lipophilic constituents were obtained. Their ¹H- and ¹³C-NMR spectra showed the characteristic features of fatty acids and fatty alcohols, and after derivatisation and GC-MS analysis they were identified as the dicarboxylic acids azelaic acid (nonanedioic acid) (1) and 4-methyl-heptanedioic acid (3), the monocarboxylic acids hexadecanoic acid (2) and stearic acid (5) and the fatty alcohol hexadecanol (4) (Table 3.2)

Table 3.2 GC-MS profile of compounds isolated from *n*-hexane fraction.

Comp	RT(min)	MW	Prob.	Formula	Peak name
1	27.73	332	45.24	C ₁₅ H ₃₂ O ₄ Si ₂	Azelaic acid-bis(trimethylsilyl)ester
2	35.57	328	72.86	C ₁₉ H ₄₀ O ₂ Si	Hexadecanoic acid trimethylsilyl ester
3	24.37	318	48.36	C ₁₄ H ₃₀ O ₄ Si ₂	Heptanedioic acid, 4-methyl (bistrimethylsilylester)
4	39.10	314	17.38	C ₁₉ H ₄₂ O ₂ Si	1-trimethylsiloxyhexadecane
5	39.36	356	48.99	C ₂₁ H ₄₄ O ₂ Si	Stearic acid trimethylsilylester

3.3.2 Seco iridoids (6-8)

3.3.2.1 7-Epiexaltoside (6):

The structure of compound **6** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) as **7-epiexaltoside** (Fig. 3.14).

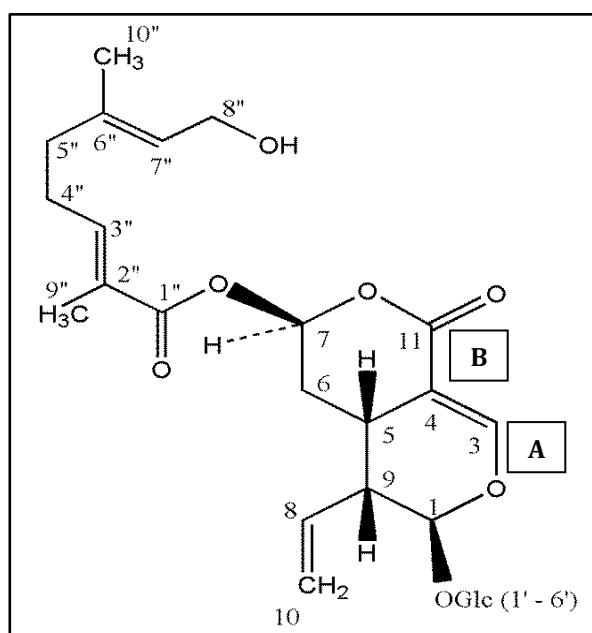


Figure 3.14 structure of 7-epiexaltoside (**6**)

Library search on the $^{13}\text{C-NMR}$ data (Table 3.3 and Fig. 3.15) suggested this compound was a seco-iridoid with a glucosyl moiety at C-1. Based on its $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ and mass spectra, and compared with data from literature (Junior, 1991) compound **6** was found to be identical to 7-epiexaltoside. The $^1\text{H-NMR}$ (Table 3.3 and Fig 3.16) and $^{13}\text{C-NMR}$ (Table 3.3 and Fig. 3.15) are in agreement with a geranyl-like side chain, with two double bonds having an *E*-configuration. The carbonyl group of the ester moiety showed a $^{13}\text{C-NMR}$ resonance at 167.1 ppm, which is typical for an α,β unsaturated ester.

The geranyl-like side chain has 2 tertiary carbons C-2'' and C-6'' showing signals at 128.3 ppm and 137.3 ppm, respectively. These tertiary carbons were substituted by a methyl group at C-9'' and C-10'' that showed signals at 12.4 ppm and 16.2 ppm, respectively.

Two secondary carbons in position C-4"and C-5" showed signals at 28.1 ppm and 39.1 ppm, respectively. The signal at 59.4 ppm was assigned to a primary carbon at position 8".

The compound has a typical bicyclic ring system consisting of 2 rings A and B. In ring A C-1, showing a signal at 99.1 ppm, was attached to a β -D-glucopyranoside moiety. Carbon C-3 showed a CH-signal at 155.4 ppm that is characteristic to an iridoid. C-4 showed signal at 104.4 ppm. The signals at 23.1 ppm and 29.4 ppm were assigned to C-5 and C-6, respectively.

In ring B, a secondary carbon at C-7 showed a signal at 93.6 ppm attached to the geranyl ring through -O- bond with. The ring has a characteristic carbonyl group (C-11) of the ester moiety that resonates at 167.1 ppm. Finally, an unsaturated side chain is attached to ring A at C-9. The signals at 133.3 ppm and 121.5 ppm were assigned to C-8 and C-10, respectively.

The $^1\text{H-NMR}$ spectrum showed the characteristic H-3 iridoid signal at 7.67 ppm, as a doublet with a coupling constant of 2.5 Hz. The singlet at 5.55 ppm was assigned to H-1. Two multiplets at 3.35-3.48 ppm and 2.77 ppm were assigned to H-5 and H-9, respectively. The signal for H-8 (5.54 ppm) was attributed to the X part of an ABX system of a vinylidene group, coupling with the signals at 5.29-5.38 ppm (H-10a and H-10b), as the AB part. The methylene protons (H-6) of the ring B were assigned to signals at 1.86-1.98 ppm. The methylene protons showed a coupling with H-7. The absorption of H-7 occurred at 6.63 ppm with a typical small coupling constant ($J=1.9$ Hz) between H-7 and H-6a as well as H-6b, which is consistent with an axial arrangement (or β -orientation) of the ester at C-7, as demonstrated before for menthiafolin and dihydrofoliamenthin (Junior, 1989; 1991). The geranyl side chain consisted of 3 methylene protons, i.e. H-4", H-5" and H-8". The triplets at 2.38 ppm and 2.18 ppm were assigned to H-4"and H-5", whereas a doublet at 4.09 ppm was assigned to H-8".

With regard to the methine protons in the geranyl side chain,, the doublet at 7.67 ppm was attributed to H-3", whereas the triplet at 7.67 ppm was assigned to H-7". Finally, the spectrum was completed by the signals for a glucose moiety. Based on structural features compound 7-epiexaltoside can be classified as a secoiridoid ester glucoside.

Yellow powder; UV (acetonitrile / H₂O) λ_{max} 200 and 233 nm; ESI-MS (positive ion mode): m/z 563 [M+Na]⁺ consistent with molecular formula C₂₆H₃₆O₁₂.

7-Epiexaltoside was originally obtained from *Villarsia exaltata* (Menyanthaceae) by Junior (1991), and interestingly *N. indica* is also a member of this small family of aquatic and wetland plants comprising five genera and fifty species worldwide.

Table 3.3 ¹H-NMR and ¹³C-NMR assignments for 7-epiexaltoside (6) recorded in methanol-*d*₄

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1	5.55; s	99.1
3	7.67; d; <i>J</i> =2.5	155.4
4		104.5
5	3.35-3.48; m	23.1
6	1.86-1.98; m	29.4
7	6.63; t; <i>J</i> =1.9	93.6
8	5.54-5.60; m	133.3
9	2.77; m	43.5
10	5.29-5.38; m	121.5
11		167.1
1'	4.72; d; <i>J</i> =7.7	101.0
2'	3.20-3.89, m	74.8
3'	3.20-3.89, m	78.3
4'	3.20-3.89, m	71.5
5'	3.20-3.89, m	78.4
6'	3.20-3.89, m	62.6
1''		166.1
2''		128.3
3''	6.86; m	145.7
4''	2.38; m	28.1
5''	2.18; t; <i>J</i> =7.4	39.0
6''		137.3
7''	5.40; t partly overlapped by H-10	126.0
8''	4.09; d; <i>J</i> =6.7	59.4
9''	1.85; s	12.4
10''	1.70; s	16.2

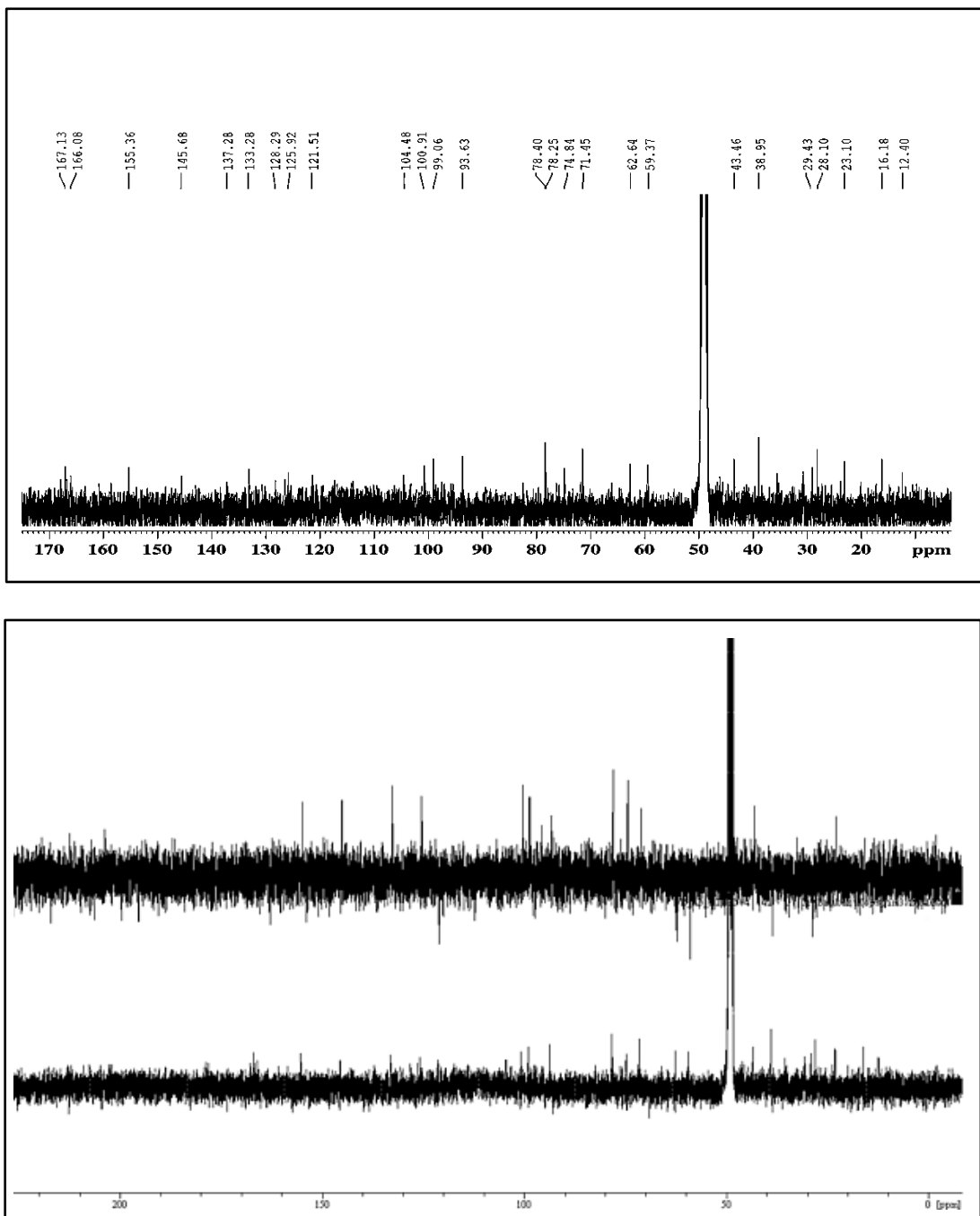


Figure 3.15 ^{13}C -NMR spectra including DEPT-135 of 7-epi exaltoside (6)

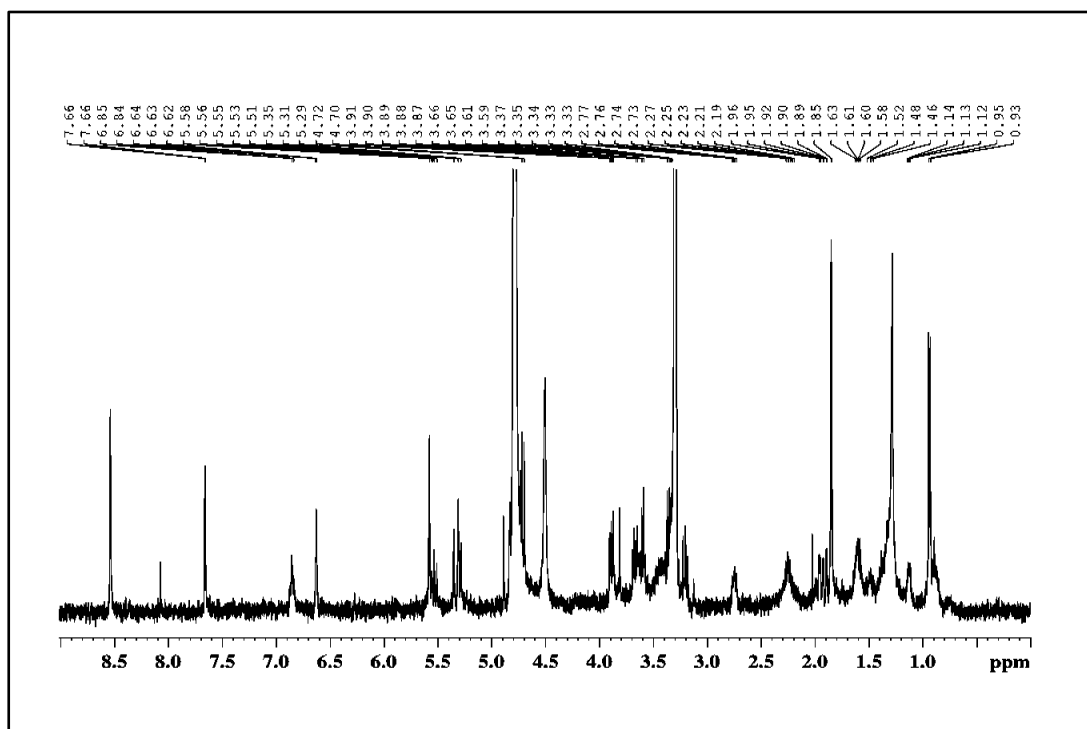


Figure 3.16 ¹H NMR spectrum of 7-epi exaltoside (6)

3.3.2.2 6'', 7''-Dihydro-7-epiexaltoside (7)

The structure of compound **7** was elucidated by ¹H-NMR and ¹³C-NMR (DEPT-135 and DEPT-90) as 6'', 7''-dihydro-7-epiexaltoside (Fig 3.17).

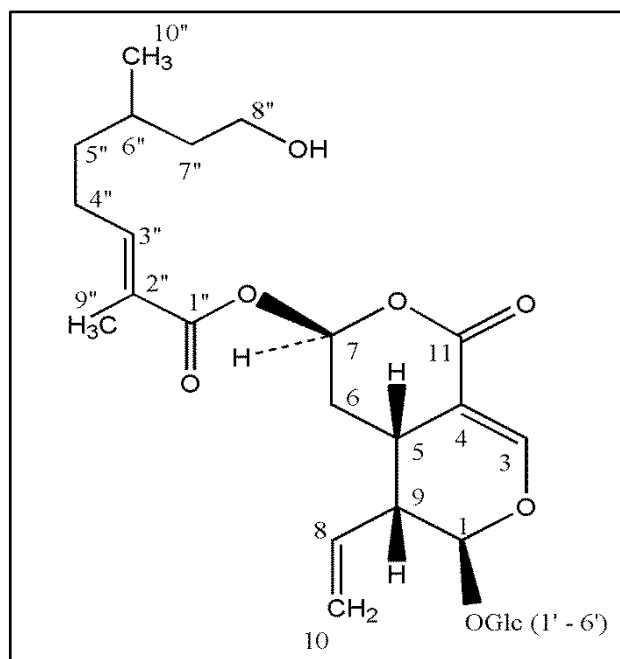


Figure 3.17 Structure of 6'', 7''-dihydro-7-epiexaltoside (**7**)

Library search on the ¹³C-NMR data (Table 3.4 and Fig. 3.18) suggested this compound is a seco-iridoid with a glucosyl moiety similar to compound **6** (Junior, 1991). Based on its ¹H-NMR (Table 3.4 and Fig. 3.19), ¹³C-NMR (Table 3.4 and Fig. 3.20) and mass spectral data compound **7** was found to be 6'', 7''-dihydro-7-epiexaltoside, containing a geranyl-like side chain, with only one double bond having an *E*-configuration. The carbonyl group of the ester moiety showed a ¹³C-NMR resonance at 167.1 ppm, which is typical for an α,β -unsaturated ester as in compound **6**.

The NMR spectra of compound **7** were very similar to those observed for compound **6**, but it contained one double bond less, which was confirmed by the [M+Na]⁺ signal observed in MS at *m/z* 565, rather than *m/z* 563 for compound **6**. Since the signals assigned to the seco-iridoid moiety in compound **6** were also observed in compound **7**, it was concluded that the saturation was located in the geranyl-like side chain. Because

the ester carbonyl was observed at δ 166.1, as expected in case of an α,β -unsaturation, it was concluded that, compared to compound **6**, the 6'',7''- double bond was saturated rather than the 2'',3''-double bond. A compound named dihydrofoliamenthin, containing a geranyl-like side chain saturated in position 2'',3'' and a *Z*-configuration at the 6'',7''- double bond, has been reported before from *Villarsia exaltata* by Junior (1991). Hence, compound **7**, for which the name 6'',7''-dihydro-7-epiexaltoside was adopted, is reported here for the first time.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 200 and 233 nm; ESI-MS (positive ion mode): *m/z* 565 [M+Na]⁺ consistent with a molecular formula C₂₆H₃₈O₁₂.

Table 3.4 ¹H-NMR and ¹³C-NMR assignments for 6'',7''-dihydro-7-epiexaltoside (7) recorded in methanol-*d*₄.

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1	5.57; d; <i>J</i> = 1.7	99.0
3	7.66; d; <i>J</i> = 2.3	155.3
4		104.6
5	3.35-3.45; m	27.4
6	1.88-1.98; m	30.6
7	6.62; t; <i>J</i> = 2.0	93.6
8	5.50; m	133.1
9	2.74; m	43.5
10	5.29-5.38; m	121.4
11		167.1
1'	4.70; d; <i>J</i> = 7.8	100.7
2'	3.32-3.90; m	74.8
3'	3.32-3.90; m	78.4
4'	3.32-3.90; m	71.5
5'	3.32-3.90; m	78.3
6'	3.32-3.90; m	62.7
1''		166.1
2''		127.9
3''	6.86; m	146.3
4''	2.24; m	27.4
5''	1.45-1.59; m	40.5
6''	1.40; m	23.1
7''	1.49-1.60; m	36.6
8''	3.60; m	61.0
9''	1.84; s	12.3
10''	0.93; d; <i>J</i> = 6.7	19.8

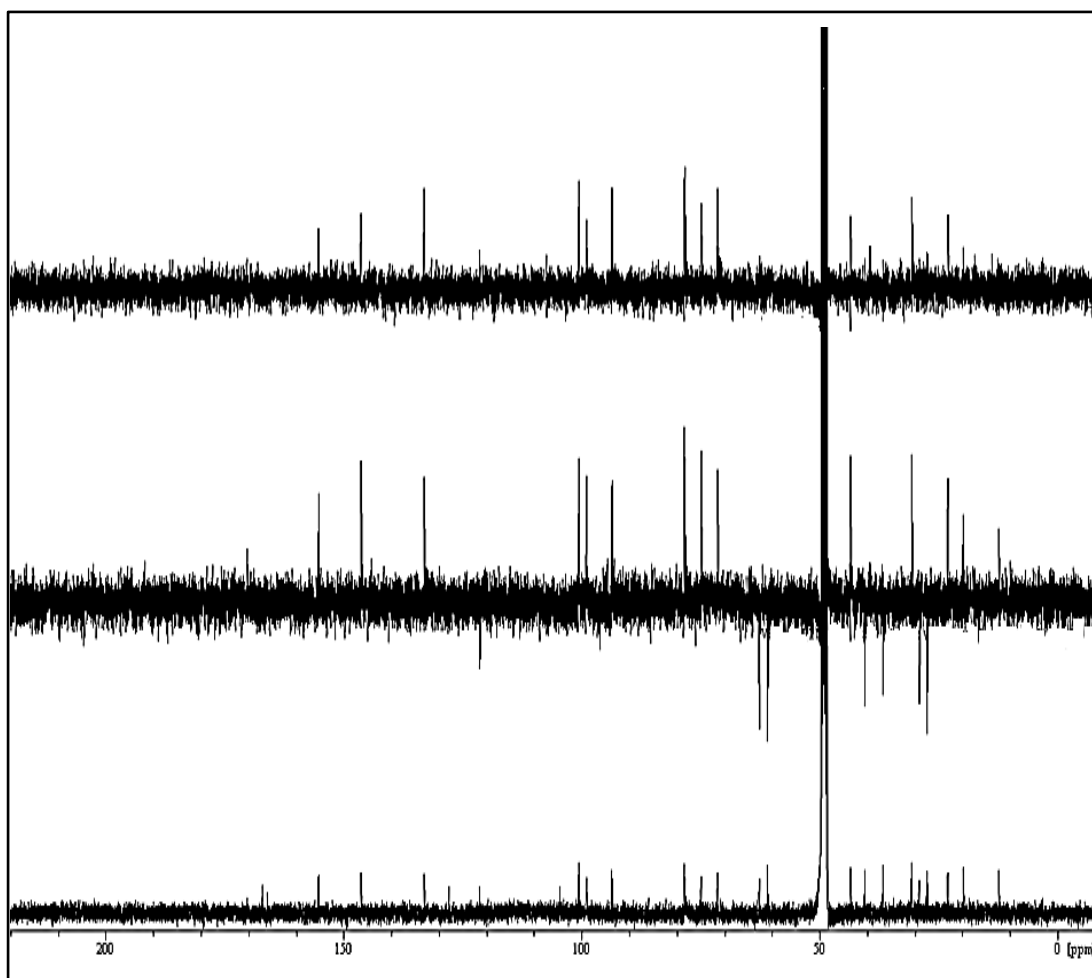
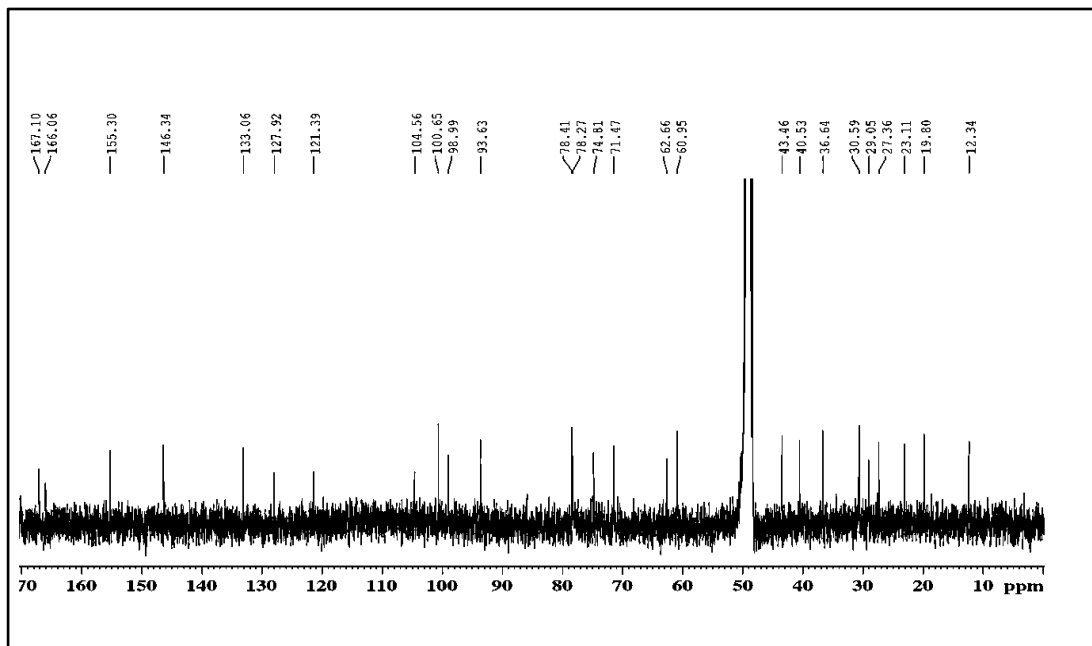


Figure 3.18 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of 6'',7''-dihydro-7-epiexaltoside (7)

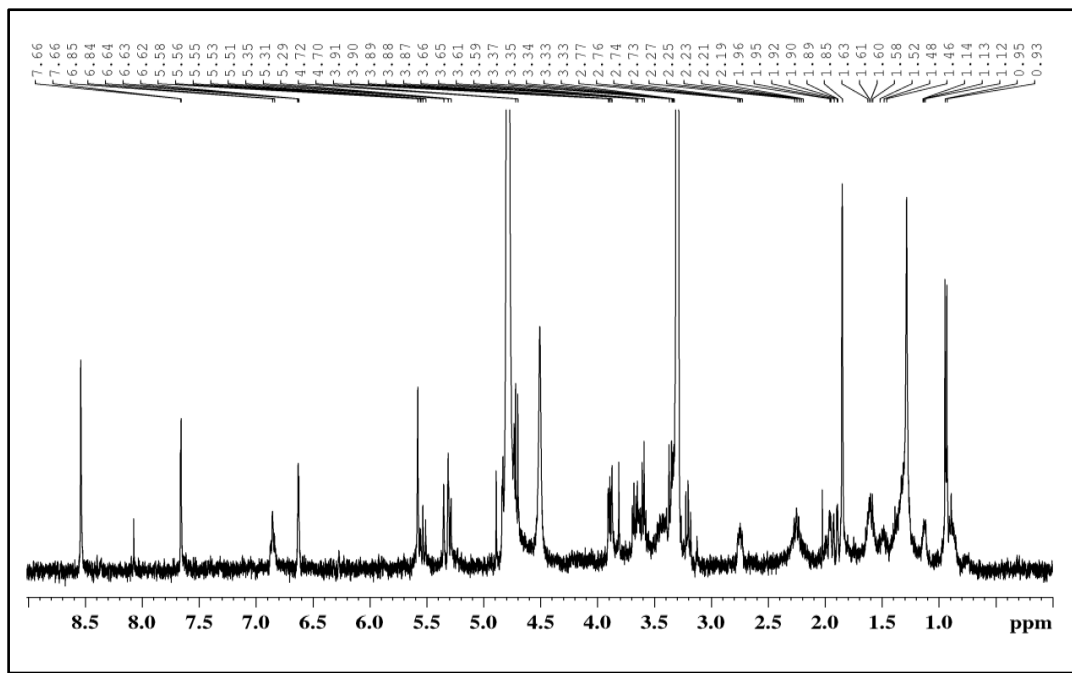


Figure.3.19 ¹H-NMR spectrum of 6'',7''-dihydro-7-epiexaltoside (7)

3.3.2.3 Menthiafolin (8)

The structure of compound **8** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) as **menthiafolin (8)** (Fig. 3.20).

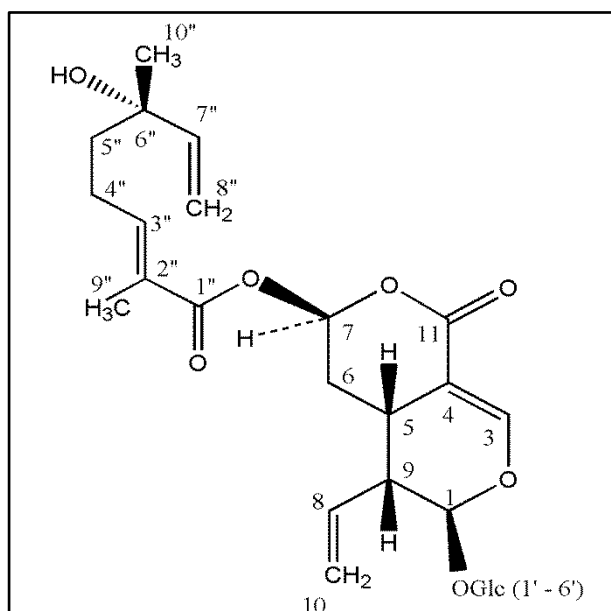


Figure 3.20 Structure of menthiafolin (**8**)

Library search on the $^{13}\text{C-NMR}$ data (Table 3.5 and Fig. 3.21) suggested this compound is a seco-iridoid with a glucosyl moiety. The $^{13}\text{C-NMR}$ spectrum of compound **8** revealed that it contained 26 carbon atoms, 16 of which belonging to the seco-iridoid core being in very good agreement with compounds **6** and **7**. However, a C10 side chain with a terminal double bond could be assigned as reported before by Junior (1989). Based on its $^1\text{H-NMR}$ (Table 3.5 and Fig. 3.22) and $^{13}\text{C-NMR}$ (Table 3.5 and Fig. 3.21) and mass spectral data compound **8** was found to be menthiafolin reported before from *Menyanthes trifoliata* (Junior, 1989).

In fact the acyl substituent present in compound **8** is 6-hydroxy-2,6-dimethyl-2-(E),7-octa-dienoic acid, which was also isolated as a pure compound from the chloroform fraction (see below). The signals at 1.28 ppm (s) and 1.85 ppm (s) can be assigned to the methyl groups C-9" and C-10".

The vinyl protons at C-7 "and C-8" act as a typical ABX spectrum. The signal for H-7" (5.93; dd; $J=17.4$; 10.7 Hz) constitutes the X part of an ABX system, showing a coupling with signals at 5.07 ppm (dd; $J=10.8$; 1.5 Hz) and 5.24 ppm (dd; $J=17.3$; 1.5; Hz) (H- 8"a and H-8"b) as the AB part.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 200 and 233 nm; ESI-MS (positive ion mode): m/z 563 [M+Na]⁺ consistent with a molecular formula C₂₆H₃₆O₁₂.

This is the first report on the occurrence of seco-iridoid in *N. indica*. Menthiafolin was previously isolated from *Menyanthes trifoliata* (Menyanthaceae). *Nymphoides indica* also belongs to the Menyanthaceae family that comprises mainly aquatic and wetland herbs.

Table 3.5 ¹H-NMR and ¹³C-NMR assignments for menthiafolin (**8**) recorded in methanol-*d*₄.

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1	5.61; d; <i>J</i> = 1.8	96.6
3	7.64; d; <i>J</i> = 2.4	153.1
4		103.3
5	3.30-3.45 (1H, m, overlapped by sugar protons H-5)	23.8
6	1.60-1.76; m	28.9
7	6.60; m	93.3
8	5.53; m	131.5
9	2.77; m	42.3
10	5.33-5.38; m, H-10a and H-10b	120.0
11		165.1
1'	4.70; d; <i>J</i> = 7.8	98.3
2'	3.18-3.95; m	73.4
3'	3.18-3.95; m	77.0
4'	3.18-3.95; m	70.2
5'	3.18-3.95; m	76.5
6'	3.18-3.95; m	61.3
1''		165.5
2''		126.3
3''	6.92; dt; <i>J</i> = 7.5;1.4	145.0
4''	2.23; m	23.3
5''	1.65; m	40.1
6''		73.30
7''	5.93; dd; <i>J</i> =17.4; 10.7	144.5
8a	5.07; dd; <i>J</i> =10.7;1.5	111.0
8b	5.24; dd; <i>J</i> =17.4;1.5	
9''	1.85; s	10.8
10''	1.28; s	26.4

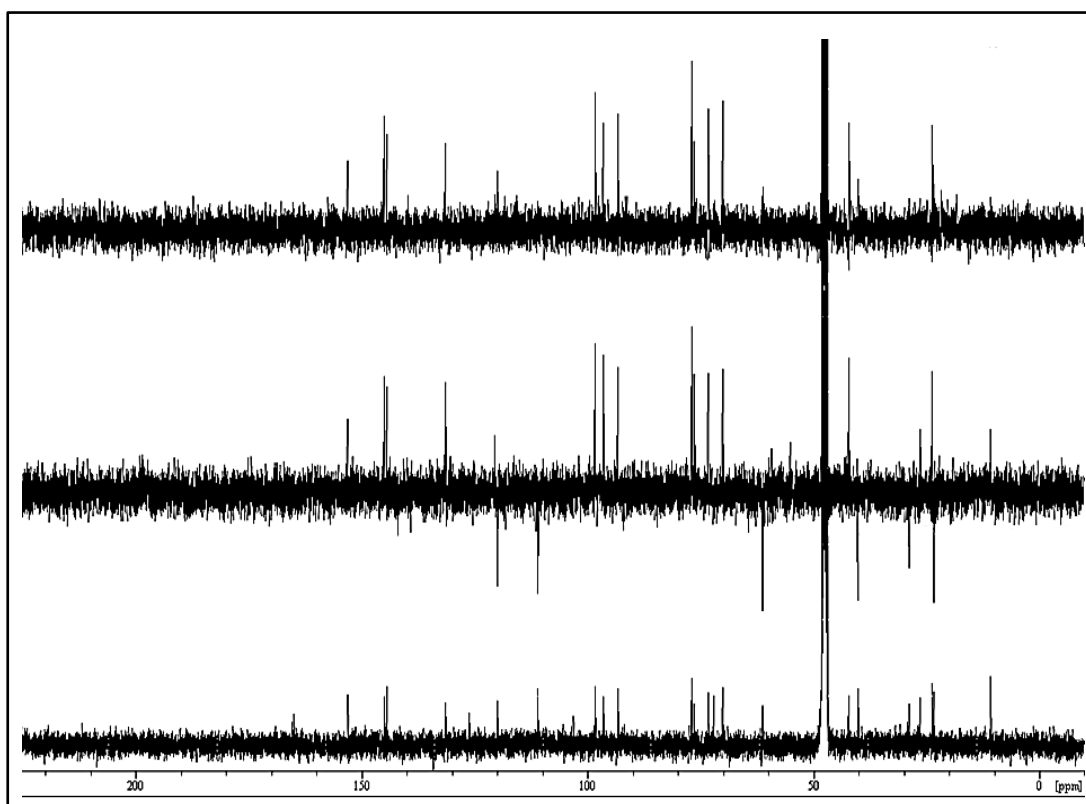
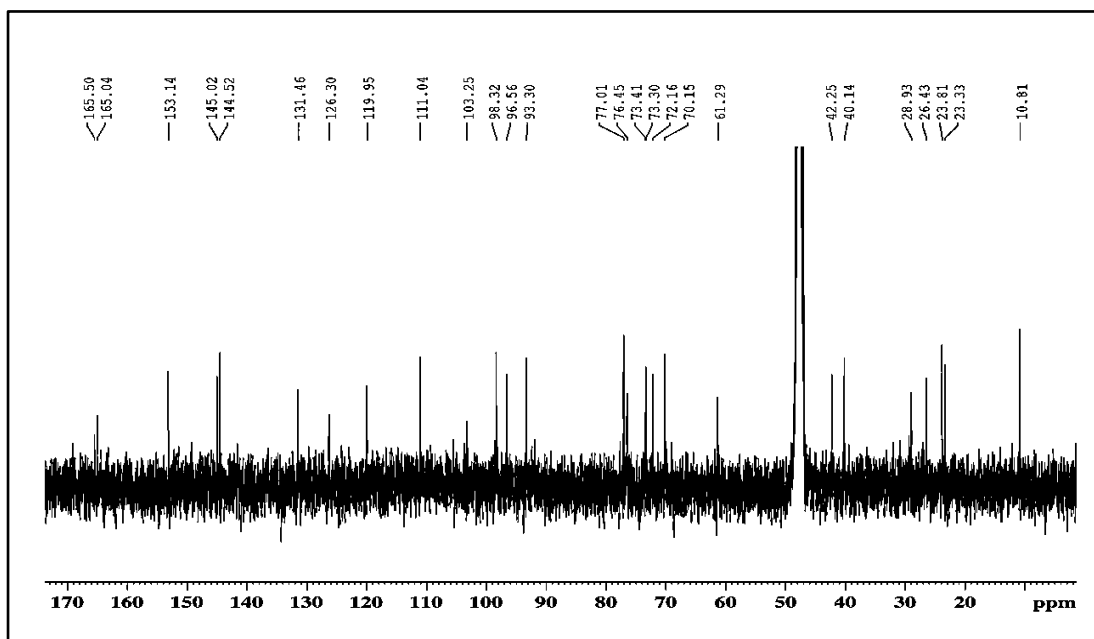


Figure 3.21 ^{13}C -NMR spectra including DEPT 135 and DEPT 90 of menthiafolin (**8**)

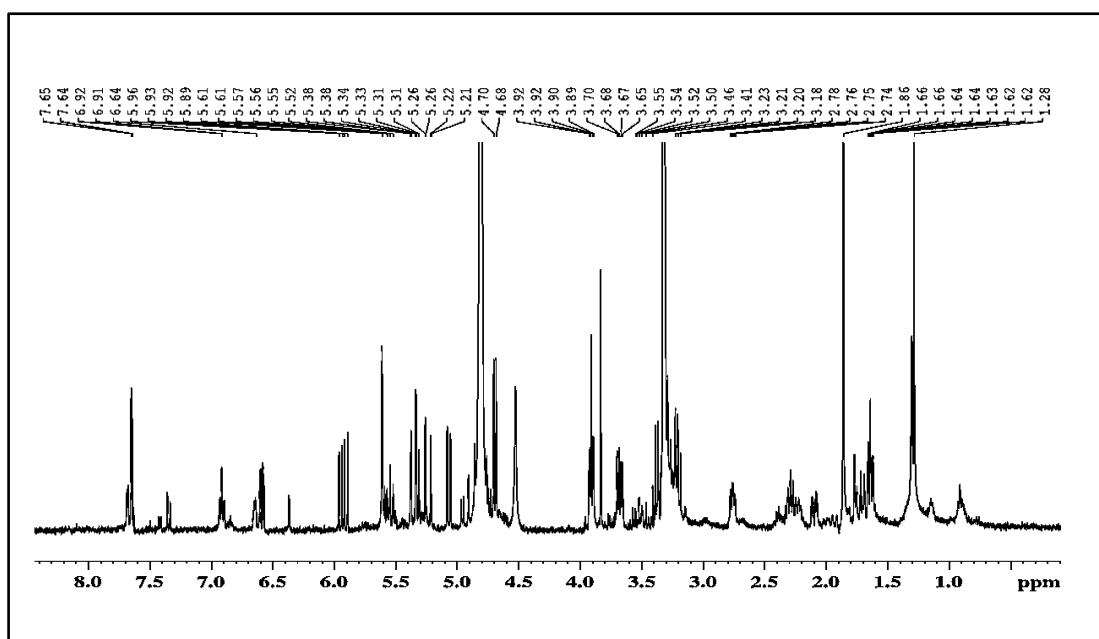


Figure 3.22 ¹H-NMR spectrum of menthiafolin (8)

3.3.3 Flavonoids (9-11)

3.3.3.1 3,7-Di-*O*-methylquercetin-4'-*O*- β -glucoside (9)

The structure of compound **9** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) and 2D-NMR as **3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (9)** (Fig. 3.23).

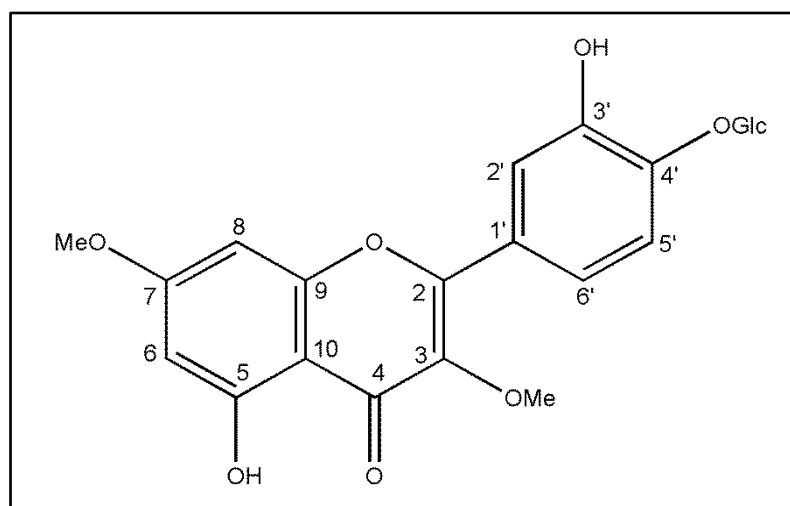


Figure 3.23. Structure of 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (**9**)

Library search on the $^{13}\text{C-NMR}$ data (Table 3.6 and Fig. 3.24) suggested this compound is a flavonoid with glucosyl moiety. Its ^1H , $^{13}\text{C-NMR}$ and mass spectral data were compared with published data (Agarwal and Rastogi, 1981; Guvenalp and Demirezer, 2005), allowing its complete identification.

The $^1\text{H-NMR}$ (Table 3.6 and Fig. 3.25) and $^{13}\text{C-NMR}$ (Table 3.6 and Fig. 3.24) spectra of compound **9** exhibited resonances due to aromatic ring systems, typical for a quercetin moiety and a sugar residue. In the $^1\text{H-NMR}$ spectrum the aromatic region exhibited an ABX system at 7.66 ppm (1H, d, $J = 2.2$ Hz, H-2'), 7.63 ppm (1H, dd, $J = 2.2$ and 8.6 Hz, H-6'), and 7.33 ppm (1H, d, $J = 8.6$ Hz, H-5') due to a 3',4' substitution of ring B, and a typical meta-coupled pattern for H-6 (6.37 ppm, d, $J = 2.2$ Hz) and H-8 (6.56 ppm, d, $J = 2.2$ Hz). An anomeric proton signal appeared at 5.05 ppm (d, $J = 7.3$ Hz, H-1') and the resonances in the region of 3.44 -3.96 ppm (6H, m, H-2'', H-3'', H-4'', H-5'', H-6'') together

with the corresponding carbon resonances inferred from the HSQC (Fig 3.26) spectrum suggested the presence of an *O*-glucopyranosyl unit. The signals at 3.82 ppm (s) and 3.91 ppm (s) are assigned to the methoxyl-groups at C-3 and C-7.

The ¹³C-NMR spectrum of compound **9** showed the presence of 23 aromatic carbon signals. In the HMBC (Fig.3.27) spectrum, a cross-peak between C-4' and H-1" established the linkage point of quercetin and the sugar moiety. The ¹³C-NMR signals at 55.1 and 59.3 ppm were assigned to the methoxyl groups at C-7 and C-3, respectively.

Yellow powder; UV (acetonitrile / H₂O) λ_{max} 200, 251 and 345 nm ESI-MS (positive ion mode): *m/z* 493 [M+H]⁺ consistent with a molecular formula C₂₃H₂₄O₁₂.

3,7-Di-*O*-methylquercetin-4'-*O*-β-glucoside (**9**) was the major flavonoid isolated from leaves of *Nymphoides indica* during the current investigation. Previous records also suggested its occurrence in *Nymphoides indica* leaves (Bhom *et al.*, 1986).

Table 3.6. ^1H -NMR and ^{13}C -NMR assignments for 3,7-di-*O* methylquercetin-4'-*O*- β -glucoside (**9**) recorded in methanol- d_4 .

Position	δ ^1H ppm, multiplicity, <i>J</i> Hz,	δ ^{13}C ppm
1		
2		146.7
3		138.9
4		178.8
5		156.0
6	6.37; d; <i>J</i> = 2.2	97.7
7		166.0
8	6.56; d; <i>J</i> = 2.2	91.8
9		156.9
10		105.4
1'		
2'	7.66; d; <i>J</i> = 2.2	125.9
3'		116.2
4'		147.8
5'	7.33; d; <i>J</i> =8.6	115.8
6'	7.63; dd, <i>J</i> =8.6; 2.0	120.6
1''	5.05; d; <i>J</i> =7.3	101.8
2''	3.44-3.96; m	73.4
3''	3.44-3.96; m	69.9
4''	3.44-3.96; m	76.1
5''	3.44-3.96; m	61.1
6''	3.44-3.96; m	77.1
OMe (C-7)	3.91; s	55.1
OMe (C-3)	3.82; s,	59.3

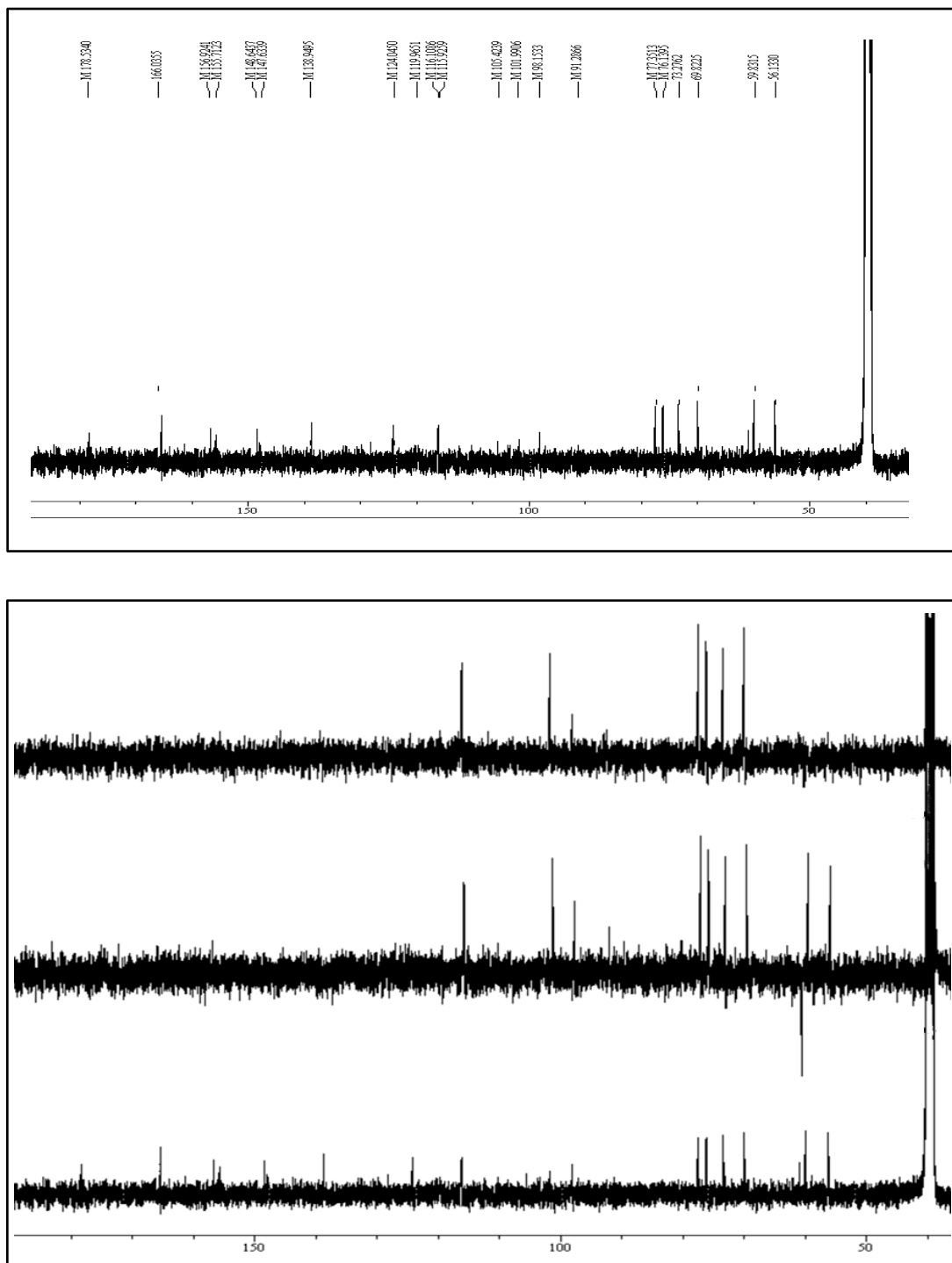


Figure. 3.24 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of 3,7-di-*O*-methylquercetin- 4'-*O*- β -glucoside (**9**)

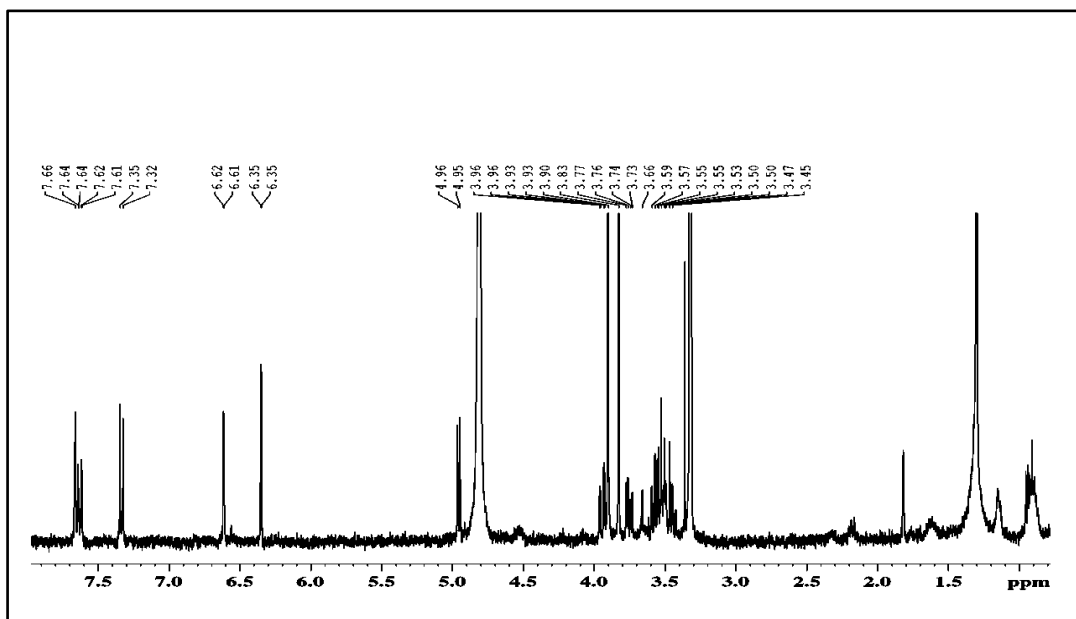


Figure.3.25 ¹H-NMR spectrum of 3,7-di-O-methylquercetin-4'-β-O-glucoside (9)

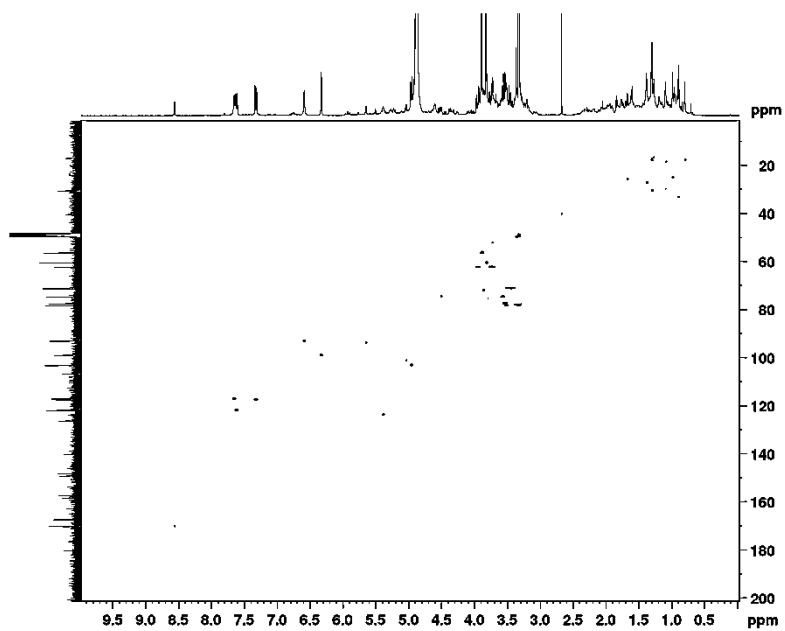


Figure.3.26 HSQC spectrum of 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (9)

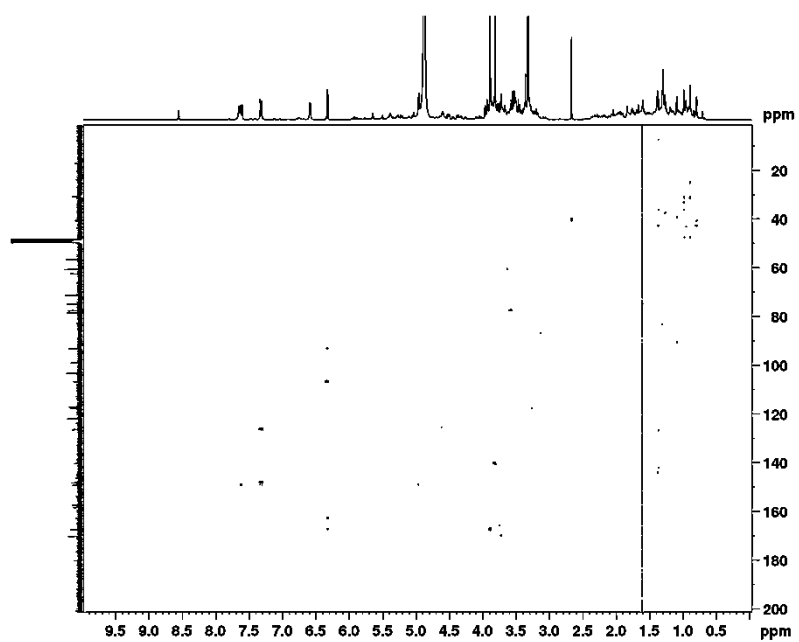


Figure.3.27 HMBC spectrum of 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (**9**)

3.3.3.2 3-O-Methylquercetin-7-O- β -glucoside (**10**)

The structure of compound **10** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) and 2D NMR (HSQC and HMBC) as **3-O-methylquercetin-7-O- β -glucoside (**10**)** (Fig. 3.28).

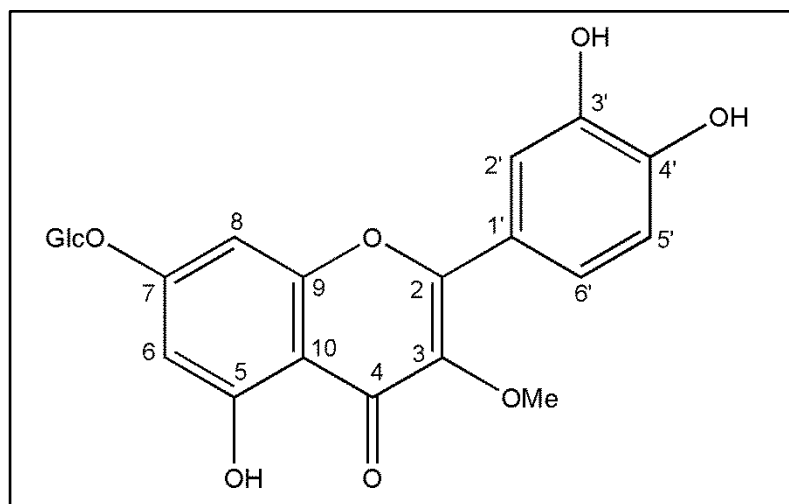


Figure 3.28 Structure of 3-O-methylquercetin-7-O- β -glucoside (**10**)

Library search on the $^{13}\text{C-NMR}$ data (Table 3.7 and Fig. 3.29) suggested this compound is a flavonoid with a glucosyl moiety. By comparison with published data (Krenn *et al.*, 2003), compound **10** was identified as 3-O-methylquercetin-7-O- β -glucoside.

The $^1\text{H-NMR}$ (Table 3.7 and Fig. 3.30) and $^{13}\text{C-NMR}$ (Table 3.7 and Fig. 3.29) spectra of compound **10** exhibited resonances due to aromatic ring systems, typical for a quercetin moiety and a sugar residue as discussed earlier for compound **9**. Similar to the $^1\text{H-NMR}$ spectrum of **9**, the aromatic region exhibited a typical pattern (an ABX system) at 7.66 ppm (1H, d, $J = 2.0$ Hz, H-2'), 7.57 ppm (1H, dd, $J = 2.0$ and 8.4 Hz, H-6'), and 6.92 ppm (1H, d, $J = 8.4$ Hz, H-5') due to a 3', 4' substitution of ring B and a typical meta-coupled pattern for H-6 (6.50 ppm, d, $J = 2.0$ Hz) and H-8 (6.77 ppm, d, $J = 2.0$ Hz). An anomeric proton appeared at 5.07 ppm (d, $J = 7.3$ Hz, H-1') and the resonances in the region of 3.41-3.96 ppm (6H, m, H-2'', H-3'', H-4'', H-5'', H-6'') suggested the presence of an O-

glucopyranosyl unit. The signals at 3.83 ppm (s) was assigned to a methoxyl group at C-3.

The ^{13}C -NMR spectrum of compound **10** showed the presence of 22 aromatic carbon signals. The chemical shift of C-7 (163.4 ppm) indicated substitution with a sugar moiety. The ^{13}C -NMR signal at 59.1 ppm was assigned to the methoxyl group at C-3. Based on its ^1H and ^{13}C -NMR assignments and literature (Krenn *et al.*, 2003) the structure was confirmed as 3-*O*-methylquercetin-7-*O*- β -glucoside. The spectral assignments were confirmed by HSQC (Fig. 3.31) and HMBC (Fig. 3.32).

Yellow amorphous powder; UV (acetonitrile / H_2O) λ_{max} at 200, 251 and 345 nm; ESI-MS (positive ion mode): m/z 479 $[\text{M}+\text{H}]^+$ consistent with a molecular formula $\text{C}_{22}\text{H}_{22}\text{O}_{12}$.

This is a common flavonoid, previously isolated from a variety of medicinal plants, for instance *Achillea nobilis* (Asteraceae) and *Leonotis leonurus* (Lamiaceae).

Table 3.7 ¹H-NMR and ¹³C-NMR assignments for 3-*O*-methylquercetin-7-*O*-β-glucoside (9) recorded in methanol-*d*₄.

Position	δ ¹ H ppm, multiplicity, J Hz,	δ ¹³ C ppm
2		156.6
3		138.4
4		178.8
5		161.4
6	6.50; d; J=2.0	99.4
7		163.4
8	6.77; d; J=2.0	94.4
9		157.2
10		106.3
1'		121.1
2'	7.66; d; J=2.0	115.2
3'		145.1
4'		148.9
5'	6.92; d; J=8.4	115.1
6'	7.57; dd; J=8.4; 2.0	121.3
1''	5.07; d; J=7.3	100.3
2''	3.41-3.96; m	73.3
3''	3.41-3.96; m	76.5
4''	3.41-3.96; m	69.9
5''	3.41-3.96; m	77.0
6''	3.41-3.96; m	61.1
OMe (C-3)	3.83; s	59.1

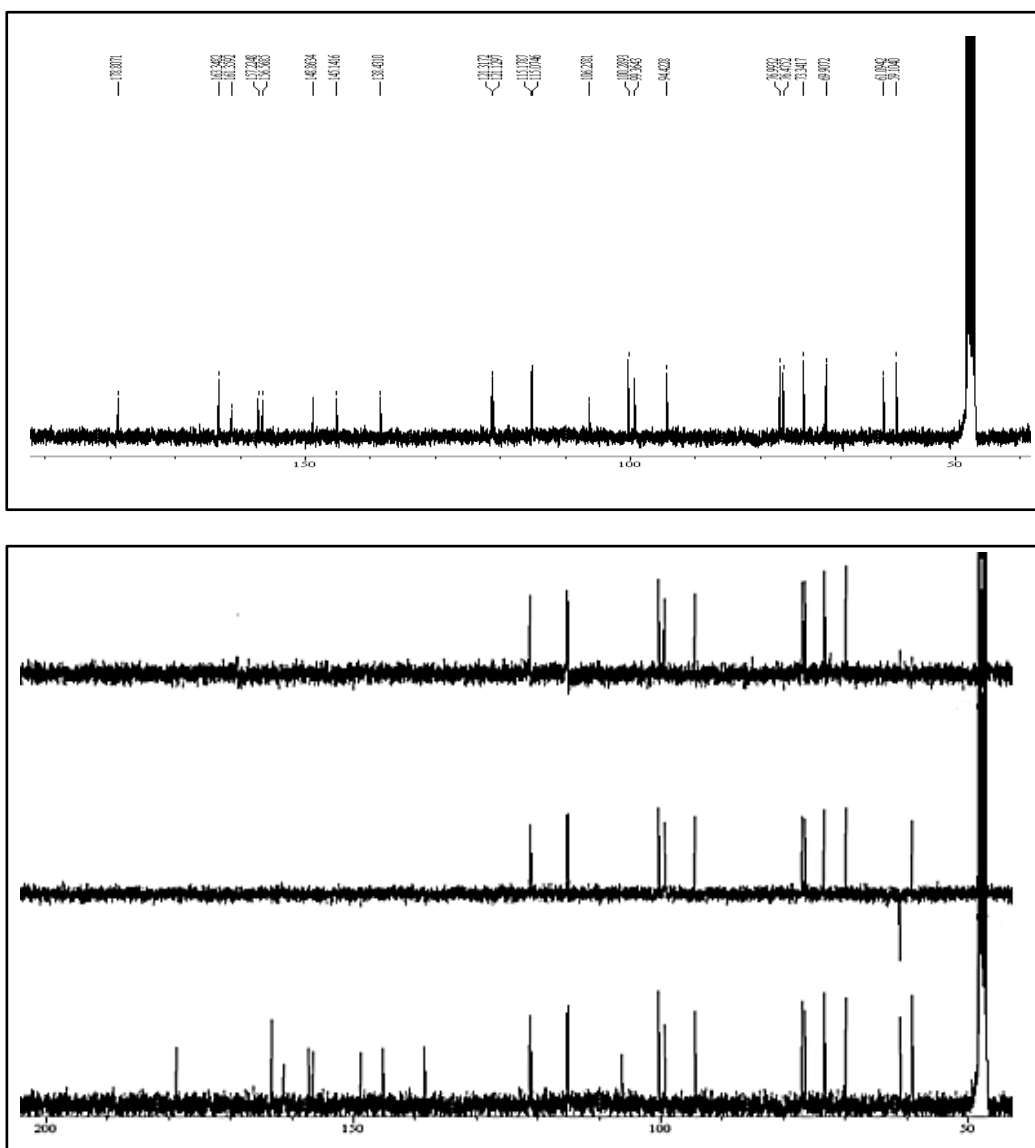


Figure.3.29 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of 3-*O*- β -methylquercetin-7-*O*-glucoside (**10**)

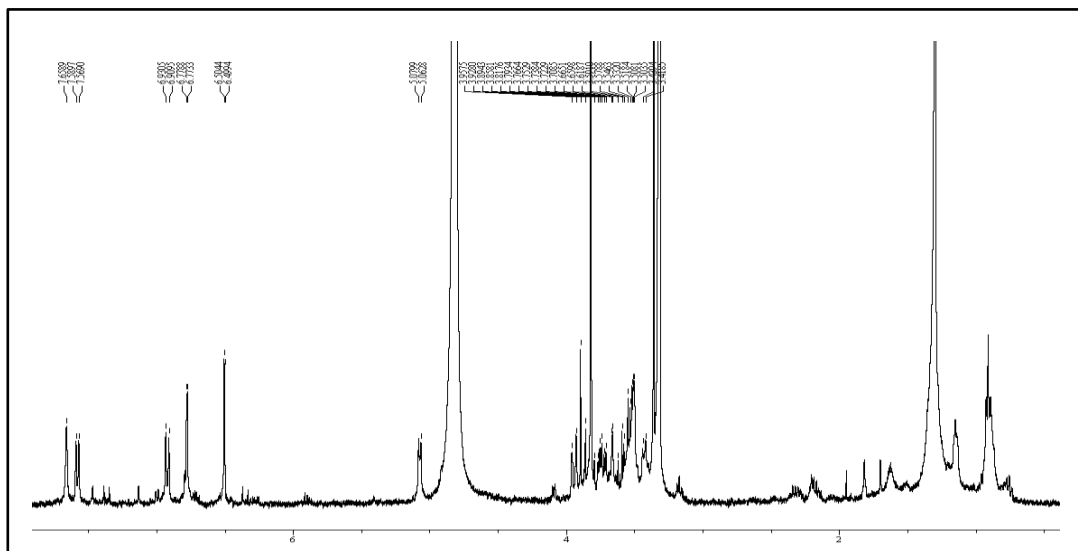


Figure.3.30 ¹H-NMR spectrum of 3,7-di-O-methylquercetin-4'-O-β-glucoside (**10**)

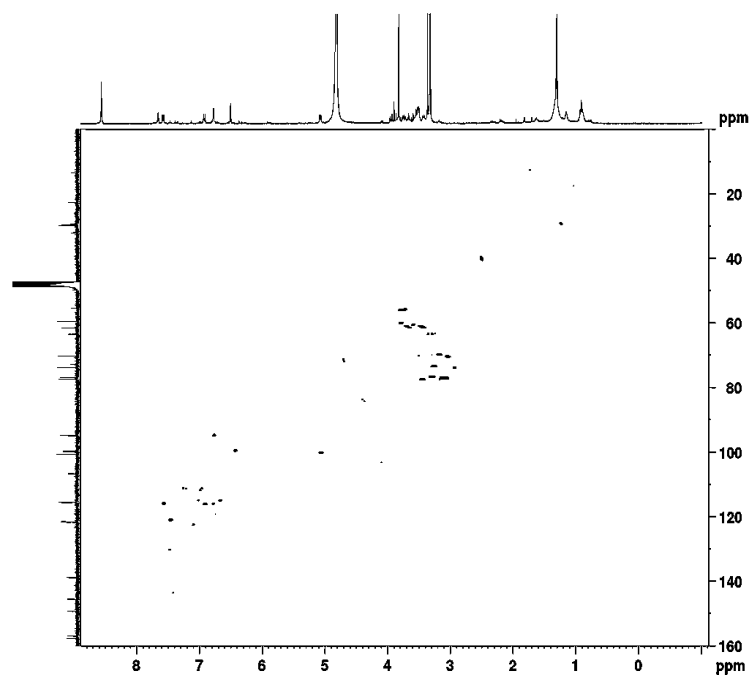


Figure.3.31 HSQC spectrum of 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (**10**)

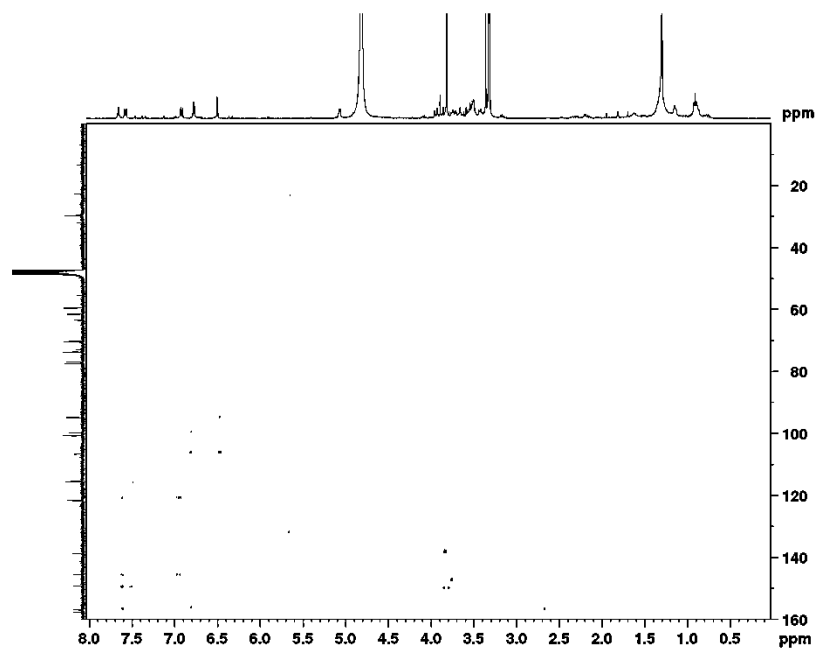


Figure 3.32 HMBC spectrum of 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside (**10**)

3.3.3.3 3,7-di-*O*-methylquercetin (**11**)

The structure of compound **10** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) as 3,7-di-*O*-methylquercetin (**11**) (Fig. 3.33).

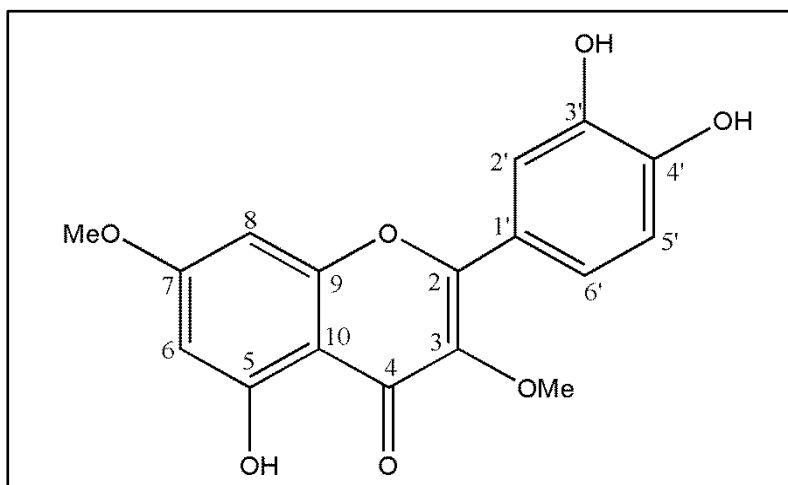


Figure 3.33 Structure of 3,7-di-*O*-methylquercetin (**11**)

Library search on the $^{13}\text{C-NMR}$ data (Table 3.7 and Fig. 3.34) suggested this compound is a flavonoid with a glucosyl moiety. By comparison with published data (Guerrero *et al.*,2002), compound **11** was identified as 3,7-di-*O*-methylquercetin.

The $^1\text{H-NMR}$ (Table 3.7 and Fig. 3.35) and $^{13}\text{C-NMR}$ (Table 3.7 and Fig. 3.34) spectra of compound **11** exhibited resonances due to aromatic ring systems, typical for a quercetin moiety. The $^1\text{H-NMR}$ spectrum exhibited a typical pattern (an ABX system) as discussed earlier (for compound **9** and **10**) at 7.65 ppm (1H, d, $J = 2.0$ Hz, H-2'), 7.56 ppm (1H, dd, $J = 2.0$ and 8.4 Hz, H-6'), and 6.92 ppm (1H, d, $J = 8.4$ Hz, H-5') due to a 3', 4' substitution of ring B and a typical meta-coupled pattern for H-6 (6.30 ppm, d, $J = 2.0$ Hz) and H-8 (6.62 ppm, d, $J = 2.0$ Hz). The signals at 3.83 ppm (s) and 3.90 ppm (s) was assigned to a methoxyl group at C-3 and C-7, respectively.

The ^{13}C -NMR spectrum of compound **11** showed the presence of 17 aromatic carbon signals. The chemical shift of C-4 (163.4 ppm) indicated the presence of a carbonyl moiety. The ^{13}C -NMR signals at 59.1 ppm and 55.1 were assigned to the methoxyl groups at C-3 and C-7, respectively. Based on its ^1H and ^{13}C -NMR assignments and literature (Guerrero *et al.*, 2002) the structure was confirmed as 3,7-di-*O*-methylquercetin. All correlations were confirmed by COSY (Fig. 3.36), HSQC (Fig. 3.37) and HMBC (Fig 3.38).

Yellow amorphous powder; UV (acetonitrile / H_2O) λ_{max} 203, 257 and 360 nm ESI-MS (positive ion mode): m/z 331 $[\text{M}+\text{H}]^+$ consistent with a molecular formula $\text{C}_{17}\text{H}_{14}\text{O}_7$.

This flavonoid was previously isolated from a variety of medicinal plants, for instance *Croton schiedeanus* Schlecht (Euphorbiaceae), *Saccopetalum horsfieldii* Benn (Annonaceae) and *Hypericum ternum* (Hypericaceae).

Table 3.8 ¹H-NMR and ¹³C-NMR assignments for 3,7-di-*O*-methylquercetin (**11**) recorded in methanol-*d*₄.

Position	δ ¹ H ppm, multiplicity, J Hz,	δ ¹³ C ppm
2		157.0
3		138.4
4		178.8
5		161.6
6	6.34; d; <i>J</i> = 2.0	97.5
7		166
8	6.62; d; <i>J</i> = 2.0	91.7
9		159.2
10		105.3
1'		121.4
2'	7.65; d; <i>J</i> =2.0	115.0
3'		145.1
4'		148.9
5'	6.92; d; <i>J</i> = 8.4	115.2
6'	7.56; dd; <i>J</i> = 8.4; 2.0	121.4
OMe (C-3)	3.81; s	55.1
OMe (C-7)	3.90; s	59.1

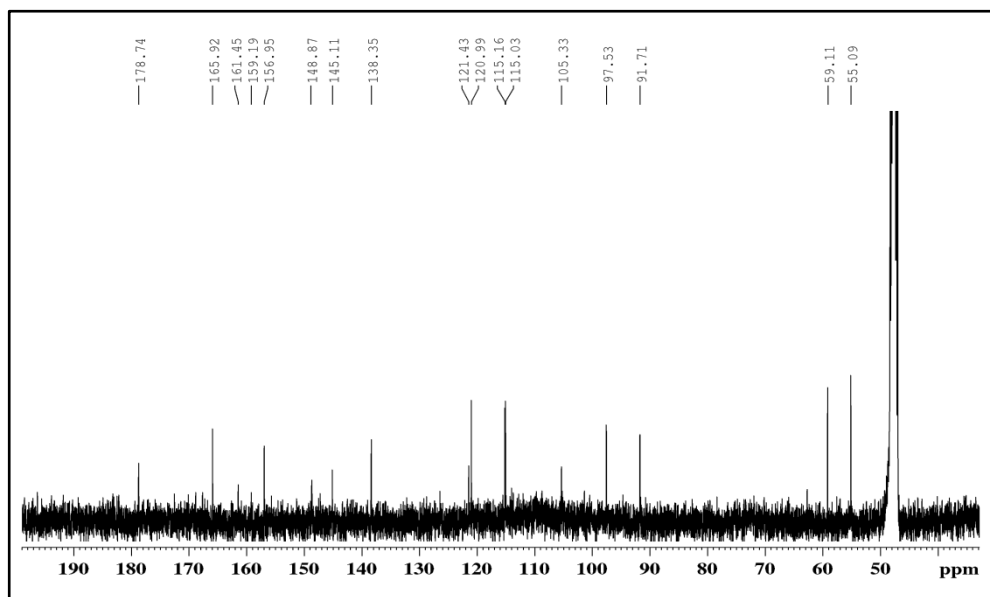


Figure.3.34 ^{13}C -NMR spectra of 3,7-di-*O*-methylquercetin (**11**)

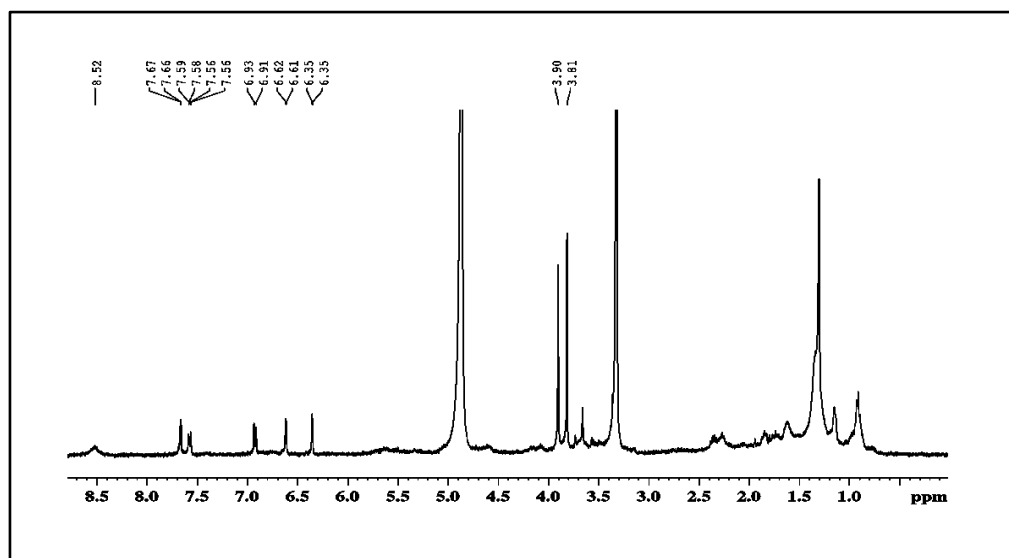


Figure.3.35 ^1H -NMR spectra of 3,7-di-*O*-methylquercetin (**11**)

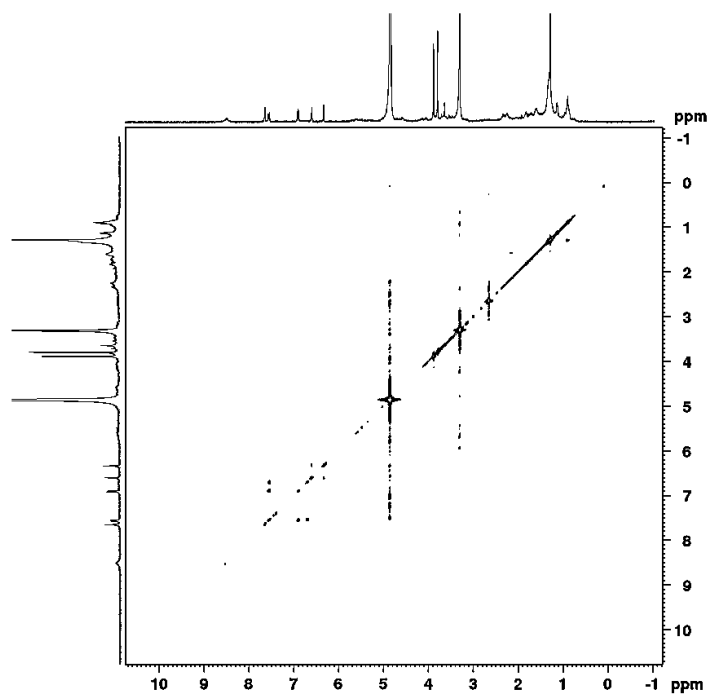


Figure. 3.36 COSY spectrum of 3,7-di-*O*-methylquercetin

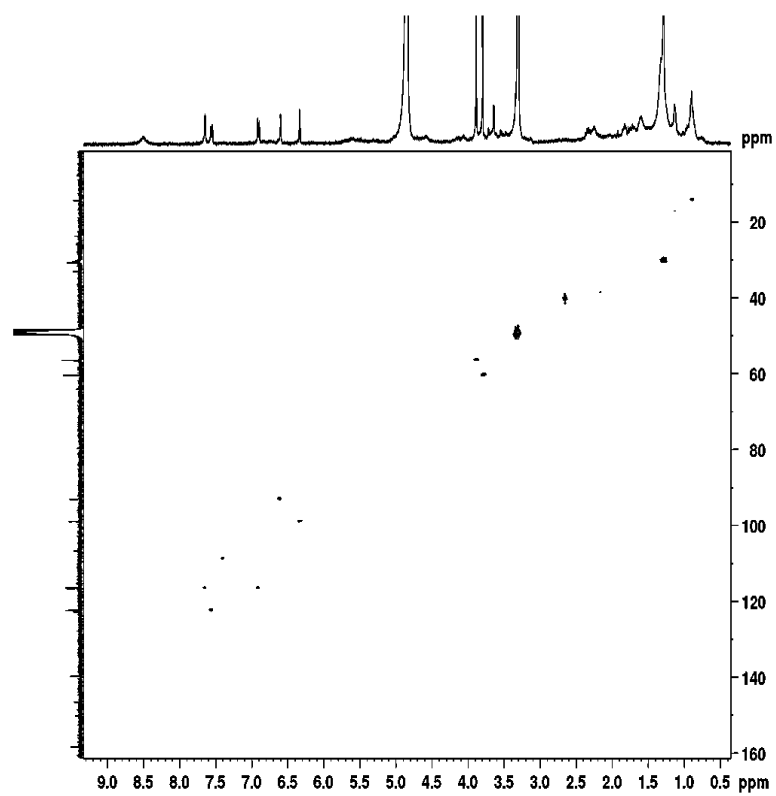


Figure 3.37 HSQC spectrum of 3,7-di-*O*-methylquercetin

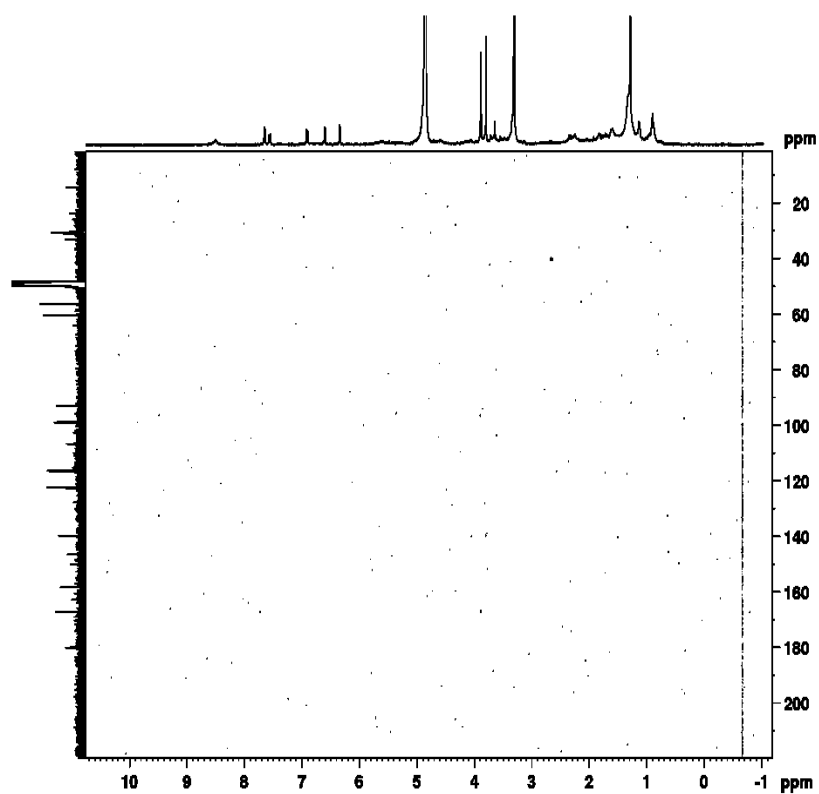


Figure 3.38 HMBC spectrum of 3,7-di-*O*-methylquercetin

3.3.4. Coumarins (12)

3.3.4.1 Scopoletin (12)

The structure of compound **12** was elucidated by ^1H - and ^{13}C -NMR as scopoletin (**12**) (Fig. 3.39).

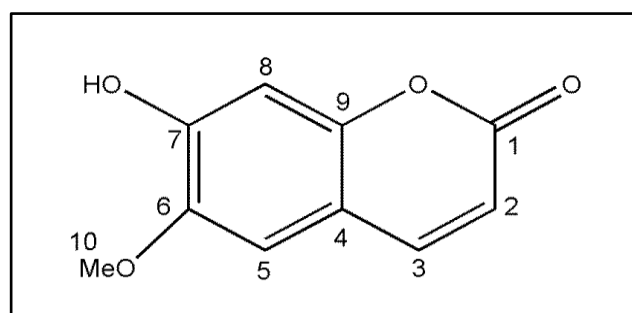


Figure 3.39 Structure of scopoletin (**12**)

Library search on the ^{13}C -NMR data (Table 3.8 and Fig. 3.40) suggested this compound is a methoxylated coumarin. Based on its ^1H , ^{13}C -NMR and mass spectral data compound **12** was found to be scopoletin (Darmawan *et al.*, 2012).

The ^1H -NMR (Table 3.8 and Fig. 3.41) and ^{13}C -NMR (Table 3.8 and Fig. 3.40) spectra of compound **12** are indicative of an aromatic ring system. The ^1H -NMR spectrum revealed two coupled doublets at 6.21 ppm ($J=9.4$ Hz) and 7.87 ppm ($J=9.4$ Hz) that were assigned to H-2 and H-3, respectively. Similarly two singlets appeared at 7.11 ppm and 6.77 ppm that were assigned to H-5 and H-8, respectively. Finally a methoxyl signal was observed at 3.92 ppm (s) that was assigned to H-10.

The ^{13}C -NMR spectrum indicated the presence of 11 carbons. The carbonyl group of the ester moiety presented a resonance at 168.8 ppm. In the same ring, the signal at 117.6 ppm was assigned to C-2 and the signal at 144.8 ppm to C-3. In the aromatic ring C-5 and C-8 were assigned 108.5 ppm and 102.6 ppm, respectively. Three quaternary aromatic carbons (C-4, C-6 and C-7) were assigned to the signals at 110.8 ppm, 145.9 ppm and 150.2 ppm, respectively. Similarly, C-9 was assigned to the signal at 152.6

ppm. The methoxylated carbon at 55.4 ppm was attributed to C-10. Further structure was confirmed by HSQC (Fig. 3.42) and HMBC (Fig 3.43).

Colourless needles; UV (acetonitrile / H₂O) λ_{max} 226, 287 and 338 nm ESI-MS (positive ion mode): m/z 193 [M+H]⁺ consistent with a molecular formula C₁₀H₈O₄.

Scopoletin is a coumarin derivative that has been found in many medicinal plant species, for instance *Macaranga gigantifolia* Merr. (Darmawan *et al.*, 2012), *Ipomoea digitata* (Khan and Hossain, 2015). This is however the first report of its presence in *Nymphoides indica*.

Table 3.8 ¹H-NMR and ¹³C-NMR assignments for the scopoletin (**12**) recorded in methanol-*d*₄.

Position	δ ¹ H ppm; multiplicity; J Hz,	δ ¹³ C ppm
1		162.8
2	6.21; d; J= 9.4	117.6
3	7.87; d; J=9.4	144.8
4		110.8
5	7.11; s	108.5
6		145.9
7		150.2
8	6.77; s	102.6
9		152.6
10	3.92; s	55.4

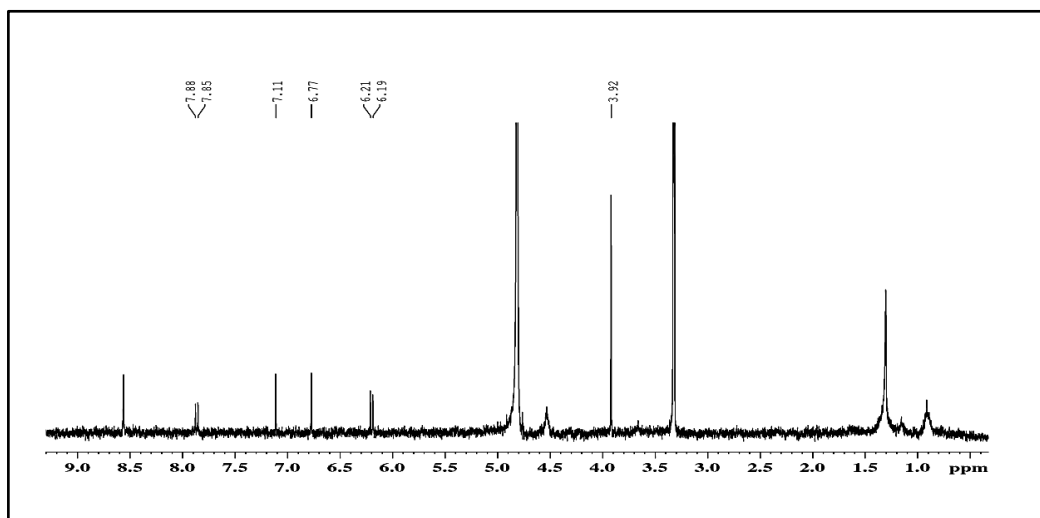


Figure.3.40 ¹H-NMR spectrum of scopoletin (12)

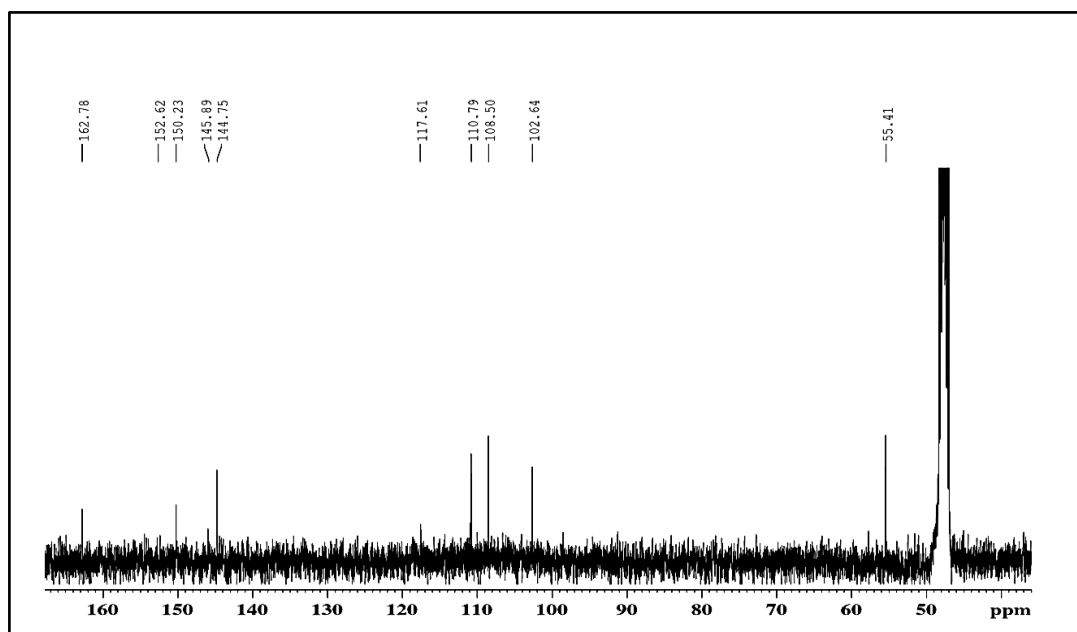


Figure.3.41 ¹³C-NMR spectrum of scopoletin (12)

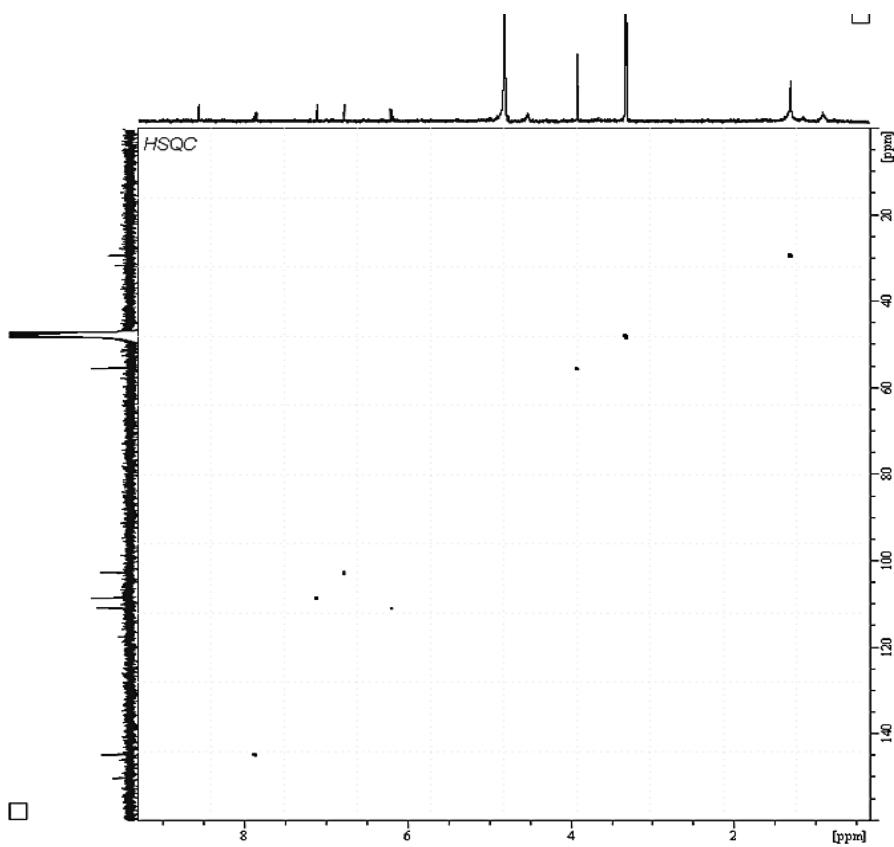


Figure 3.42 HSQC spectrum of scopoletin

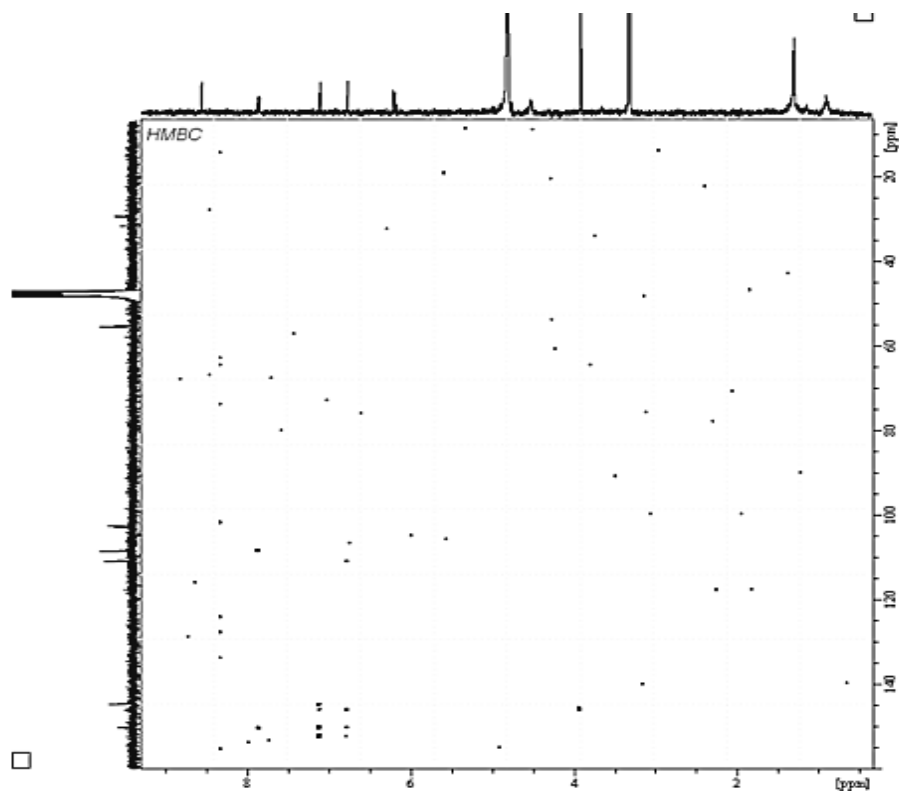


Figure 3.43 HMBC spectrum of scopoletin

3.3.5. Acyclic monoterpenes (13-14)

3.3.5.1 8-Hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (13)

The structure of compound **13** was elucidated by ¹H- and ¹³C-NMR (DEPT 135) as 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (**13**) (Fig 3.44).

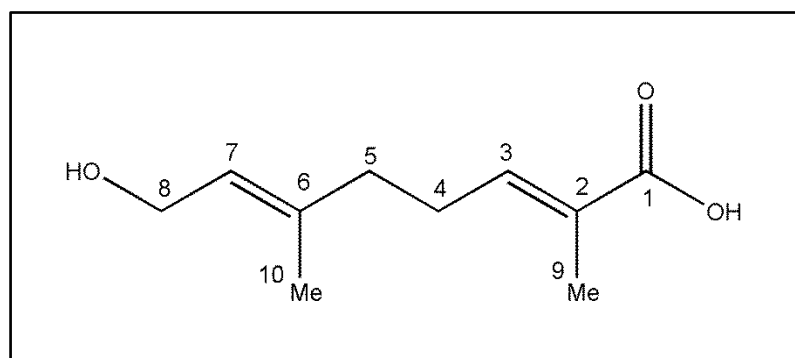


Figure 3.44 Structure of 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (**13**)

Library search on the ¹³C-NMR data (Table 3.9 and Fig. 3.45) suggested this compound is an acyclic monoterpene. Based on its ¹H, ¹³C-NMR and mass spectral data compound **13** was found to be 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (Iwagawa *et al.*, 1990).

The ¹H-NMR spectrum (Table 3.9 and Fig. 3.46) revealed signals at 1.68 ppm (s, H-10) and 1.80 ppm (s, H-9) that were assigned to two olefinic methyl groups. Two methylene protons at 2.14 ppm (t, *J*=7.3Hz, H-5) were coupled to other methylene protons at 2.30 ppm (q, *J*=7.3Hz, H-4), which were in turn coupled with a proton at 6.67 ppm (m, H-3). A doublet at 4.09 ppm (H-8) was attributed to allylic methylene protons that were found adjacent to a hydroxyl group. These protons were coupled with an olefinic proton at 5.39 ppm (m, H-7). The olefinic protons were further coupled with methyl protons at 1.68 ppm. Comparison with related monoterpenes and the acyl side chains of the seco-iridoids **6** – **8** discussed above confirmed the *E*-configuration of both double bonds.

The ^{13}C -NMR spectrum (Table 3.9 and Fig 3.45) revealed 10 carbon signals. The carbonyl group of the ester moiety showed a signal at 171.1 ppm that was assigned to C-1. The signals at 11.2 ppm and 14.8 ppm were attributed to the methyl groups at C-9 and C-10, respectively. The methylene signals at 26.6 ppm, 37.8 ppm and 58.0 ppm were assigned to C-4, C-5 and C-8, respectively. Similarly two methine signals at 140.8 ppm and 124.2 ppm were attributed to C-3 and C-7, respectively. The signals at 128.6 ppm and 137.1 ppm were assigned to C-2 and C-6.

Yellowish powder; UV (acetonitrile / H_2O) λ_{max} 222 nm; ESI-MS (positive ion mode) m/z 207 $[\text{M}+\text{Na}]^+$ consistent with molecular formula $\text{C}_{10}\text{H}_{16}\text{O}_3$.

Compound **13** was previously reported from *Radermachia sinica* (Iwagawa *et al.*, 1990) and was found as a substituent in some iridoids (Junior, 1983). Interestingly, during the current investigation, compound **13** was present as a side chain in compound **6**.

Table 3.9 ¹H-NMR and ¹³C-NMR assignments for 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (**13**) recorded in methanol-*d*₄.

Position	δ ¹ H ppm; multiplicity; J Hz,	δ ¹³ C ppm
1		171.1
2		128.6
3	6.67; m	140.8
4	2.30; q; 7.3	26.6
5	2.14; t; 7.3	37.8
6		137.1
7	5.38; m	124.2
8	4.04; d; J= 7.0	58.0
9	1.80; s	11.2
10	1.68; s	14.8

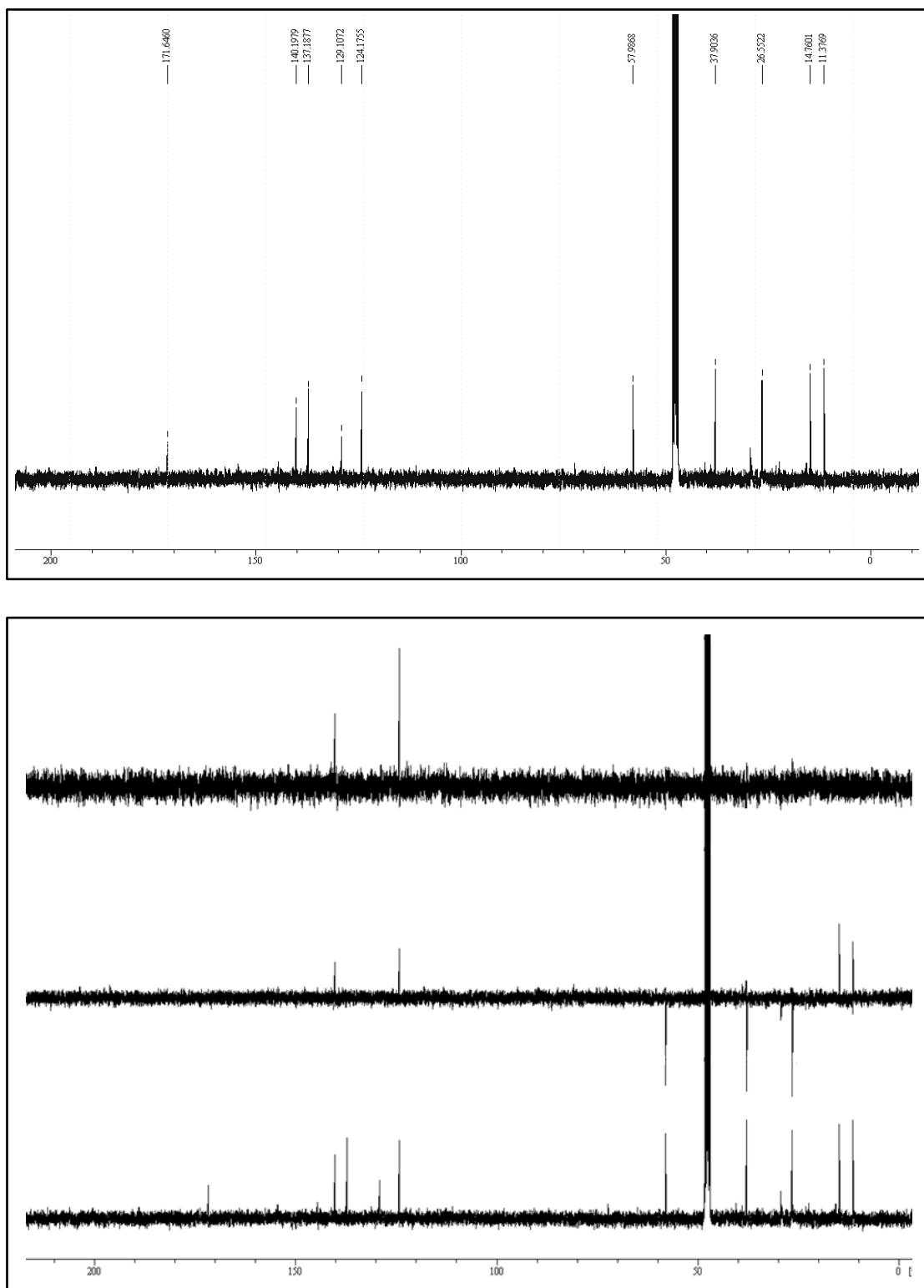


Figure 3.45 ^{13}C -NMR , DEPT-135 and DEPT-90 spectra of 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)- octadienoic acid (**13**)

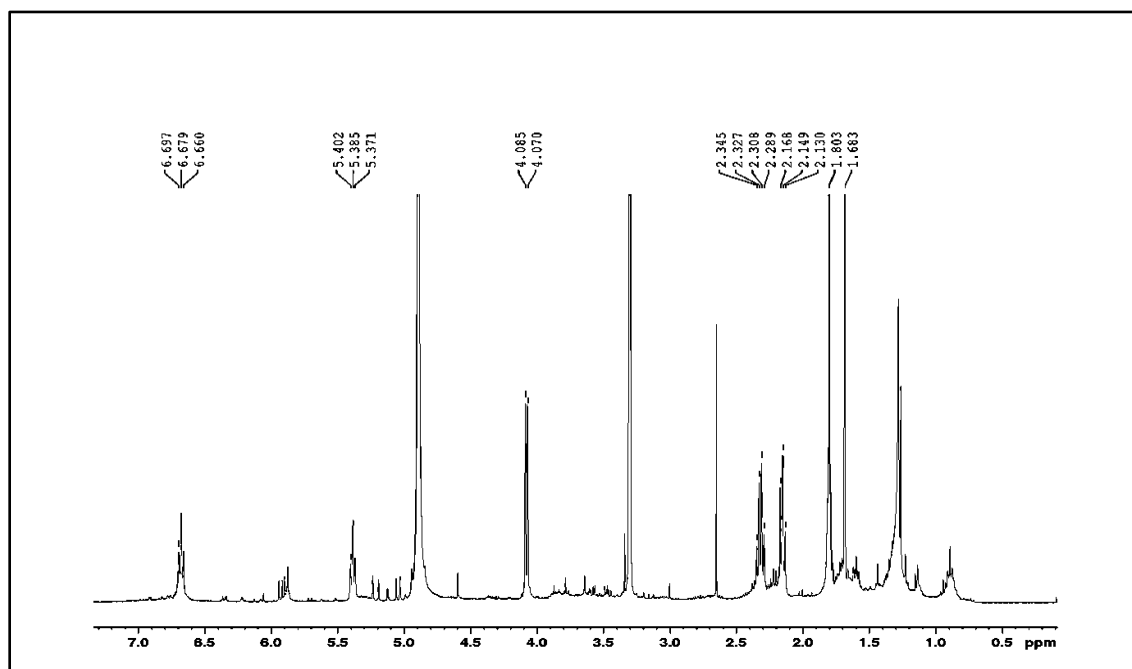


Figure 3.46 ¹H-NMR spectrum of 8-hydroxy-2,6-dimethyl-(2*E*,6*E*)-octadienoic acid (**13**)

3.3.5.2 8-Hydroxy-2,6-dimethyl-2-enoate (**14**)

The structure of compound **14** was elucidated by ^1H - and ^{13}C -NMR (DEPT-135) as 8-hydroxy-2,6-dimethyl-2-enoate (**14**) (Fig. 3.47).

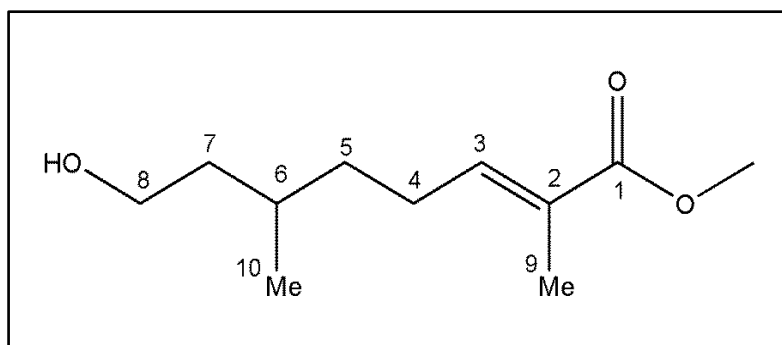


Figure 3.47 Structure of 8-hydroxy-2,6-dimethyl-2-enoate (**14**)

Library search on the ^{13}C -NMR data (Table 3.10 and Fig 3.48) suggested this compound is an acyclic monoterpene. Based on its ^1H (Table 3.10 and Fig. 3.49) and ^{13}C -NMR spectra (Table 3.10 and Fig. 3.48), and its mass spectral data compound **14** was found to be 8-hydroxy-2,6-dimethyl-2-enoate (Otsuka *et al.*, 1994).

In ^1H -NMR the signals at 0.94 ppm (d, $J=7$, H-10) and 1.84 ppm (d, $J=1$, H-9) were assigned to two olefinic methyl groups and the signal at 3.71 ppm (s) was attributed to a methoxyl group (-COOMe). A methylene group at 2.26 ppm (m, H-4) was coupled to a proton at 6.77 ppm (m, H-3). The multiplet signal at 1.25 - 1.65 ppm was assigned to H-5 (2H), H-6 (1H) and H-7 (2H), respectively. A multiplet at 3.60 ppm (H-8) was attributed to a methylene group adjacent to hydroxyl functionality.

The ^{13}C -NMR spectrum showed 11 carbon signals. The carbonyl group of the ester moiety was observed at 167.8 ppm. The signals at 11.0 ppm and 18.3 ppm were attributed to the methyl groups at C-9 and C-10, respectively, while the signal at 50.80 ppm was assigned to methoxylated protons adjacent to carbonyl moiety. The methylene signals of monoterpene at 25.7 ppm, 35.6 ppm, 39.2 ppm and 59.5 ppm were assigned to C-4, C-5, C-7 and C-8 respectively. Similarly two methane signals appeared

at 142.7 ppm and 29.1 ppm were attributed to C-3 and C-6 respectively. Finally the signal at 127.1 ppm was assigned to C-2. The *E* configuration of 2,3-double bond was defined from the ¹³C-NMR chemical shift of the C-9 methyl group at 11.0 ppm, similar to compound **13**.

Yellowish powder; UV (acetonitrile / H₂O) λ_{max} 218 nm; ESI-MS (positive ion mode) *m/z* 201 [M+H]⁺ consistent with a molecular formula C₁₁H₂₀O₃.

Compound **14** was previously reported from *Linaria japonica* (Otsuka *et al.*, 1994) in its glycosylated form. This is first report of its occurrence in *N. indica*, although the possibility cannot be excluded that the methyl ester is formed as an artefact during extraction, and the free carboxylic acid is the genuine compound.

Table 3.10. ^1H -NMR and ^{13}C -NMR assignments for 8-hydroxy-2,6-dimethyl-2-enoate (**14**) recorded in methanol- d_4 .

Position	δ ^1H ppm; multiplicity; J Hz,	δ ^{13}C ppm
1		168.8
2		127.1
3	6.77; m	142.7
4	2.26; m	25.7
5	1.25-1.65; m	35.6
6	1.25-1.65; m	29.1
7	1.25-1.65; m	39.2
8	3.60; m	59.5
9	1.84; d; $J=1.0$	11.0
10	0.94; d; $J=7.0$	18.3
OMe	3.71; s	50.8

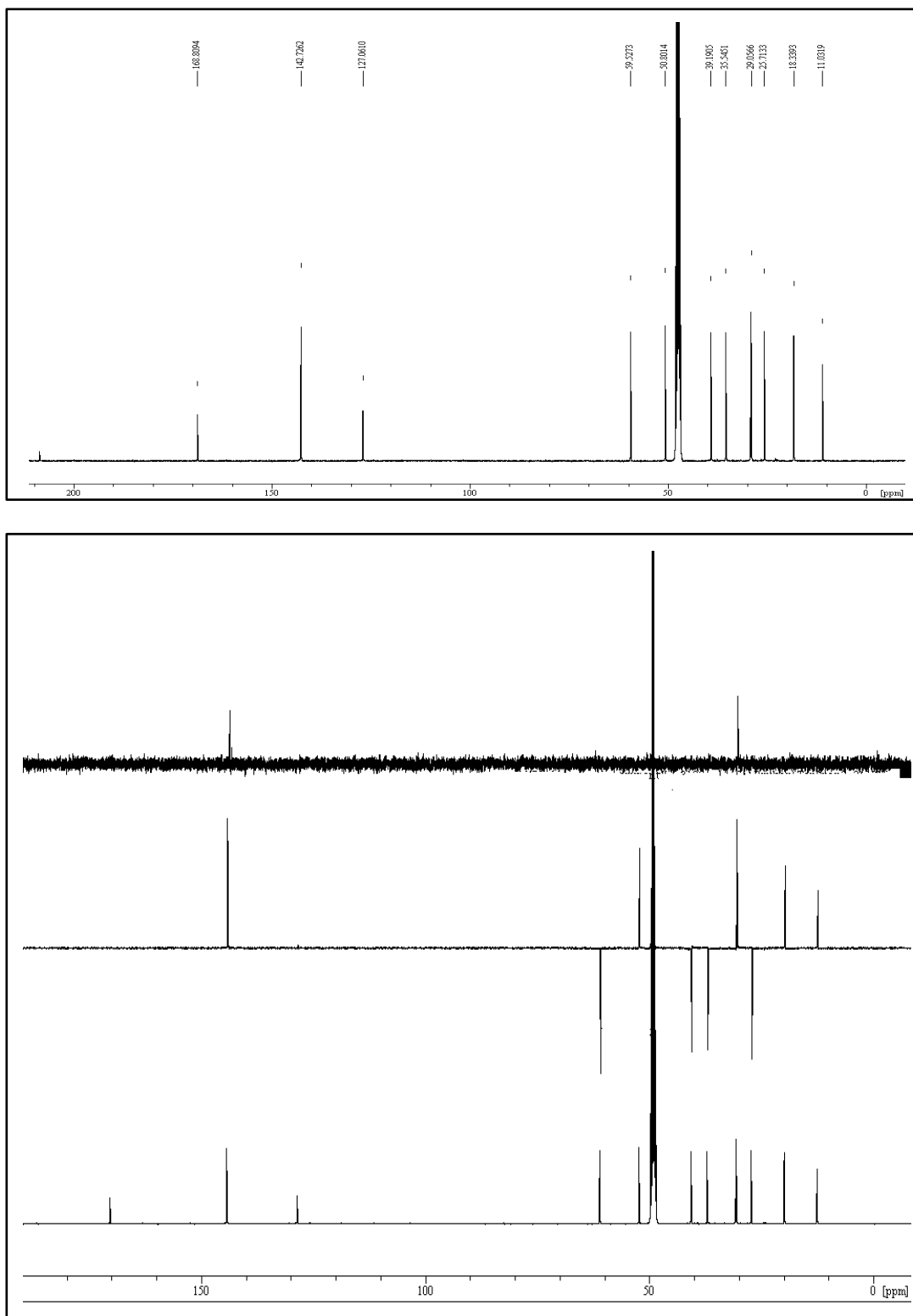


Figure 3.48 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of 8-hydroxy-2,6-dimethyl 2-enoate (**14**)

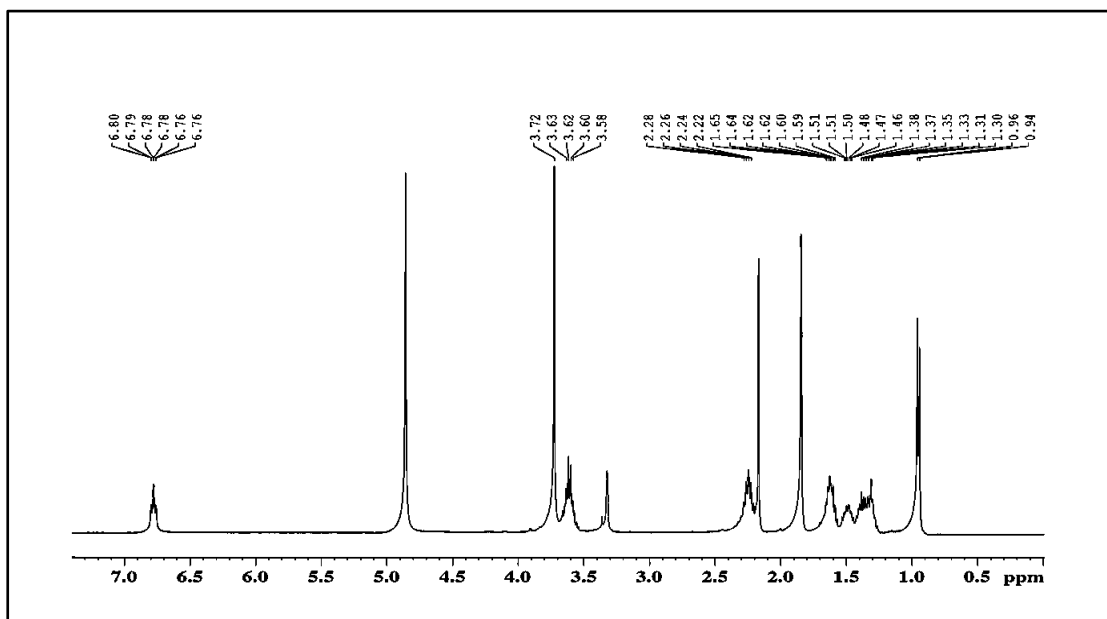


Figure 3.49 ^{13}C -NMR and DEPT-135 spectra of 8-hydroxy-2,6-dimethyl-2-enoate (**14**)

3.4. BIOLOGICAL ACTIVITIES

3.4.1. Total Phenolic Content (TPC), Total Flavonoid Content (TFC) and DPPH-scavenging activity

All fractions were analysed for their total phenolic, total flavonoid content (TPC and TFC, respectively) and DPPH scavenging activity (Table 3.11). The TFC of the crude extract was 70.28 mg RUE/g extract, as calculated from the rutin calibration curve. In the different fractions the highest flavonoid contents were observed in the ethyl acetate fraction (73.44 mg RUE/g) followed by the methanol fraction (62.34 mg RUE/g). The chloroform (17.52 mg RUE/g), *n*-butanol (10.57 mg RUE/g) had comparably low flavonoid contents. Similarly, the total phenolic content (TPC) of the total extract was 28.88 mg GAE/g as calculated from the gallic acid calibration curve, whereas the ethyl acetate (34.1 mg GAE/g) and methanol 90% (31.87 mg GAE/g) subfractions contained slightly higher amounts. This comparably high phenolic and flavonoid contents were indicative of possibly high antioxidant activities in the corresponding fractions. Indeed the ethyl acetate fraction presented moderate antioxidant potential (IC₅₀ 81 µg/ml), followed by the methanol fraction (IC₅₀ 97 µg/ml) and chloroform fractions (IC₅₀ 119 µg/ml), respectively. These findings were in accordance with previous reports that correlate the antioxidant activities of plant extracts with high flavonoid and phenolic contents (Yoo *et al.*, 2008; Pathirana and Shahidi, 2005; Goulasa *et al.*, 2014). Furthermore, in an attempt to have an estimate of TPC activity, we compared our findings with a more broader investigation that involved a wide range of medicinal plants (30 in total) and categorised total phenolic contents into ranks (very low to very high phenolic contents). Upon comparison of our TPC findings, we could emphasise that total phenolic contents of *Nymphoides indica* lie in “very low”- “low” levels (Dudonne’ *et al.*, 2009). Likewise based on the findings of above cited work for antioxidant activity (DPPH scavenging), the *N.indica* leaves fractions was considered as “moderate” in inhibition for DPPH.

Table 3.11 Total phenolic (TPC), total flavonoid contents (TFC) and antioxidant activities of *N.indica* fractions.

Sample	Extract Weight (g)	TPC	amount of GAE (mg)	TFC	amount of RUE (mg)	DPPH	
		(mg GAE/g extract) ^a		(mg RUE/g extract) ^b		% inhibition ^c	IC ₅₀ µg/mL
Total Extract	88.61	28.88 ± 0.42	2561	70.28 ± 1.0	6221	53	121
MeOH 90%	11.2	31.87 ± 0.83	360	62.34 ± 0.9	698	55	97
Chloroform	2.03	9.76 ± 0.96	19	17.52 ± 1.22	35	52	119
Ethyl acetate	1.2	34.1 ± 0.54	41	73.44 ± 1.23	88	60	81
<i>n</i> -Butanol	19.58	16.38 ± 1.07	193	10.57 ± 0.39	818	48	-
Quercetin							6

(a) Total phenolic contents, gallic acid equivalents

(b) Total flavonoid contents, rutin equivalents

(c) at 128 µg/mL (final concentration).

3.4.2 Inhibition of AGEs formation and α -glucosidase inhibition by crude extracts

The different fractions were tested for potential antiglycation and α -glucosidase inhibitory activity (Table 3.12). A mild to moderate antiglycation activity was observed for the crude extract at 100 μ g/mL (33%). Likewise all tested fractions presented moderate (24-36%) inhibition of the formation of AGEs inhibition. Similarly mild α -glucosidase inhibitory activity was observed for the total extract at 834 μ g/mL (13%), whereas the highest inhibition was shown by the methanolic (31%) and *n*-butanol fractions (25%) fractions at the same concentration. At higher test concentrations the inhibitory activity did not further increase, and the 50% inhibition level could not be reached.

Table 3.12. Anti-glycation and α -glucosidase inhibitory effects of *N. indica* fractions.

Sample	Antiglycation	α -glucosidase inhibition
	% Inhibition ^a	% Inhibition ^b
Total Extract	33	13
MeOH 90%	36	31
Chloroform	23	12
Ethyl acetate	27	17
<i>n</i> -butanol	24	25
Standard inhibitor	78	67 ^b

^a at 100 μ g/mL, Quercetin IC₅₀ 0.23 mM

^b at 834 μ g/mL, Acarbose IC₅₀ 0.26 mM

3.4.3 Antimicrobial activity of crude extracts

The crude extracts were evaluated for antibacterial and antifungal activities. The highest levels of antimicrobial and antifungal activities were observed for the *n*-hexane fraction, i.e. an IC₅₀ of 19.5 µg/mL against *S. aureus* and 32 µg/mL against *M. canis*. Similarly the methanol 90% fraction was active (IC₅₀ 36.4 µg/mL) against *S. aureus* and cytotoxic (IC₅₀ 38.9 µg/mL) against MRC-5 cells (Table 3.13).

Table 3.13 Antibacterial, antifungal and cytotoxic activities of *N. indica* fractions.

Sample	IC ₅₀ µg/mL				
	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>M. canis</i>	MRC-5
MeoH 90%	36.4	>64	>64	>64	38.9
Chloroform	>64	>64	>64	>64	>64
Ethyl acetate	>64	>64	>64	>64	>64
<i>n</i> -Hexane	19.5	>64	>64	32	>64
<i>n</i> -butanol	>64	>64	>64	>64	>64
Aqueous	>64	>64	>64	>64	>64

Reference: Tamoxifen(MRC-5) IC₅₀ 11.4 µg/mL; erythromycin (*S.aureus*)IC₅₀ 11.2 µg/mL, Trimethoprim (*E.coli*) IC₅₀ 0.25 µg/mL; Miconazole(*C. albicans*) IC₅₀ 5.99 µg/mL; Terbinafin (*M.canis*) IC₅₀ 0.11 µg/mL.

MRC-5: Human fetal lung fibroblasts; *S. aureus*: *Staphylococcus aureus*; *E.coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*

3.4.4. Biological activities of isolated constituents

In view of the antimicrobial activity of the *n*-hexane fraction, compounds **1** – **5** were evaluated in an integrated screening panel. Only azelaic acid (**1**) was active against *S. aureus* with an IC₅₀ value of 55.1 μM (vs. 11.2 μM for erythromycin) (Table 3.14). It also showed weak antiplasmodial activities against *P. falciparum* K1, *T. brucei* and *T. cruzi*, but it was also cytotoxic against MRC-5 cells in the same concentration range (IC₅₀ 32.2 μM). Compounds **2**–**5** showed comparable activity against some of the protozoal parasites. Lipophilic compounds such as medium and long chain fatty acids have been previously reported to contribute to antiprotozoal and antifungal activities (Krugliak *et al.*, 1995; Wang & Johnson, 1992; Ozcelik *et al.*, 2005a; Orhan *et al.*, 2009). It was therefore concluded that the activity of the *n*-hexane fraction of *N. indica* was mainly due to presence of long chain mono and dicarboxylic fatty acids and fatty alcohols that may act synergistically.

Three seco-iridoid glucosides (**6**–**8**) were isolated from the methanolic fraction. In the antimicrobial screening panel, none of the secoiridoids presented activity in the set test ranges. Nevertheless, the possibility cannot be excluded that after oral intake the glycosidic moiety is hydrolysed, and that the resulting aglycones and their metabolites are active. However in the antidiabetic assays, compound **7** presented moderate antiglycation (54% inhibition at 0.67 mM, IC₅₀ 0.36 mM) and mild α-glucosidase inhibitory (37% at 1 mM) activities, whereas only moderate antiglycation activities were observed for compound **6** (49% inhibition at 0.67 mM) and **8** (41% inhibition at 0.67 mM). The Compounds **6** and **8** did not present any α-glucosidase inhibitory activity (Table 3.15). Based on the observed antiglycation activities however, which are higher than the positive control aminoguanidine, a compound with proven clinical activity, it could be assumed that the isolated secoiridoids may have potential anti-AGEs activities. Also in this case the possibility cannot be excluded that after oral intake the glycosidic moiety is hydrolysed, and that the resulting aglycones and their metabolites are active.

In the antimicrobial screening panel, among the flavonoids, compound **9** showed weak antiprotozoal activities against *Trypanosoma brucei* (IC₅₀ 8.4 μM), *Trypanosoma cruzi*

(IC₅₀ 30.0 μM) and *Leishmania infantum* (IC₅₀ 32.5 μM), accompanied however by PMM (peritoneal murine macrophages) cytotoxicity.

Compound **9** is considered as the major flavonoid in *N. indica* (Bohm *et al.*, 1986) with no previous evidence of pharmacological properties. Also during the current investigation, it was collected in comparatively large amount that shows its relative profusion in leaves and this may contribute to the antiprotozoal activities of *N. indica* extracts. A number of reports (Muzitano *et al.*, 2006_a; Muzitano *et al.*, 2006_b) covering the anti-leishmanial activities of flavonoids support these findings. Despite the fact that flavonoids are major contributors of plants to a wide array of activities including cytotoxicity and antimicrobial activities (Lovkova *et al.*, 2001; Cushnie and Lamb 2005), the isolated flavonoids, however, presented rather moderate IC₅₀ values and were therefore not considered for further antimicrobial evaluation.

In the antidiabetic assays the isolated flavonoids presented moderate activities. The highest antiglycation activity was shown by compound **10** (54% inhibition at 0.67 mM, IC₅₀ 0.42 mM) followed by compound **9** (44% inhibition). These compounds also presented mild to moderate α-glucosidase inhibitory activities. Compound **10**, however, did not present any α-glucosidase inhibition (Table 3.15). Compound **11** was inactive in antimicrobial screening, but presented mild antiglycation (39% inhibition) and α-glucosidase inhibition (27% inhibition). Indeed it has been established that for antiglycation activity of flavonoids there are some essential features supporting the activity. Phenolic groups at C-3', C-4', C-5, and C-7 positions and methylation or glucosylation of the hydroxyl groups are considered important for the activity of flavones, flavonols and flavanones (Matsuda *et al.*, 2003). In addition it has been established that oxidative injury associated with degenerative diseases can be inhibited by antioxidants and radical scavengers (Nakagawa *et al.*, 2002). Compounds or plant extracts having antioxidative properties may also prevent AGEs formation (Yamaguchi *et al.*, 2000, Kiho *et al.*, 2004; Tung *et al.*, 2007). It has been reported before that extracts containing compounds with different mechanisms of action may be more effective than isolated constituents (Xi *et al.*, 2008, Yagi *et al.*, 2013), as also observed during this investigation.

The coumarins are known for their cytotoxic and antimicrobial activities (De Souzaa *et al.*, 2005; Smyth *et al.*, 2009). The isolated coumarin (compound **12**) was identified as scopoletin. Antimicrobial activities have been reported for scopoletin (Ragasa and Hofilena 2011; More *et al.*, 2012) but during current investigation this could not be confirmed for the test organisms (IC₅₀ values >64 µM) (Table 3.13). However, as the crude methanolic fraction of *N. indica* showed significant cytotoxic and antimicrobial activities (Table 3), this could possibly due to the presence of flavonoids and coumarins. It could therefore be hypothesised that various constituents may contribute synergistically towards the antimicrobial activity of *N. indica* as indeed sometimes isolation of the components from the extract may cause a decrease in desired activity (Orhan *et al.*, 2009).

Finally two acyclic monoterpenes (**13-14**) were isolated from the *n*-butanol fractions of *N. indica* leaves. These monoterpenes were not active in the antimicrobial screening panel. However, moderate antiglycation activity and α-glucosidase inhibition (Table 3.15) was observed. As the *n*-butanol fraction also showed antiglycation activity (24% inhibition) (Table 3.12), it can be hypothesised that the AGEs inhibitory potential of the *n*-butanol fraction is mainly contributed by flavonoids and monoterpenes (Al-Musayeib *et al.*, 2011; Balasubramaniam and Anuradha, 2011; Joglekar *et al.*, 2013).

Table. 3.14 Antimicrobial, antifungal, cytotoxic and antiprotozoal activities (IC₅₀ μM) of isolated compounds from *N. indica* fractions.

Compound no.	IC ₅₀ μM									
	<i>MRC5</i>	<i>Pf-K1</i>	<i>T. bruc</i>	<i>T. cruz</i>	<i>L. inf</i>	<i>PMM</i>	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>A. fumigatus</i>
1	32.2	62.2	32.6	34.6	> 64	> 64	55.1	> 64	> 64	> 64
2	28.5	> 64	> 64	33.5	> 64	> 64	> 64	> 64	> 64	> 64
3	32.2	> 64	> 64	32.8	> 64	> 64	> 64	> 64	> 64	> 64
4	> 64	> 64	32	31.5	50.8	> 64	> 64	> 64	> 64	> 64
5	32.6	> 64	> 64	33.6	32.4	32.0	> 64	> 64	> 64	> 64
6	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
7	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
8	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
9	> 64	> 64	8.4	30.0	32.5	32.0	> 64	> 64	> 64	> 64
10	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
11	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
12	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
13	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
14	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64

Reference: Tamoxifen (*MRC-5*), IC₅₀ 11.35 μM; Suramine (*T. bruc*), IC₅₀ 0.03 μM, Fungizone (*L. inf*), IC₅₀ 1.15 μM; Chloroquine (*Pf-K1*), IC₅₀ 0.16 μM, Benznidazole (*T. cruz*), IC₅₀ 3.27 μM; Erythromycin (*S. aureus*), IC₅₀ 11.31 μM; Chloramphenicol (*E. coli*), IC₅₀ 4.88 μM; Miconazole (*C. albicans*), IC₅₀ 10.5 μM; Terbinafine (*A. fumigatus*), IC₅₀ 0.79 μM.

MRC-5: human fetal lung fibroblasts; *Pf-K1*; *Plasmodium falciparum*; *T. bruc*: *Trypanosoma brucei*; *L.inf*: *Leishmania infantum*; *PMM*: peritoneal murine macrophages; *T. cruz*: *Trypanosoma cruzi*; *S. aureus*: *Staphylococcus aureus*; *E. coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*; *A. fumigatus*: *Aspergillus fumigatus*

Table 3.15 Anti-glycation and α -glucosidase inhibitory effect of isolated compounds from *N. indica*

Compound no.	Protein glycation		α -glucosidase	
	% inhibition ^a	IC ₅₀ mM	% inhibition ^b	IC ₅₀ mM
6	49	-	na	-
7	54	0.36	37	-
8	41	-	na	-
9	44	-	45	-
10	54	0.42	na	-
11	39	-	27	-
12	nt	-	nt	-
13	47	-	38	-
14	51	0.61	40	-
Standard		0.23 ^c		
Inhibitor		1.75 ^d		0.26 ^e

na not active

nt not tested (low amount)

^a (at 0.67 mM)

^b (at 1 mM)

^c Quercetin

^d Aminoguanidine

^e Acarbose

3.5. SUMMARY AND CONCLUSION

Nymphoides indica (L.) Kuntze (Menyanthaceae) is traditionally used in many ethnomedicinal formulations in the Indian subcontinent including Pakistan. However, scientific data reporting its constituents are poor. This study aimed at evaluating its traditional use and phytochemical constituents, as a part of running programmes on antimicrobial, antioxidant and antidiabetic properties (inhibition of alpha-glucosidase activity; inhibition of Advanced Glycation Endproducts - AGEs) of medicinal plants and natural products. Various extracts and fractions of different polarity of *N. indica* leaves showed moderate activities in these assays. Phytochemical investigations of the extracts and fractions resulted in the isolation and identification of 5 lipophilic compounds, i.e. the dicarboxylic acids azelaic acid (nonanedioic acid) (**1**) and 4-methyl-heptanedioic acid (**3**), the monocarboxylic acids hexadecanoic acid (**2**) and stearic acid (**5**) and the fatty alcohol hexadecanol (**4**); 3 seco-iridoids, i.e. 7-epiexaltoside (**6**), 6",7"-dihydro-7-epiexaltoside (**7**) (reported for the first time from nature) and menthiafolin (**8**); 3 flavonoids, i.e. 3,7-di-*O*-methylquercetin-4'-*O*-glucoside (**9**), 3-*O*-methylquercetin-7-*O*-glucoside (**10**) and 3,7-di-*O*-methylquercetin (**11**); scopoletin (**12**) and the monoterpenoids foliamenthic acid (**13**) and 6,7-dihydrofoliamenthic acid methyl ester (**14**). Compounds **1-5** showed moderate antimicrobial activities, whereas compound **9** presented mild antiprotozoal activities against *Trypanosoma brucei* (IC₅₀ 8 μM), *Leishmania infantum* (IC₅₀ 32 μM) and *Trypanosoma cruzi* (IC₅₀ 30 μM). Likewise promising antiglycation activities were shown by compound **7** (IC₅₀ 0.36 mM), **10** (IC₅₀ 0.42 mM) and **14** (IC₅₀ 0.61 mM). Finally mild to moderate α-glucosidase inhibition was shown by compounds **7, 9, 11** and **13 - 14**. It could be concluded that *N. indica* leaf extracts possess mild to moderate antimicrobial, antiprotozoal, antioxidant and antidiabetic activities. The traditional use of this plant species could be due to synergistic effects of its constituents, or in combinations with other plant species in traditional formulations.

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CHAPTER 4

PHYTOCHEMICAL AND BIOLOGICAL INVESTIGATIONS ON *KICKXIA RAMOSISSIMA*

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4.1 INTRODUCTION

Although *Nanorrhinum ramosissimum* (Wall.) Betsche is the accepted name of the species in the genus *Nanorrhinum* (family Plantaginaceae) (Adopted in 1984) (The Plant List, 2016), common synonyms are *Linaria ramosissima* Wall. (family Scrophulariaceae)(1831); *Linaria somalensis* Vatke (family Scrophulariaceae)(1882); *Kickxia ramosissima* (Wall.) Janchen (family Scrophulariaceae) (1933); *Kickxia somalensis* (Vatke) Cuf. (family Scrophulariaceae)(1963) and *Pogonorrhinum somalense* (Vatke) Betsche (family Scrophulariaceae)(1984) (Plants Jstore, 2016). Due to the fact that synonym *Kickxia ramosissima* is more commonly used in literature, it was adopted during current project. *Kickxia* (Scrophulariaceae) is a small genus of herbs geographically distributed from West-Africa to India (Khan and Aqil, 1993). The genus *Kickxia* is comprised of 47 species worldwide (Mabberley, 1997). *Kickxia ramosissima* (Wall.) Janch. is a perennial herb with numerous filiform branches, membranous leaves and yellow flowers (Pandya *et al.*, 2012). It is mainly found in rocky areas with shady places (Kirtikar and Basu 2005) and high salt contents (Mathur *et al.*, 1980) (Fig. 4.1). In Pakistan it has been reported from various places including the district Attock (Ahmad *et al.*, 2009), the district Karak (Khan *et al.*, 2011), and the Nara desert (Qureshi *et al.*, 2010).



Figure 4.1 *Kickxia ramosissima* natural habitat and herbarium sheet

Kickxia ramosissima has been used in the indigenous system of treatment of the Indian subcontinent (Vardhana, 2008). In Pakistan it is known as “Wal”, “Shin beeta” or “Khunger booti”. It is used for a number of ailments, for instance as diuretic, against kidney stones (Pandya et al., 2012), fever and rheumatism (Jain et al., 2008) and during management of snake and scorpion bites (Bole and Pathak, 1988). Traditionally in the Indian subcontinent including Pakistan this species has been reported as an effective remedy for diabetes mellitus (Qureshi and Bhatti, 2008; Ahmad et al., 2009; Qureshi et al., 2010; Patel and Sachdeva, 2014).

Despite its ethnomedicinal importance only a few *Kickxia* species worldwide were chemically investigated. This has resulted in the isolation of mainly flavonoids and iridoid glycosides (Yuldashev *et al.*, 1996; Khan and Aqil 1993; Khan *et al.*, 2001; Al-Rehaily *et al.*, 2006; Ahmad *et al.*, 2006; Ferhat *et al.*, 2010), fatty acids (Morteza-Semnani *et al.*, 2008) and a hexitol, D-mannitol (Khan and Aqil, 1993) (Fig. 4.2). *Kickxia ramosissima*, however, is one of the least explored species.

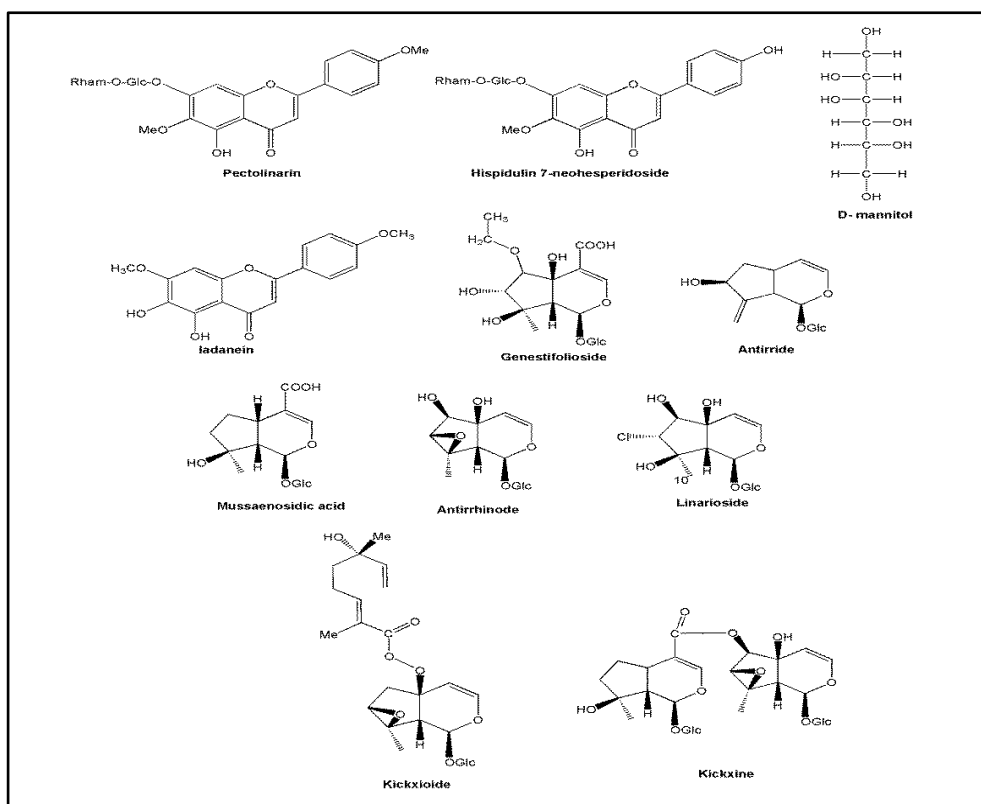


Figure 4.2. Compounds isolated from *Kickxia* sp.

4.2 EXTRACTION AND ISOLATION

4.2.1 Plant material

Kickxia ramosissima whole plant was collected in October 2012 from Takht-e-Nusrati, district Karak (KPK), Pakistan. Afterwards the herbarium sheets of the collected whole plant were submitted for identification at the Islamabad Herbarium of the Taxonomy Department, Quaid-I-Azam University, Islamabad, Pakistan, where the voucher specimen was deposited (voucher No. 48 CJ, accession no. ISL-44586). The whole plant was dried under shade followed by powdering and sieving through a 20 mesh filter (Fig. 4.3). All powdered material was stored below 20 °C till further use.



Figure 4.3 *Kickxia ramosissima* dried whole plant powder.

4.2.2. Extraction and fractionation

Powdered plant material (0.955 kg) was extracted with 90% (v/v) methanol by double cold maceration. The extract was filtered through Whatman No.1 filter paper using a vacuum pump. The collected filtrate was dried using a rotary evaporator under reduced pressure below 40 °C. The resultant semisolid material was lyophilised with a final yield of 83.54 g, and stored below 20 °C. The liquid-liquid partitioning was performed on the crude extract according to a standard extraction scheme (see chapter 2). After partitioning with different solvents as shown in the scheme, *n*-hexane (1.0 g), methanol

90% (3 g), chloroform (1.69 g), ethyl acetate (1.22 g), *n*-butanol (9.63 g) and aqueous fractions (67.0 g) were obtained. The collected fractions were dried under reduced pressure at 40 °C, lyophilized and stored below 20 °C.

4.2.3 Phytochemical analysis

Phytochemical screening of extracts and fractions for various compound classes was carried out using published methods (Sofowora, 1993, Trease and Evans, 1989, Harborne, 1973; Egwaikhide and Gimba, 2007; Farnsworth, 1966) with slight modifications for sterols, triterpenes, carbohydrates and saponins flavonoids, alkaloids, and tannins (chapter 2). Phytochemical screening of extracts and fractions indicated that flavonoids, tannins, saponins and triterpenes were present at different concentrations according to the fraction (Table 4.1). No alkaloids were detected. Fractions of medium polarity contained flavonoids and terpenes. The residual aqueous fraction was positive for tannins and saponins.

Table 4.1 Phytochemical analysis of *Kickxia ramosissima* fractions

Extract/ fraction	Phytochemicals				
	Alkaloids	Flavonoids	Tannins	Steroids/ triterpenoids	Saponins
MeoH 90%	-	+++	-	+++	-
Chloroform	++	+++	-	+++	-
Ethyl acetate	-	+++	-	+	-
<i>n</i>-Hexane	-	-	-	-	-
<i>n</i>-butanol	-	++	-	++	+++
Aqueous	-	-	+++	-	+++

profundly present: high degree of precipitation (dark coloration), moderately present: less degree of precipitation (medium colouration), slightly present: very low precipitation (very little colouration), absent: no change in colour (Senguttuvan et al., 2014)

Phytochemicals: ++++ profoundly present; ++ moderately present; + slightly present; - absent.

4.2.4. Thin layer chromatography

The NP (normal phase) TLC for all obtained fractions was performed using various solvent systems as mobile phase, including CH₂Cl₂/MeOH (70:30) with a few drops of NH₄OH and CH₂Cl₂/MeOH (75:25) for the methanolic fraction; CH₂Cl₂ / MeOH (78 : 22) for the chloroform and ethyl acetate fractions; *n*-hexane/CHCl₃ (93:7) for the *n*-hexane fraction; CH₂Cl₂/MeOH (25:75 or 36:64) the for *n*-butanol fraction and MeOH/CH₂Cl₂ (85:15) for the aqueous fraction, respectively. Developed TLC plates were examined under UV at 254 nm and 366 nm and after spraying with *p*-anisaldehyde. For instance, TLCs for the methanolic and ethyl acetate fractions are shown in Fig 4.4.

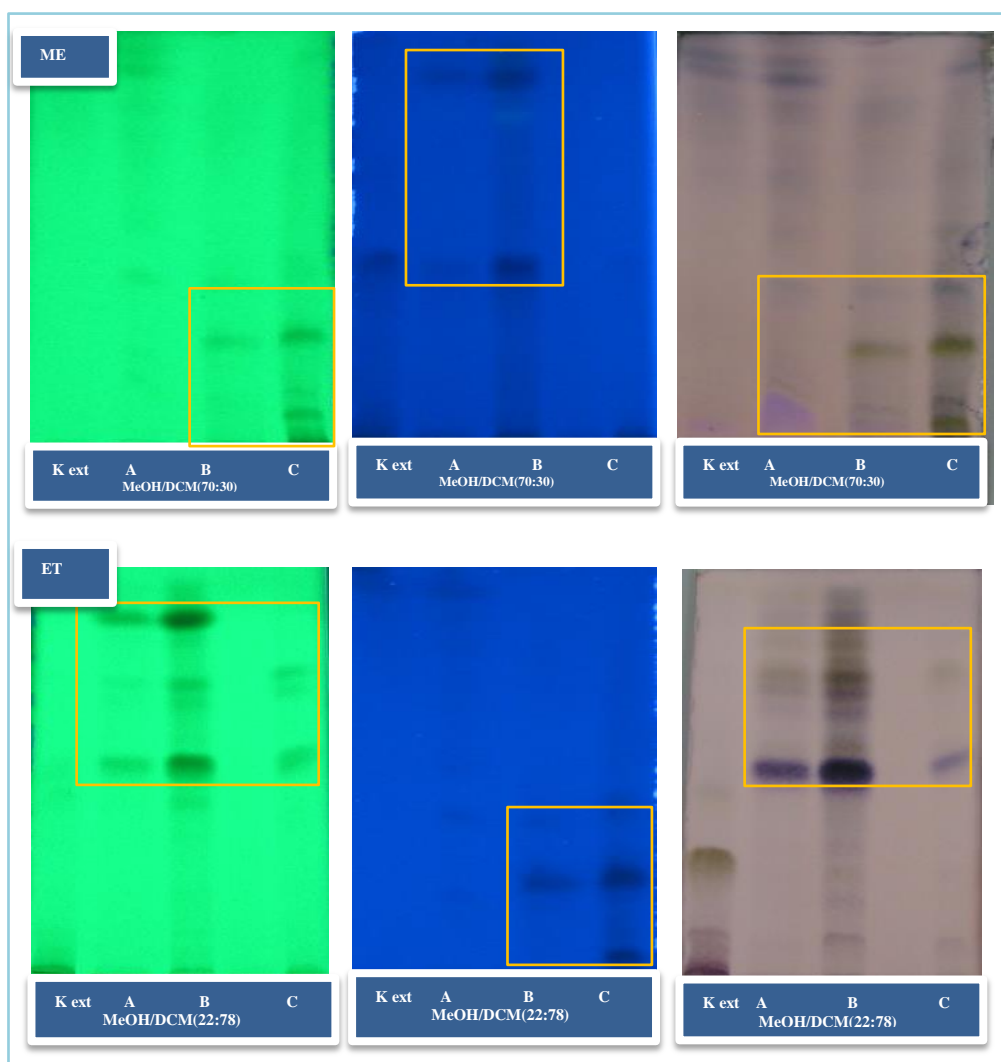


Figure 4.4. TLC profile of the total plant extract (K ext), methanolic (ME) and ethyl acetate (ET) fractions A, B, C on silica TLC (NP) plate at 254 and 366 nm after spraying with *p*-anisaldehyde reagent.

4.2.5. Flash chromatography

An aliquot of 0.8 g from the methanol 90% fraction was dissolved in 2 mL methanol and mixed with 1.1g silica; the mixture was dried with nitrogen gas. The dried extract was loaded on a prepacked Flash Grace Reveleris® silica cartridge of 80 g. The compounds were eluted using a gradient from ethyl acetate over methanol to methylene chloride. Based on UV and ELSD detection, multiple subfractions were collected (Fig. 4.5). All fractions were further analysed by TLC and similar fractions were combined. In this way 14 subfractions were obtained. Flash chromatography with similar conditions was repeated when necessary. Finally, based on TLC profiling, subfractions **KRM2** (200 mg), **KRM3**(150 mg), **KRM5** (210 mg), **KRM7** (200 mg) and **KRM14** (110 mg) were selected for further HPLC profiling.

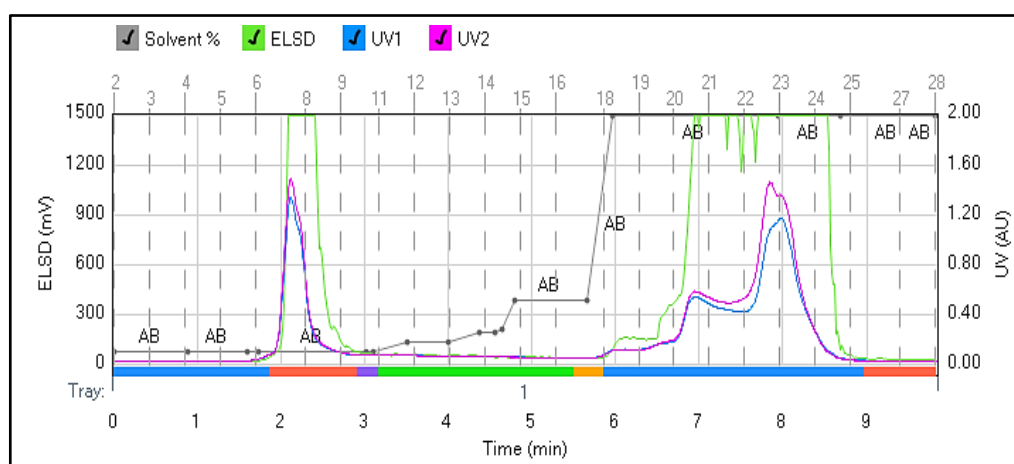


Figure 4.5 Flash chromatogram of the methanolic fraction.

Similarly an aliquot of 0.8 g from the ethyl acetate fraction was subjected to flash chromatography as above. The gradient used was from ethyl acetate over methanol to methylene chloride. Based on UV and ELSD detection different subfractions (8 in total) were collected (Fig 4.6). Finally a total of 4 subfractions was obtained and subfractions **KRET2** (210 mg), **KRET3** (160 mg), **KRET4** (100mg) and **KRET5** (270 mg) were selected for further HPLC profiling.

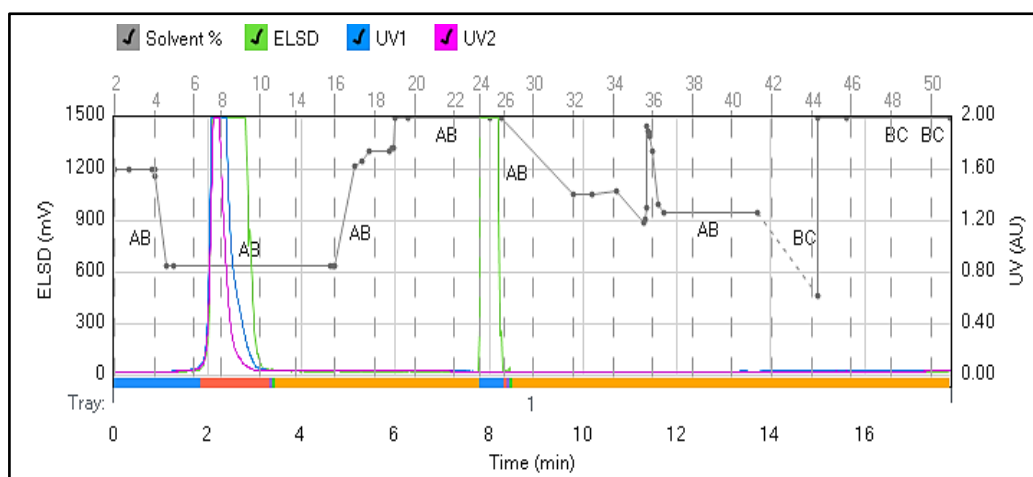


Figure 4.6. Flash chromatogram of the ethyl acetate fraction.

Likewise an aliquot of 0.8 g from the chloroform fraction was loaded on a flash column as discussed above. The compounds were eluted using a gradient from ethyl acetate over methylene chloride to methanol. Based on UV and ELSD detection different subfractions were collected (Fig 4.7). All fractions were analysed by TLC and similar fractions were combined as described above; in this way 8 subfractions were obtained. Based on TLC profiling subfractions **KRCL1** (135 mg), **KRCL2** (125 mg), **KRCL4** (160 mg), **KRCL7** (150 mg) and **KRCL8** (120 mg) were selected further HPLC analysis.

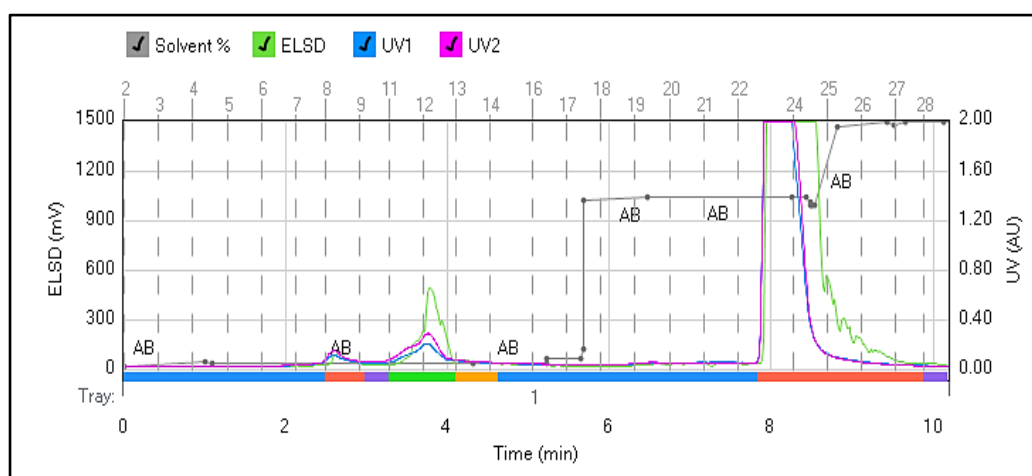


Figure 4.7 Flash chromatogram of the chloroform fraction.

Similarly an aliquot of 0.8 g from the *n*-butanol fraction was subjected to flash chromatography with a gradient from ethyl acetate over methylene chloride to methanol as previously explained. Finally 10 subfractions were obtained (Fig 4.8). Based on TLC analysis, subfractions **KRB2** (110 mg), **KRB4** (127 mg), **KRB5** (131 mg), **KRB6** (120 mg) and **KRB7** (126 mg) was selected for HPLC analysis.

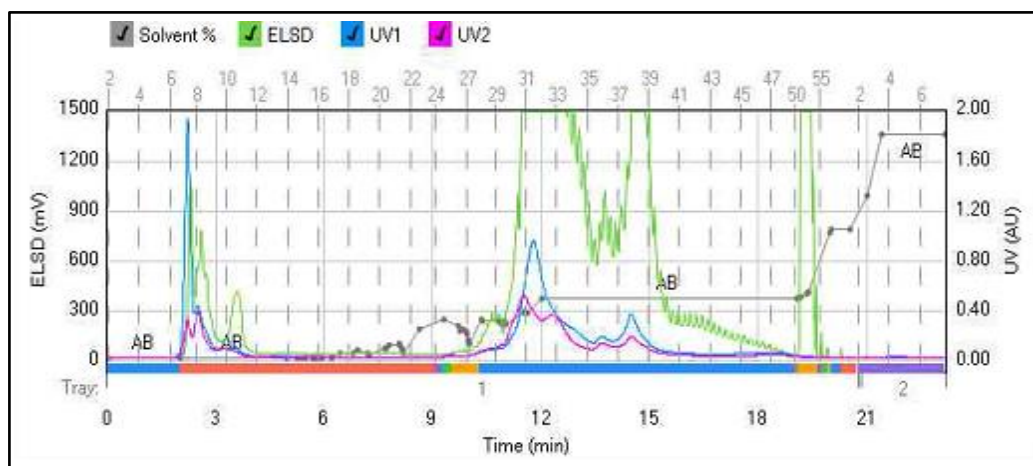


Figure 4.8 Flash chromatogram of the *n*-butanol fraction.

4.2.6. HPLC analysis and isolation of compounds

The chloroform fraction and all obtained subfractions **KRCL1**, **KRCL2**, **KRCL4**, **KRCL7** and **KRCL8** were analysed by HPLC using an optimized methanol/H₂O + 0.1% formic acid gradient, ranging from 15% acetonitrile to 80% in 50 min at a flow rate of 1 mL/min. Samples were prepared in a concentration range from 1-10 mg/ mL in methanol (Fig 4.9). The isolation of pure compounds was performed by means of semi-preparative HPLC-DAD-MS using the same gradient at 3 mL/min, yielding compounds **1** (7.5 mg), **2** (5.2 mg) and **3** (5.2 mg).

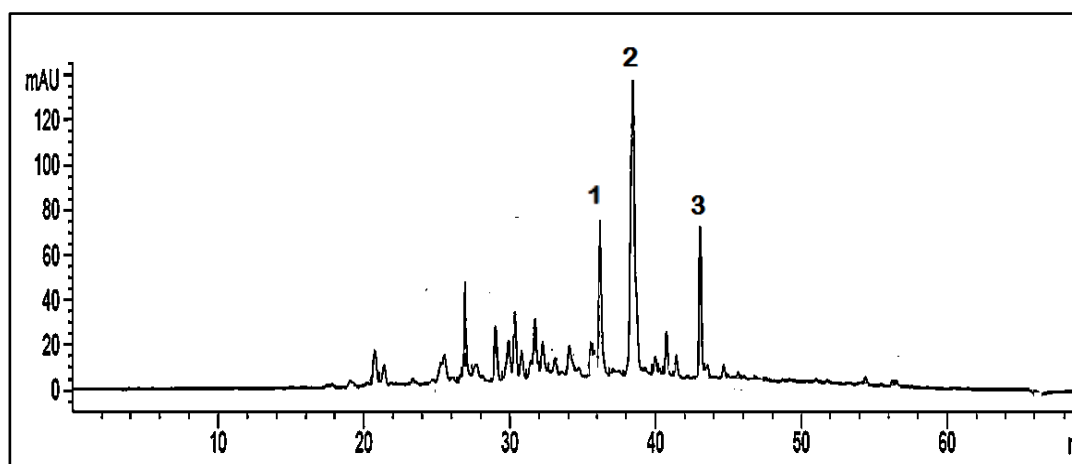


Figure 4.9 HPLC chromatogram of chloroform fraction **KRCL2** at 254 nm.

The crude ethyl acetate fraction and subfractions **KRET3** (250 mg), **KRET4** (180 mg) and **KRET5** (150 mg) were analysed by HPLC using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 15% acetonitrile to 100% in 60 min at a flow rate of 1 mL/min (Fig. 4.10). The isolation of pure compounds was performed by means of semi-preparative HPLC-DAD-MS using the same gradient at 3 mL/min. Compounds **2** (2.2 mg), **3** (5.2 mg) and **4** (5.7 mg) were finally obtained.

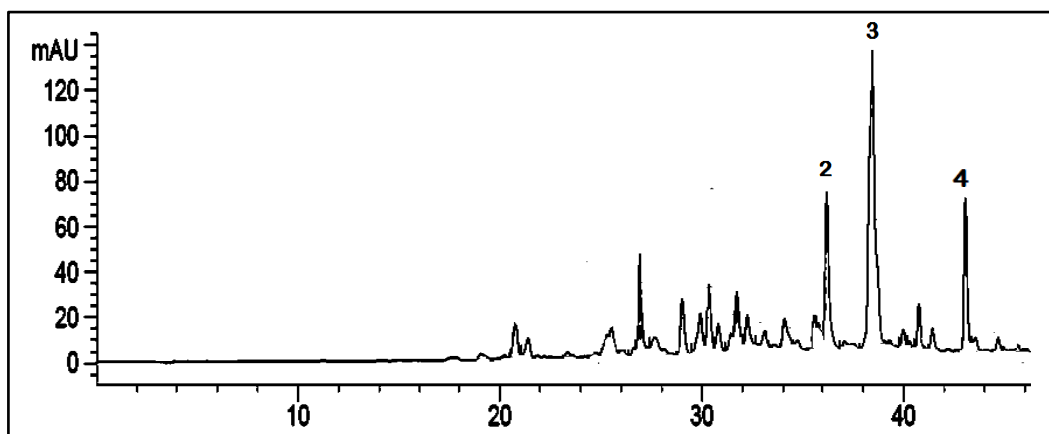


Figure 4.10 HPLC chromatogram of ethyl acetate fraction **KRET3** at 254 nm.

Similarly the crude methanol fraction and subfractions **KRM2**, **KRM3**, **KRM5**, **KRM7** and **KRM14** were analysed by HPLC using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 35% acetonitrile to 70% in 50 min at a flow rate of 1 mL/min (Fig. 4.11). The compounds were isolated by means of semi-preparative HPLC-DAD-MS system using the same gradient at a flow rate of 3 mL/min. Finally compounds **2** (3.3mg), **5** (4.5 mg) and **6** (3.2 mg) were isolated.

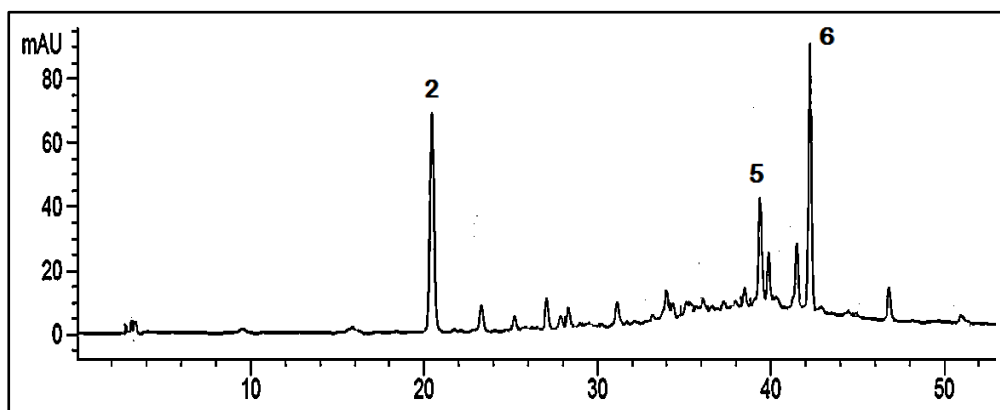


Figure 4.11 HPLC chromatogram of methanol fraction **KRM2** at 254 nm

The crude *n*-butanol and subfraction **KRB2**, **KRB4**, **KRB6** and **KRB7** was analysed by HPLC using an optimized acetonitrile/H₂O + 0.1% formic acid gradient ranging from 39% acetonitrile to 65% in 45 min at a flow rate of 1 mL/min (Fig. 4.12). Compound isolation was performed by means of semi-preparative HPLC-DAD-MS system using the same solvent gradient as HPLC at a flow rate of 3 mL/min. Compounds **3** (3.2 mg), **5** (3.2 mg) and **7** (3.8 mg) were finally isolated.

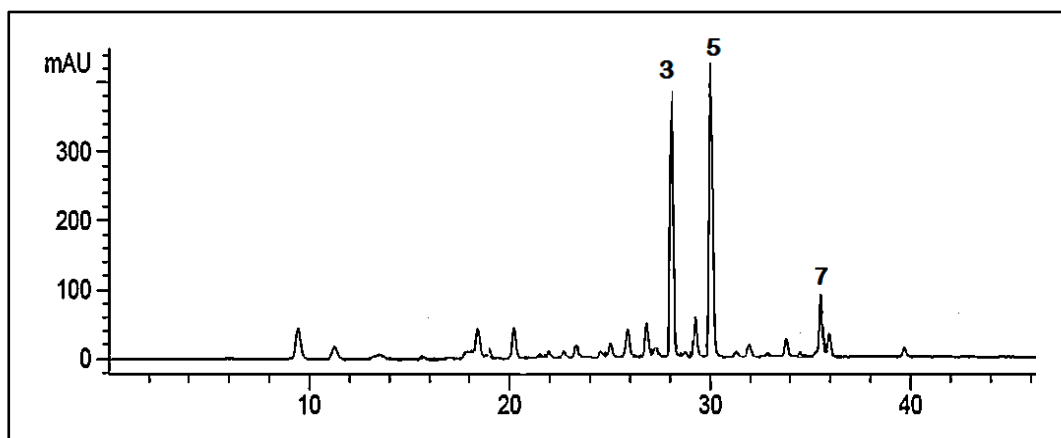


Figure 4.12 HPLC chromatogram of *n*-butanol fraction **KRB2** at 254 nm

4.3. STRUCTURE ELUCIDATION

Structures of the isolated compounds from *Kickxia ramosissima* were elucidated using ^1H - and ^{13}C - NMR (including DEPT-135 and DEPT-90) as well as 2D-NMR (COSY, HSQC and HMBC) spectroscopy. The molecular ion was derived from the high resolution mass spectra obtained with an Agilent™ 6530 quadrupole-time-of-flight mass spectrometer (QTOF-MS) and from the semi-preparative HPLC-DAD-MS system.

4.3.1 Iridoids (1-4)

4.3.1.1 Kickxiasine (1)

The structure of compound **1** was elucidated by ^1H -NMR, ^{13}C -NMR (including DEPT-135 and DEPT-90) and 2D-NMR (COSY, HSQC and HMBC). Since this compound had not been reported before the name **kickxiasine** was adopted (Fig. 4.13).

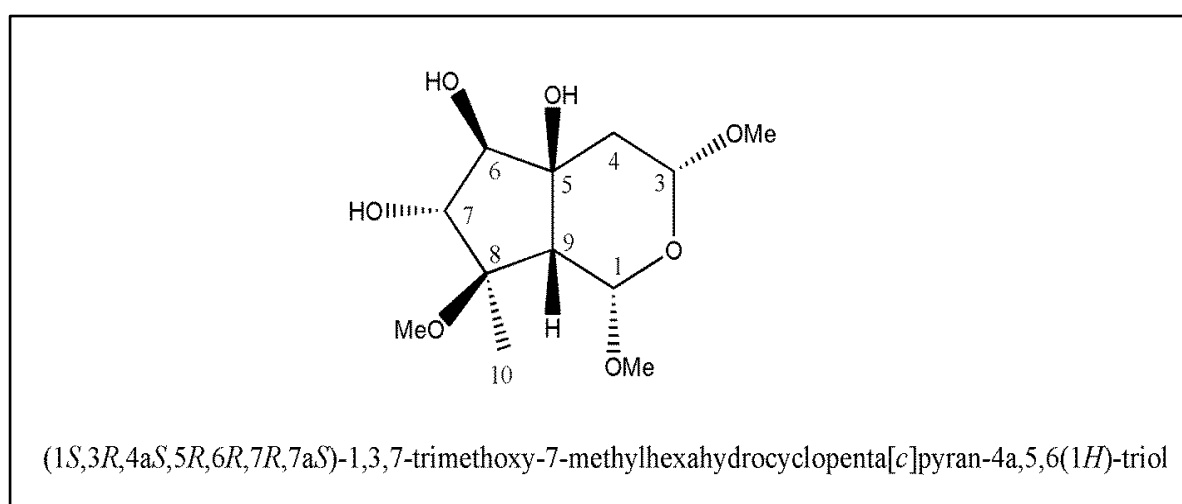


Figure 4.13 Structure of kickxiasine (**1**)

Library search on the ^{13}C -NMR data (Table 4.2 and Fig 4.14) using the NMR Predict database and ^1H -NMR (Table 4.2, Fig 4.15 and Fig 4.16) suggested that this compound has a similarity with 5-deoxy-8-deacetyl-3-*O*-methyl-6-*O*-*p*-hydroxybenzoyl-1,3-di-*epi*-clandonensine (Awale *et al.*, 2005) with some structural modifications on the cyclopentanopyran ring system. By the analysis of COSY (Fig 4.17), HSQC (Fig 4.18) and HMBC (Fig 4.19) correlations the structure was elucidated as shown.

The $^1\text{H-NMR}$ spectrum (Fig 4.15 and Fig 4.16) showed signals due to one methyl, one methylene, six methines and three methoxyls. The methylene signals at 1.51 ppm (dd) and 2.13 ppm (m) were assigned to H-4. A methine at 4.88 ppm (dd, $J=5.3, 8.4$ Hz) was attributed to H-3. A methine signal at 2.05 ppm (d, H-9) was coupled with another methine at 4.94 ppm (d, H-1), an upfield signal that was indicative of substitution with oxygen. Finally two methine protons showed coupled doublets at 3.76 ppm ($J=4.3$ Hz) and 3.92 ppm ($J= 4.3$ Hz), assigned to H-7 and H-6, respectively. The methyl group at 1.35 ppm (s, H-10) was attached to an oxygen-bearing quaternary carbon as determined by HMBC correlations.

The $^{13}\text{C-NMR}$ spectrum showed 12 carbon signals. The signals at 98.7 ppm and 98.8 ppm were assigned to carbons C-1 and C-3, respectively. In the cyclopentane ring the signals at 75.9 ppm and 77.3 ppm were attributed to C-6 and C-7, respectively. In the same ring the signals at 83.2 ppm, 77.4 ppm and 60.3 ppm were assigned to C-8, C-5 and C-9, respectively. The methyl group (C-10) showed a resonance signal at 17.1 ppm. Three methoxyl groups in position 1, 3 and 8 showed signals at 50.3, 55.6, and 56.3 ppm, respectively. The protons of these methoxyl groups were correlated in the HMBC spectrum with C-1, C-3 and C-8. The methyl group (C-10) at 1.35 ppm was attached to C-8, which was also established by HMBC correlations of the methyl protons with C-8 and C-9.

HMBC correlations from C-1 to the methoxyl group attached to C-8 and to H-9, and from C-6 to H-10 confirmed that the cyclopentane ring was substituted by a methyl group (C-10) and a methoxyl group. Similarly in the second ring, the signals at 97.3 ppm, 97.4 ppm and 39.5 ppm were assigned to C-1, C-3 and C-4, respectively. Also the HMBC correlations from H-1 to the methoxyl group attached to C-3 and from C-8 to the methoxyl group attached to C-1 confirmed the position of these methoxyl groups at C-1 and C-3.

The relative configuration was assigned on the basis of chemical shifts and coupling constants reported for related compounds (Morota *et al.*, 2010; Ahmad *et al.*, 2006; Awale *et al.*, 2005), and by analogy with linarioside (compound **4**, see below). The $^{13}\text{C-}$

NMR chemical shifts and 2D NMR experiments support the structure of compound **1** as a new compound for which the name kickxiasine was adopted.

Yellow powder; $[\alpha]^{20}_D$ -19.072 (c = 0.0011, MeOH); UV (acetonitrile / H₂O) λ_{\max} 202, 209, 216 and 225 nm; HR-MS (positive ion mode): m/z 301.1274 [M+Na]⁺ (calculated 301.1258) consistent with a molecular formula C₁₂H₂₂O₇Na, or C₁₂H₂₂O₇ for kickxiasine.

Compound **1** (kickxiasine) is reported here for the first time from nature. Although a number of polymethoxylated iridoids have been reported in nature from different medicinal plants, for instance *Tabebuia avellanedae* (Awale *et al.*, 2005), *Rehmannia glutinosa* (Morota *et al.*, 2010; Liu *et al.*, 2014) and *Gonocaryum calleryanum* (Kaneko *et al.*, 1995), however, the possibility that compound **1** is an artefact formed during extraction and isolation cannot be excluded.

Table 4.2 ¹H and ¹³C NMR assignments for kickxiasine (**1**) recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1	4.94; d; <i>J</i> =7.3	98.7
3	4.88; dd; 8.4, 5.3	98.8
4_a	2.13; m	40.9
4_b	1.51; dd; <i>J</i> = 14.0, 8.4*	
5		77.4
6	3.93 ; d; <i>J</i> =4.3	75.9
7	3.77; d; <i>J</i> =4.3	77.3
8		83.2
9	2.05; d, <i>J</i> = 7.3	60.3
10	1.35; s	17.1
OMe (C-1)	3.47; d; <i>J</i> =6.7	50.3
OMe (C-3)	3.47; s	55.6
OMe (C-8)	3.36; s	56.3

* Multiplicity and coupling constants were calculated on a spectrum recorded in acetone-d₆ because of overlapping with the HDO peak in the spectrum recorded in MeOH-d₄.

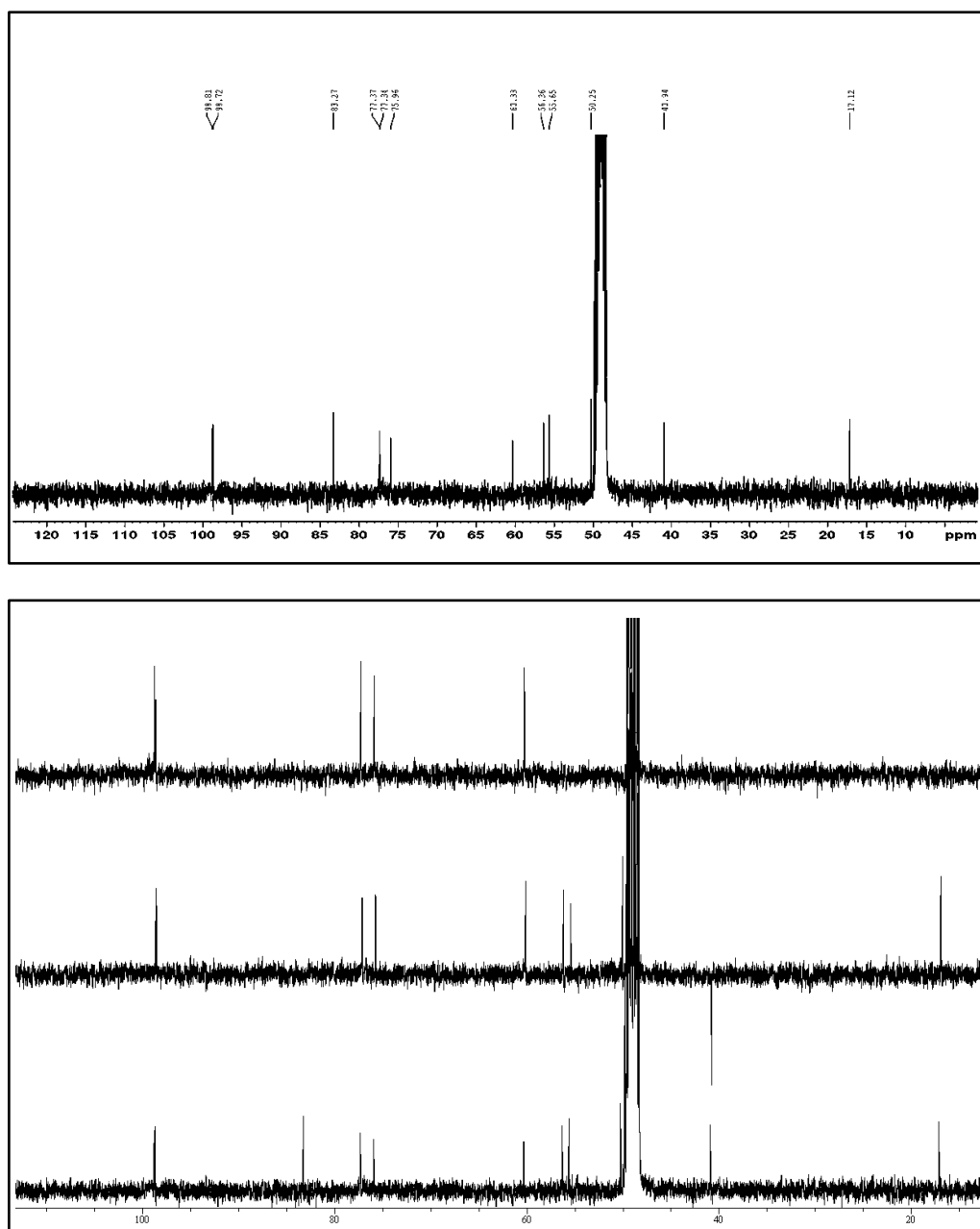


Figure 4.14 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of kickxiasine **1** in methanol- d_4 .

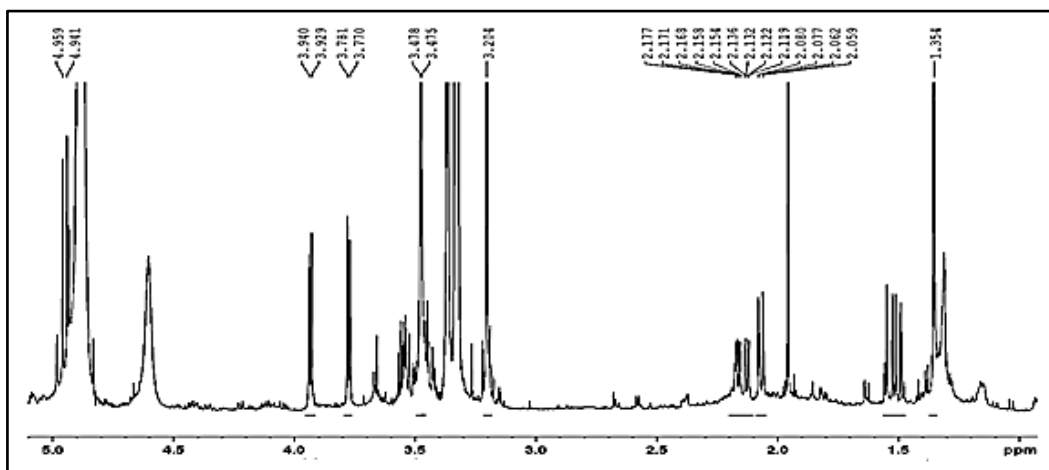


Figure 4.15 $^1\text{H-NMR}$ spectrum of kickxiasine **1** in methanol- d_4

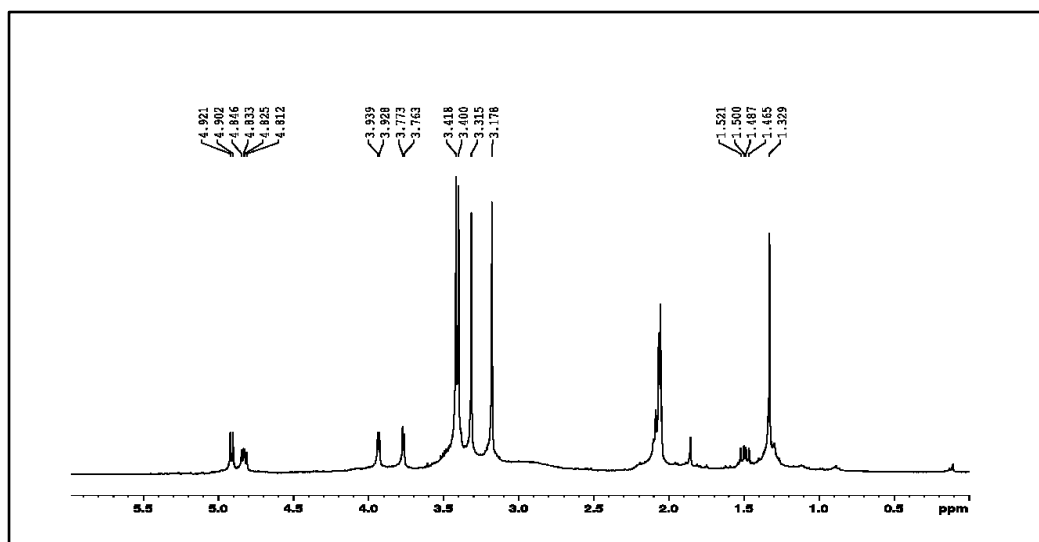


Figure 4.16 $^1\text{H-NMR}$ spectrum of kickxiasine **1** in acetone- d_6

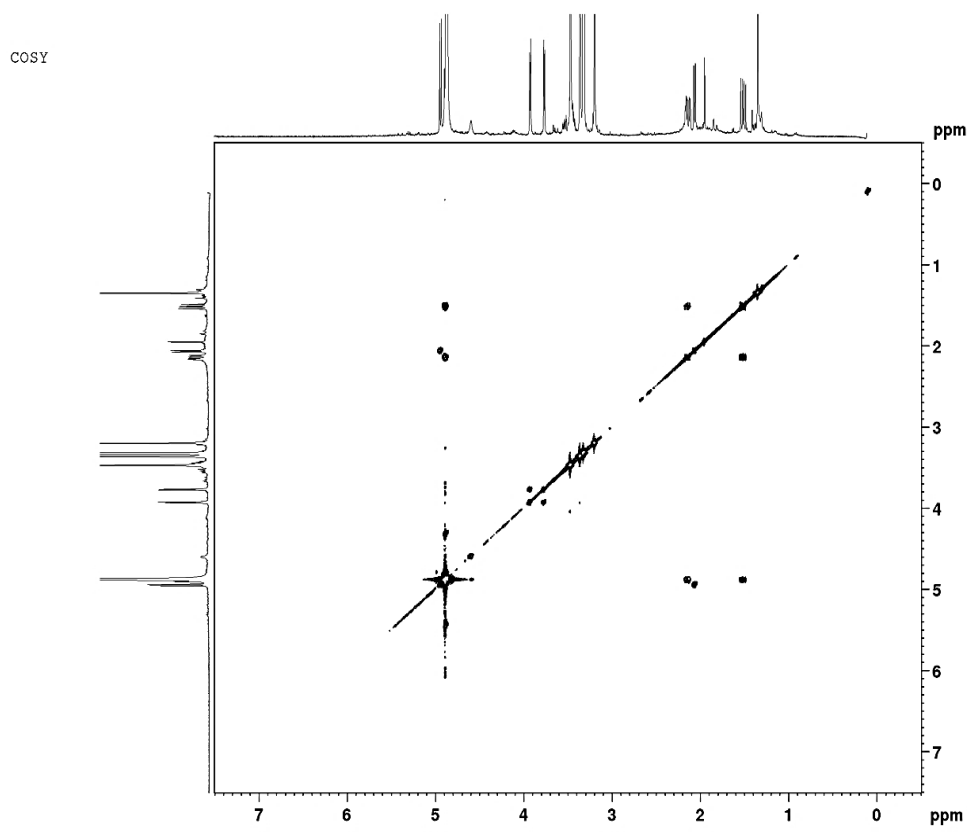


Figure 4.17 COSY spectrum of kickxiasine **1**

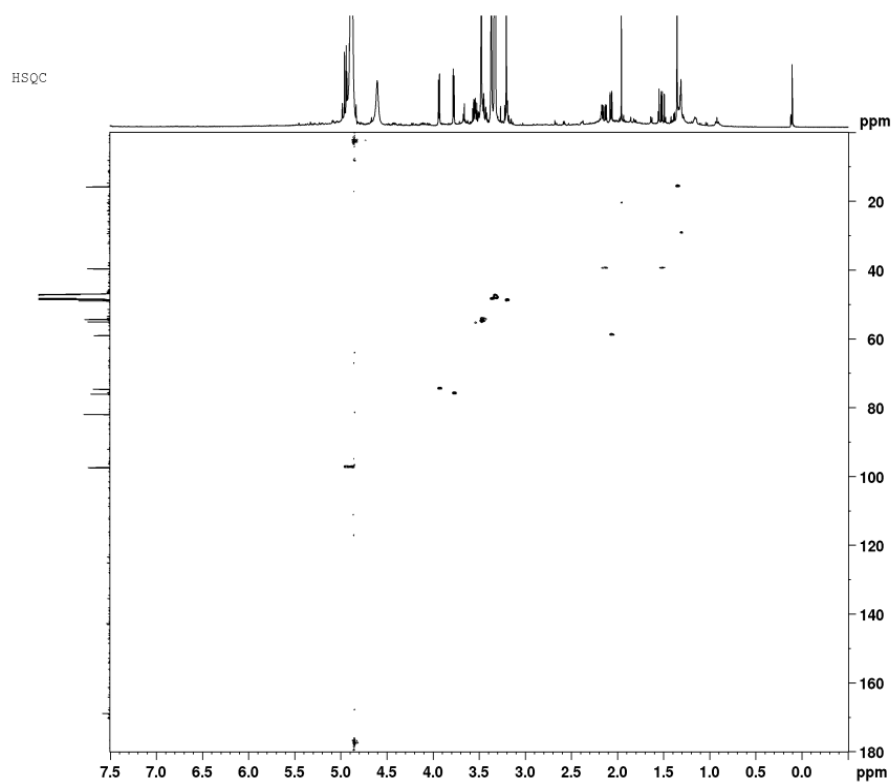


Figure 4.18 HSQC spectrum of kickxiasine **1**

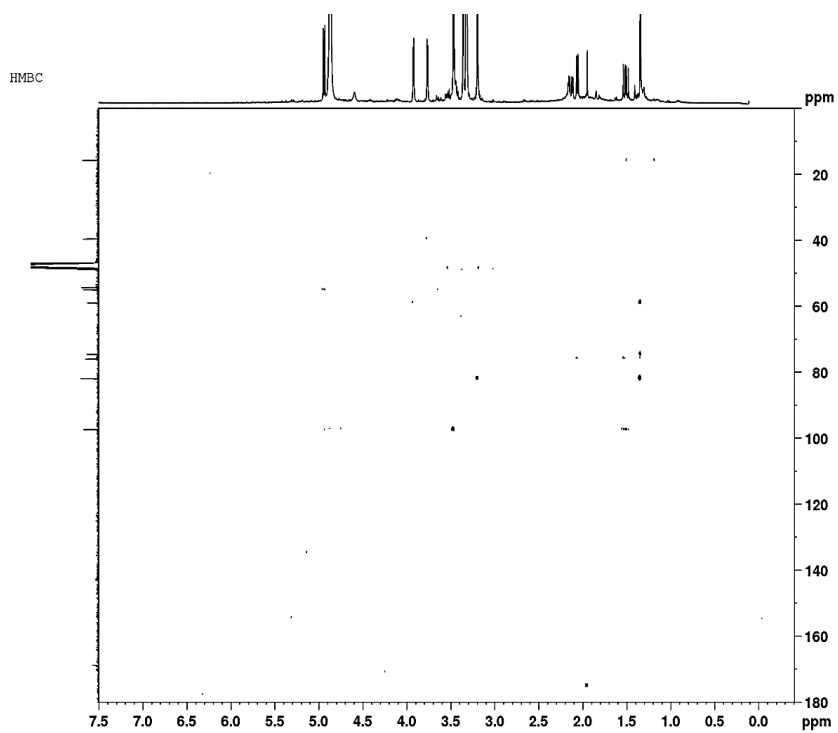


Figure 4.19 HMBC spectrum of kickxiasine **1**

4.3.1.2 Mussaenosidic acid (2)

The structure of compound **2** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$, DEPT-135 and DEPT-90 as mussaenosidic acid (Fig. 4.20).

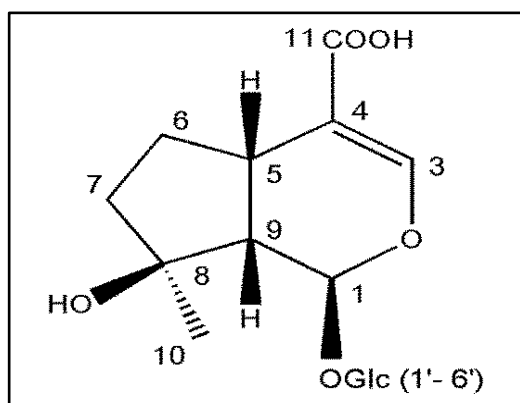


Figure 4.20 Structure of mussaenosidic acid (**2**)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.3 and Figure 4.21) using the NMR Predict database and $^1\text{H-NMR}$ (Fig 4.22) suggested that this compound was an iridoid glucoside. Comparison with NMR data from literature (Venditti *et al.*, 2016) suggested that compound **2** was mussaenosidic acid.

The $^1\text{H-NMR}$ spectrum (Fig. 4.22) of compound **2** displayed two methylene signals (multiplets) at 2.23 ppm (assigned to H-6) and at 1.74 / 1.52 ppm assigned to H-7a and H-7b, respectively. Another multiplet at 3.10 ppm was attributed to H-5. A singlet at 1.30 ppm was assignable to a methyl group attached to a hydroxyl bearing carbon. Similarly a multiplet appearing at 2.36 ppm was assigned to H-9 whereas a doublet at 5.60 ppm ($J=2.8$ Hz) was attributed to H-1. The anomeric proton H-1' resonated at 4.80 ppm (partly overlapping with the HDO signal). Finally the sugar protons were observed between 3.0-3.52 ppm (H-2'; H-3'; H-4'; H-5'). The methylene protons of the sugar moiety (H-6') resonated at 3.92 ppm (dd; $J=2.0, 12.0$ Hz, H-6a') and 3.72 ppm (dd; $J=6.0, 12.0$ Hz, H-6b').

The $^{13}\text{C-NMR}$ spectrum showed signals of 16 carbons and a β -glucopyranose moiety. The signals at 94.3 ppm and 29.5 ppm were assigned to C-1 and C-5, respectively. The

¹³C-NMR spectrum showed the presence of a double bond with the olefinic carbons at 151.0 ppm and 112.6 ppm that were assigned to C-3 and C-4, respectively. The resonance at 169.4 ppm was assigned to a carbonyl group (C-11). Two methylene signals were recorded at 28.7 ppm and 39.6 ppm, which were attributed to C-6 and C-7, respectively. Finally the signals at 79.6 ppm and 50.6 ppm were attributed to C-8 and C-9, respectively. The methyl signal appearing at 24.6 ppm was assigned to C-10.

Yellow powder; UV (acetonitrile / H₂O) λ_{max} 237 nm; ESI-MS (positive ion mode): *m/z* 399 [M+Na]⁺ consistent with a molecular formula C₁₆H₂₄O₁₀.

Mussaenosidic acid has been reported from various medicinal plants including *Pedicularis kernerii* (Venditti *et al.*, 2016), *Kickxia elatine* (L.) Dum., *Kickxia spuria* (L.) Dum. (Handjieva *et al.*, 1995), *Lagochilus ilicifolius* (Guangzhou *et al.*, 2012) and *Vitex negundo* (Sehgal *et al.*, 1982). Mussaenosidic acid (**2**) is reported here for the first time in *Kickxia ramosissima*.

Table 4.3 ¹H-NMR and ¹³C-NMR assignments for mussaenosidic acid (**2**) recorded in D₂O.

Position	δ_{H} (ppm), multiplicity, <i>J</i> (Hz)	δ_{C} (ppm)
1	5.6; d; <i>J</i> = 2.8	94.3
3	7.47; s	151.0
4		112.6
5	3.10; m	29.5
6	2.23 (2H; m)	28.7
7	1.74 (1H, m):1.52 (1H, m)	39.6
8		79.6
9	2.36; m	50.6
10	1.30; s	22.9
11		171.2
1'	4.80 (1H, overlapped with solvent)	98.2
2'	3.07-3.52; m	72.6
3'	3.07-3.52; m	77.2
4'	3.07-3.52; m	69.5
5'	3.07-3.52; m	76.2
6a'	3.92; dd; <i>J</i> =12.0; 2.0	60.6
6b'	3.72; dd; <i>J</i> =12.0; 6.0	

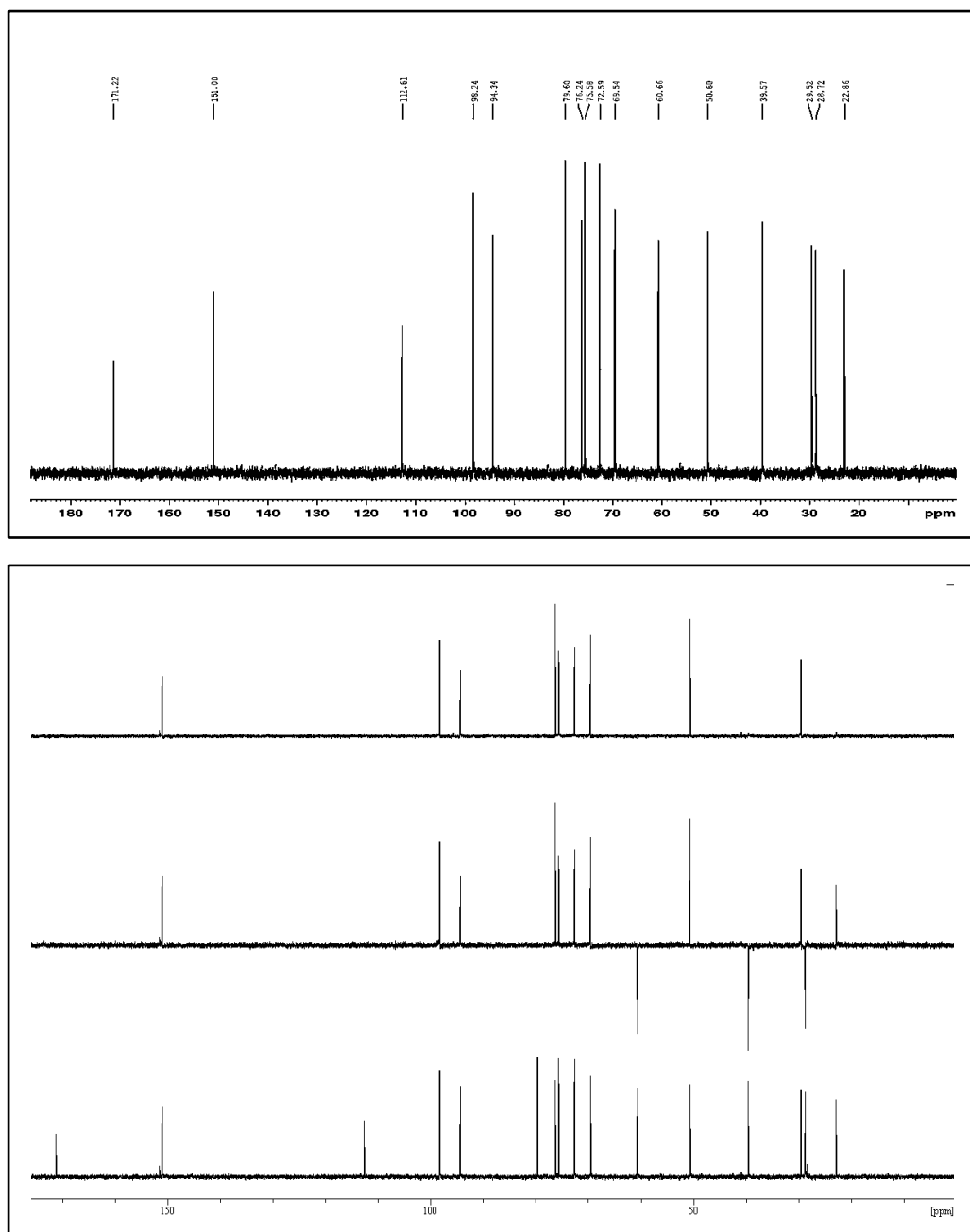


Figure 4.21 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of mussaenosidic acid (**2**) in D_2O .

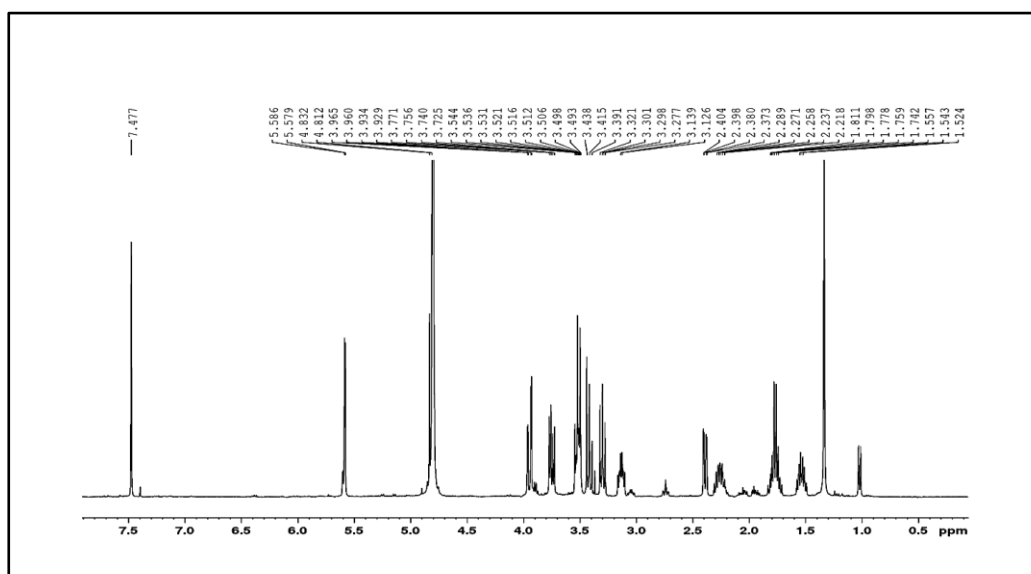


Figure 4.22 $^1\text{H-NMR}$ spectrum of mussaenosidic acid (**3**) in D_2O .

4.3.1.3 Mussaenoside (3):

The structure of compound **3** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$, DEPT-135 and DEPT-90 as mussaenoside (Fig. 4.23).

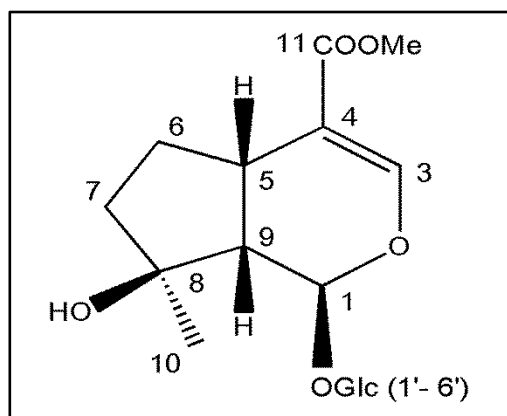


Figure 4.23 Structure of mussaenoside (**3**)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.4 and Fig 4.24) using the NMR Predict database and $^1\text{H-NMR}$ (Fig 4.25) suggested that this compound was an iridoid glucoside. Comparison with NMR data from literature (Ersoz *et al.*, 1998; Haznagy-Radni *et al.*, 2014) suggested that compound **2** was mussaenoside.

The $^1\text{H-NMR}$ (Fig 4.25) spectrum of compound **3** displayed two methylene signals (multiplets) at 2.30 ppm and 1.45 ppm assigned to H-6_a and H-6_b, respectively, and 1.75 ppm assigned to H-7_a and H-7_b. Another multiplet at 3.20 ppm was attributed to H-5. A singlet at 1.33 ppm was assignable to a methyl group attached to a hydroxyl bearing carbon, whereas a methoxyl group appeared at 3.71 ppm. Similarly a doublet of doublet appearing at 2.24 ppm ($J=4.2, 9.3$ Hz) was assigned to H-9 whereas a doublet at 5.48 ppm ($J=4.2$ Hz) was attributed to H-1. The anomeric proton H-1' resonated at 4.25 ppm (d; $J=7.9$ Hz). Finally sugar protons were observed at 3.0-3.72 ppm (H-2'; H-3'; H-4'; H-5'). The methylene protons of the glucose moiety (H-6'_a and H-6'_b) were observed at 3.92 ppm (dd; $J= 2.0, 11.8$ Hz) and 3.65 ppm (dd; $J= 6.2, 11.8$).

The ^{13}C -NMR spectrum showed signals of 17 carbons and a β -glucopyranose moiety. Two olefinic carbons were observed at 152.0 ppm and 113.3 ppm that were assigned to C-3 and C-4, respectively. The carbon resonances at 169.4 ppm (C-11) and 51.6 ppm confirmed the presence of the COOCH_3 ester group. The signals at 95.3 ppm and 32.0 ppm were assigned to carbons C-1 and C-5, respectively. The methylene signals at 30.7 ppm and 40.7 ppm were assigned to C-6 and C-7, respectively. Finally signals at 80.5 ppm and 52.3 ppm were attributed to C-8 and C-9, respectively. The methyl signal (C-10) appeared at 24.6 ppm.

Yellow amorphous solid; UV (acetonitrile / H_2O) λ_{max} 237 nm; ESI-MS (positive ion mode): m/z 413 $[\text{M}+\text{Na}]^+$ consistent with a molecular formula $\text{C}_{17}\text{H}_{26}\text{O}_{10}$.

Mussaenoside is a common iridoid and has been isolated from a number of plants species, for instance *Melampyrum* sp. (Damtoft *et al.*, 1984), *Bellardia trixago* (Ersoz *et al.*, 1998) and *Mussaenda incana* (Dinda *et al.*, 2005). This is the first report of its occurrence in *Kickxia ramosissima*. Since mussaenoside is the methyl ester of mussaenosidic acid (**2**), the possibility that it is an artefact formed during extraction procedures in which methanol is used cannot be excluded.

Table 4.4 ¹H-NMR and ¹³C-NMR assignments for mussaenoside (**3**) recorded in methanol-*d*₄.

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1	5.48; d; <i>J</i> =4.2	95.3
3	7.4; s	152.0
4		113.3
5	3.20; m	32.0
6	2.30(1H, m) / 1.45(1H, m)	30.7
7	1.75 (2H, m)	40.7
8		80.5
9	2.24; dd; <i>J</i> =9.3; 4.2	52.3
10	1.33; s	24.6
11		169.4
1'	4.25; d; <i>J</i> =7.8	99.8
2'	3.0-3.72; m	74.7
3'	3.0-3.72; m	78.4
4'	3.0-3.72; m	71.7
5'	3.0-3.72; m	78.0
6_a'	3.92; dd; <i>J</i> = 11.8; 2.0	62.9
6_b'	3.65; dd; <i>J</i> = 11.8; 6.2	
OMe	3.71; s	51.6

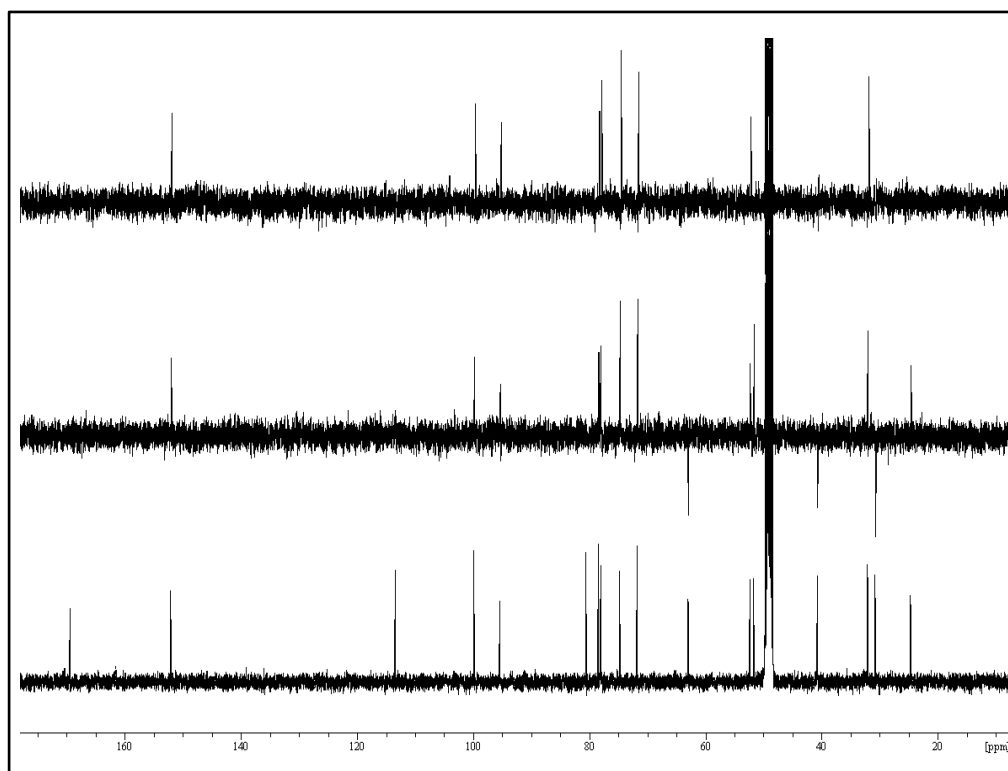
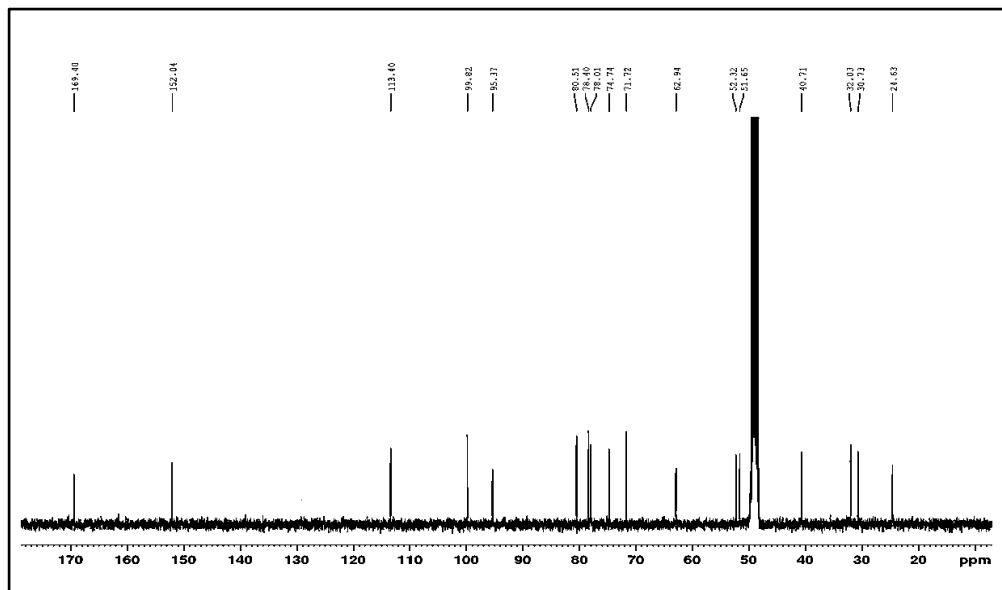


Figure 4.24 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of mussaenoside(3) in methanol- d_4 .

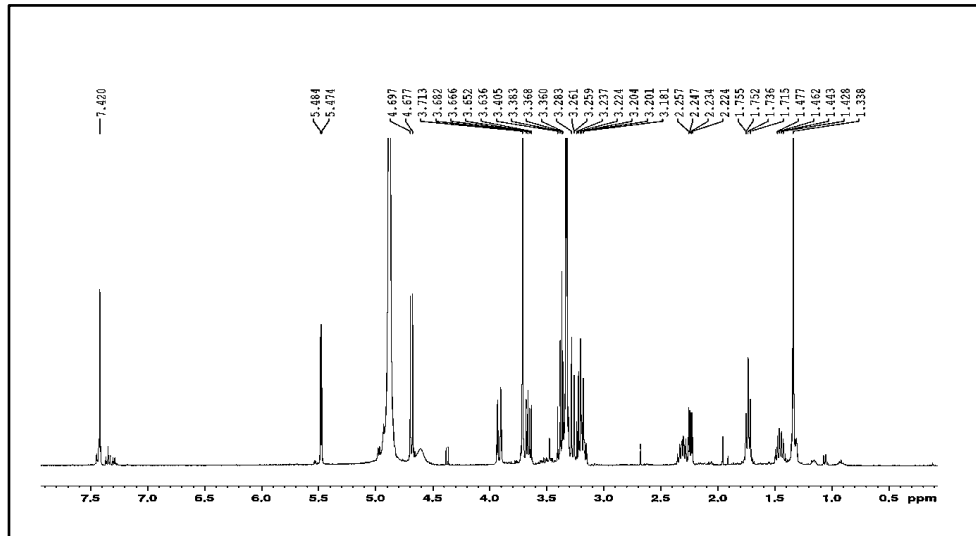


Figure 4.25 ¹H-NMR spectrum of mussaenoside(3) in methanol-d₄.

4.3.1.4 Linarioside (4):

The structure of compound **4** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$, DEPT-135 and DEPT-90 as linarioside (Fig. 4.26).

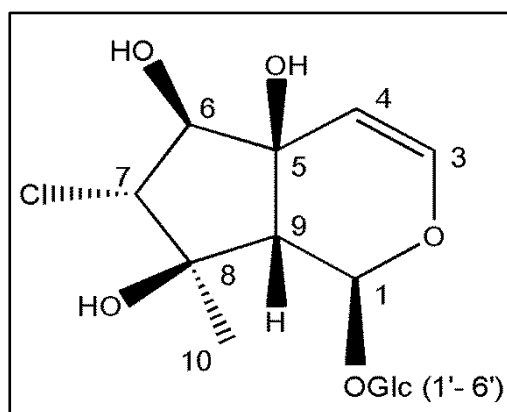


Figure 4.26 Structure of linarioside (**4**)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.5 and Fig 4.27) using the NMR Predict database and $^1\text{H-NMR}$ (Fig 4.28) suggested that this compound was an iridoid glucoside. Comparison with spectral data obtained from literature (Kitagawa *et al.*,1973; Otsuka, 1993) suggested compound **4** was linarioside.

The signals at 6.42 ppm and 5.30 ppm were assigned to the olefinic protons H-3 and H-4 that were coupled with each other. The singlets appearing at 5.80 ppm and 2.57 ppm were attributed to H-1 and H-9, respectively. These singlets were indicative of the absence of coupling (i.e. very small J) between H-1 and H-9. Multiplets at 3.42-4.2 ppm represented H-6 and H-7, respectively. The methyl signal was recorded at 1.30 ppm and assigned to H-10. The anomeric proton (H-1') showed a resonance at 4.20 ppm (d; $J= 7.8$ Hz). Finally the sugar protons were observed at 3.42-4.2 ppm (H-2'; H-3'; H-4'; H-5' and H-6').

The $^{13}\text{C-NMR}$ spectrum showed signals of 15 carbons and a β -glucopyranose moiety. The signals at 91.0 ppm, 139.2 ppm and 108.1ppm were assigned to C-1, C-3 and C-4, respectively. Similarly C-5 and C-6 were assigned to the signals at 64.6 ppm and 79.2

ppm, respectively. The methyl group appearing at 17.1 ppm was attributed to C-10. Finally the signals at 72.5 ppm and 74.0 ppm were assigned to C-7 and C-8, respectively.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 210 nm; ESI-MS (positive ion mode): m/z 422 [M+Na]⁺ consistent with a molecular formula C₁₅H₂₃O₁₀ Cl.

Linarioside was the first example of a chlorine containing iridoid glucoside in nature isolated from *Linaria japonica* that was reported by Kitagawa *et al.* (1973). A detailed investigation proved that the compound linarioside is a naturally occurring compound and not formed as an artefact during the extraction process. Afterwards a number of other medicinal plants were reported as having linarioside as one of their constituents. These include *Cymbalaria muralis* (Kapoor and Reisch, 1974), *Asystasia Bella* (Demuth *et al.*, 1989), *Linaria aegyptiaca* (Ferhat *et al.*, 2010) and *Linaria genistifolia* (Ilieva *et al.*, 1992). We are reporting linarioside for the first time in *Kickxia ramosissima*.

Table 4.5 ¹H-NMR and ¹³C-NMR assignments for linarioside(4) recorded in D₂O.

Position	δ_H (ppm); multiplicity; <i>J</i> (Hz)	δ_C (ppm)
1	5.80; s	91.0
3	6.42; d; <i>J</i> =6.5	139.2
4	5.30; dd; <i>J</i> =1.2; 6.5	108.1
5		64.6
6	3.42-4.2; m	79.2
7	3.42-4.2; m	72.5
8		74.0
9	2.57; s	56.2
10	1.30; s	17.1
1'	4.20; d; <i>J</i> = 7.8	97.8
2'	3.42-4.2; m	75.2
3'	3.42-4.2; m	79.2
4'	3.42-4.2; m	69.7
5'	3.42-4.2; m	76.2
6'	3.42-4.2; m	61.6

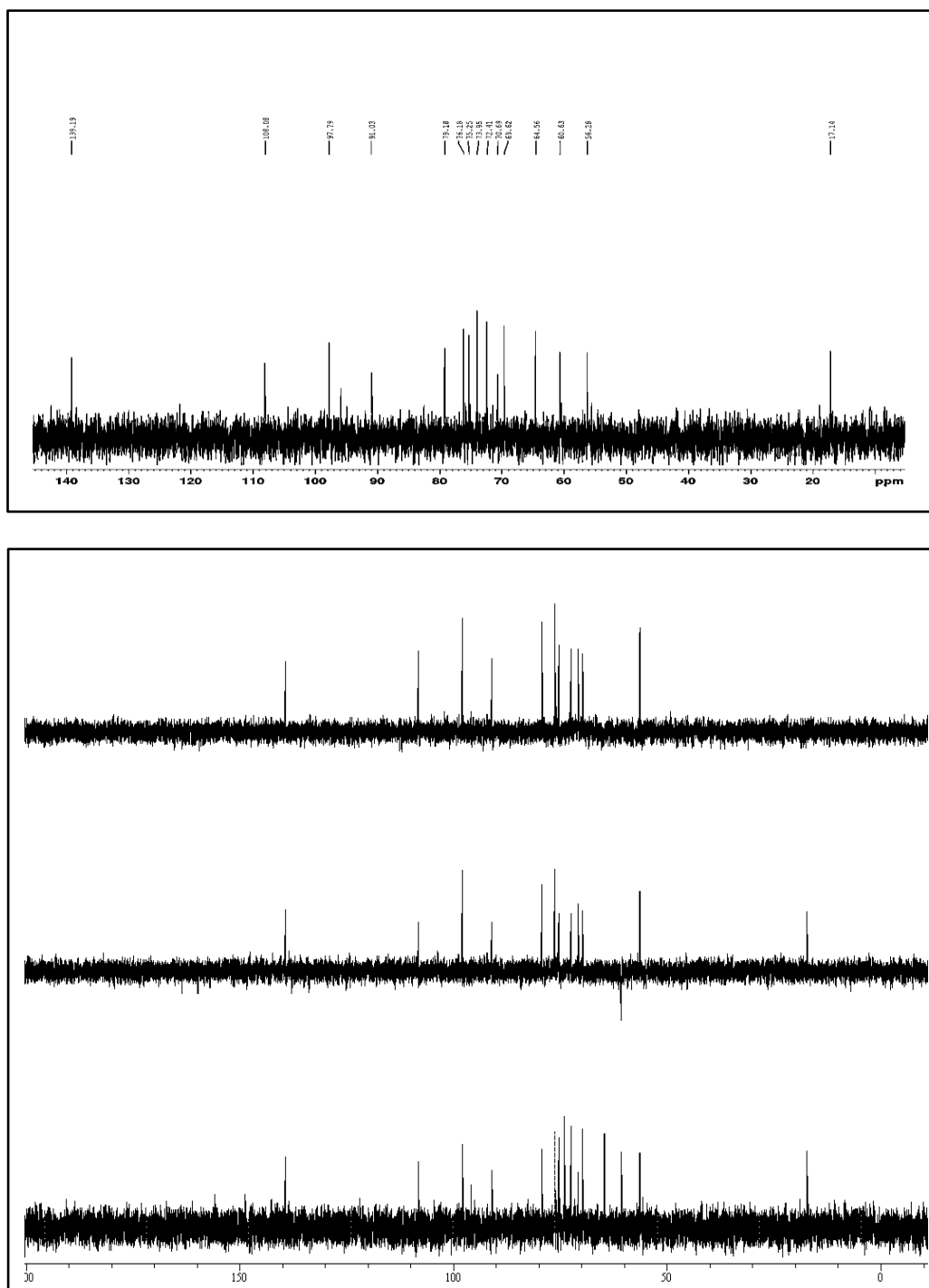


Figure 4.27 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of linarioside(4) in D_2O .

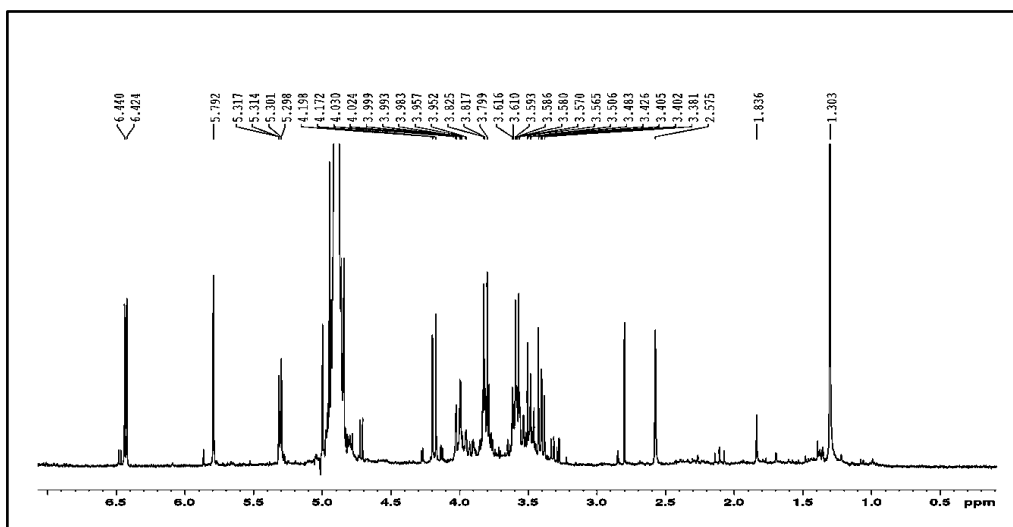


Figure 4.28 ¹H-NMR spectrum of linarioside(4) in D₂O.

4.3.2 Flavonoids

4.3.2.1 Pectolarigenin (5):

The structure of compound **5** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$, DEPT-135 and DEPT-90 as pectolarigenin (Fig. 4.29).

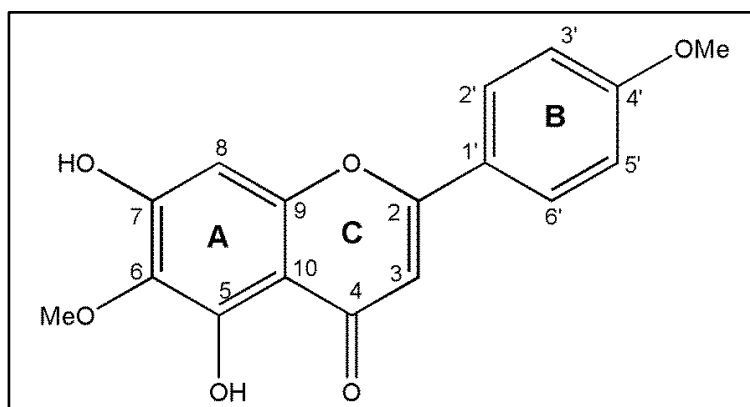


Figure 4.29 Structure of pectolarigenin (**5**)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.6 and Fig 4.30) suggested this compound is a flavonoid with a glucosyl moiety. By comparison with published data (Lim *et al.*, 2008), compound **5** was identified as pectolarigenin.

The $^1\text{H-NMR}$ (Table 4.6 and Fig 4.31) and $^{13}\text{C-NMR}$ (Table 4.6 and Fig 3.30) spectra of compound **5** exhibited resonances due to aromatic ring systems, typical for a flavonoid. In the $^1\text{H-NMR}$ spectrum typically 4 protons (H-2', H-3', H-5' and H-6') appearing as 2 doublets ($J=9.0$ Hz each) at 7.77 ppm and 6.97 ppm suggested an A_2B_2 pattern of ring B. Two singlets at 6.53 ppm and 6.51 ppm were assigned to H-3 and H-8, respectively. The typical pattern of singlets suggested the presence of a 5, 6, 7-oxygenated flavone ring. Similarly the proton spectrum exhibited 2 methoxyl signals at 3.84 ppm and 3.94 ppm that were assigned to the substituents at C-6 and C-4', respectively.

The $^{13}\text{C-NMR}$ spectrum of compound **5** showed the presence of 17 unsaturated carbon signals. The signals at 61.0 ppm and 55.6 were assigned to the methoxyl groups at C-6 and C-4', respectively. The carbonyl group (C-4) showed a resonance at 183.0 ppm, indicating the absence of a hydroxyl group at C-3 and a free hydroxyl group at C-5. A

hydrogen bond between the latter hydroxyl and the carbonyl group leads to relative deshielding of C-4. In ring C the chemical shifts at 164.4 ppm, 103.9 ppm and 152.2 were assigned to C-2, C-3 and C-5 respectively. The signals at 130.4 ppm and 155.0 ppm were attributed to C-6 and C-7, respectively, whereas in ring A the resonances at 93.4 ppm, 153.1 ppm and 105.8 ppm were assigned to C-8, C-9 and C-10, respectively. Finally in ring B the signals at 123.6 ppm, 128.2 ppm and 114.5 ppm were attributed to C-1', C-2'/C-6' and C-3'/C-5', respectively, and the signal at 162.8 ppm to C-4'.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 275, 335 nm; ESI-MS (positive ion mode): m/z 315 [M+H]⁺ consistent with a molecular formula C₁₇H₁₄O₆.

Pectolinarigenin is a common flavonoid that has been reported from various medicinal plants including *Linaria reflexa* (Cheriet *et al.*, 2014) *Cirsium chanroenicum* (Lim *et al.*, 2008), and it has also been reported before from *Kickxia ramosissima* (Singh and Parakash, 1987).

Table 4.6 ¹H-NMR and ¹³C-NMR assignments for pectolinarin (**5**) recorded in CDCl₃.

Position	δ_{H} (ppm), multiplicity, <i>J</i> (Hz)	δ_{C} (ppm)
1		
2		164.4
3	6.53; s	103.9
4		183.0
5		152.2
6		130.4
7		155.0
8	6.51; s	93.4
9		153.1
10		105.8
1'		123.6
2'	7.77; d; <i>J</i> =9.0	128.2
3'	6.97; d; <i>J</i> =9.0	114.5
4'		162.8
5'	6.97; d; <i>J</i> =9.0	114.5
6'	7.77; d; <i>J</i> =9.0	128.2
OMe (C-6)	3.98; s	61.0
OMe (C-4')	3.84; s	55.6

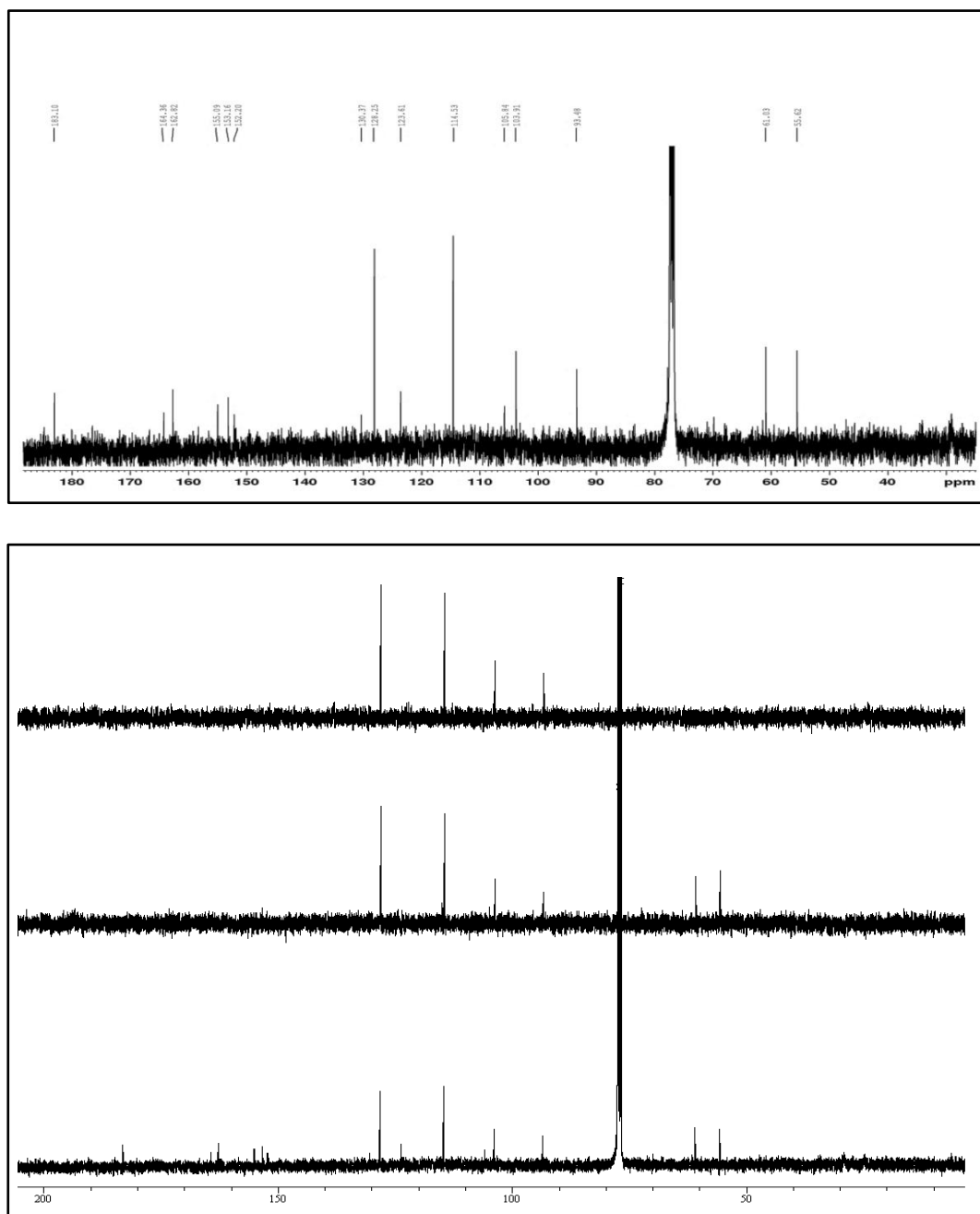


Figure 4.30 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of pectolinarigenin (**5**) in CDCl_3 .

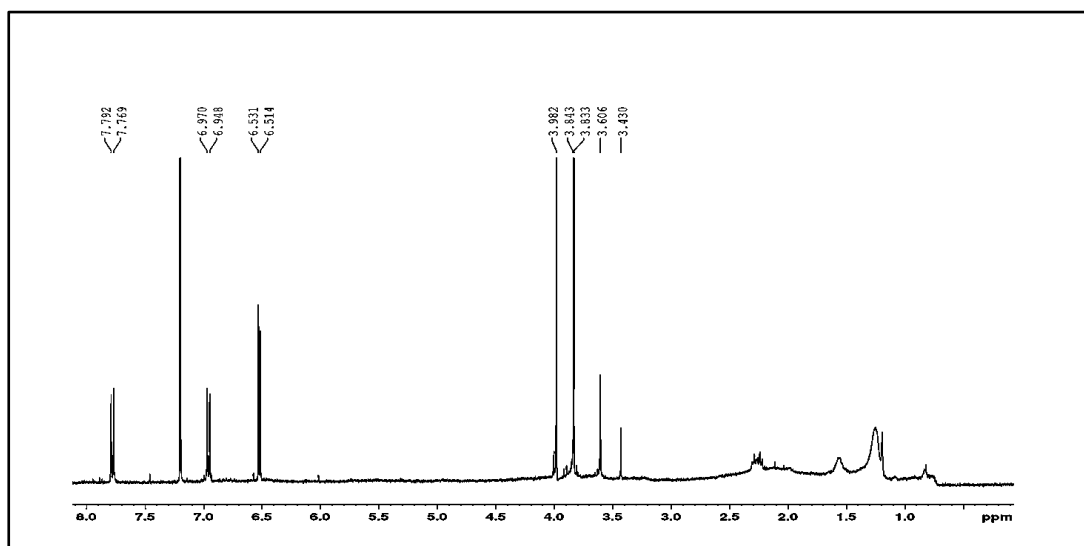


Figure 4.31 ¹H-NMR spectrum of pectolarigenin (5) in CDCl₃.

4.3.2.2 Pectolinarin (6):

The structure of compound **6** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$, DEPT-135 and DEPT-90 as pectolinarin (Fig. 4.32).

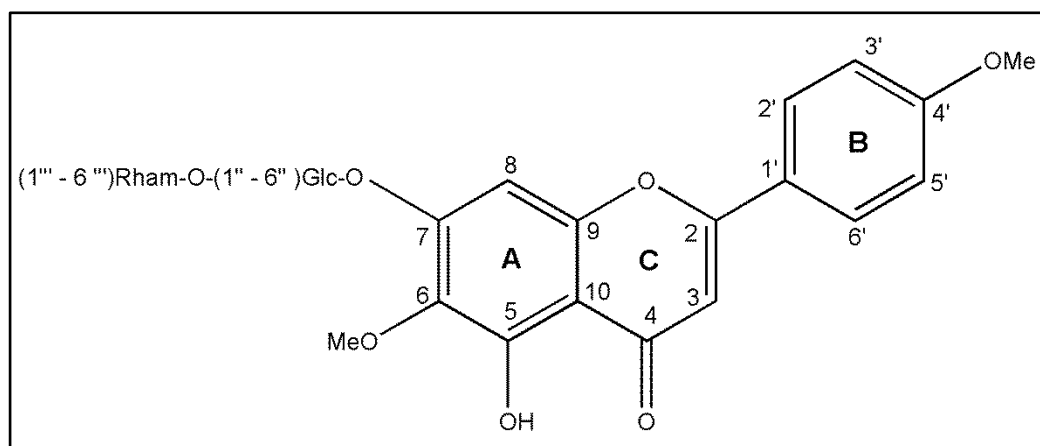


Figure 4.32 Structure of pectolinarin(6)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.7 and Fig 4.33) suggested this compound is a flavonoid with a rutinosyl moiety. Its ^1H , $^{13}\text{C-NMR}$ and mass spectral data were compared with published data (Lim *et al.*, 2008), allowing its complete identification.

The $^1\text{H-NMR}$ (Table 4.6 and Fig 4.34) and $^{13}\text{C-NMR}$ (Table 4.6 and Fig 4.33) spectra of compound **9** exhibited resonances due to aromatic ring systems, typical for a flavonoid moiety and sugar residues. In the $^1\text{H-NMR}$ spectrum the aromatic region exhibited an AA'XX' system of spins at 8.04 ppm (2H each; d; $J= 8.9$ Hz) attributed to H-2' and H-6', and 7.16 ppm (2H each, d; $J=8.9$ Hz) assigned to H-3' and H-5', respectively, indicating a 4'-oxygenated B ring.

The doublets at 4.56 ppm ($J = 1.1$ Hz, H-1''') and 5.12 ppm ($J = 7.0$ Hz, H-1'') are typical of anomeric hydrogens. According to the coupling constants, the configurations of the D-glucosyl- and L-rhamnosyl moieties were determined as β ($J= 7.0$ Hz) and α ($J= 1.1$ Hz), respectively. The resonances in the region of 3.12 -3.75 ppm (m, H-2'', H-3'', H-4'', H-5'', H-6'' and H-2''', H-3''', H-4''', H-5''') can be assigned to the remaining sugar protons. The doublet at 1.06 ppm (3H, $J=6.0$ Hz, H-6''') is related to the rhamnose methyl group. The singlets at 6.92 ppm and 6.94 ppm were assigned to H-8 and H-3, respectively. Two

methoxyl signals at 3.85 ppm (s) and 3.76 ppm(s) were assigned to C-4' and C-6, respectively.

The ^{13}C -NMR spectrum of compound **6** showed the presence of 29 carbon signals. Nine signals were observed in the range of 66.4 - 76.9 ppm, corresponding to two sugar units i.e glucose and rhamnose. The ^{13}C -NMR signals at 56.0 ppm and 60.8 ppm were assigned to the methoxyl groups at C-4' and C-6, respectively. The methyl group of rhamnose was observed at 18.2 ppm.

The chemical shifts of the carbon signals for D-glucose and L-rhamnose, indicated L-rhamnose as the terminal sugar, linked to C-6" of glucose (66.0 ppm), i.e. rutinose. The DEPT experiment confirmed that this signal was a methylene group.

Yellow powder; . UV (acetonitrile / H₂O) λ_{max} 275, 330 nm; ESI-MS (positive ion mode): m/z 623 [M+Na]⁺ consistent with a molecular formula C₂₉H₃₄O₁₅.

Pectolarine has already been isolated from a number of plants species including *Cirsium japonicum* (Liu *et al.*, 2007), *Cirsium chanroenicum* (Lim *et al.*, 2008) and *Kickxia abhaica* (Al-Rehaily *et al.*, 2006). It has also been reported from *Kickxia ramosissima* before (Ahmad *et al.*, 2006).

Table 4.7 ¹H-NMR and ¹³C-NMR assignments for pectolinarin (**6**) recorded in DMSO-*d*₆.

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1		
2		164.6
3	6.92; s	103.9
4		182.8
5		152.6
6		133.3
7		157.0
8	6.94; s	94.8
9		152.7
10		106.3
1'		123.2
2'	8.04; d; <i>J</i> =8.9	129.0
3'	7.16; d; <i>J</i> = 8.9	115.3
4'		163.0
5'	7.16; d; <i>J</i> = 8.9	115.2
6'	8.04; d; <i>J</i> =8.9	128.9
1''	5.12; d; <i>J</i> =7.0	100.9
2''	3.12-3.75; m	73.6
3''	3.12-3.75; m	76.9
4''	3.12-3.75; m	69.9
5''	3.12-3.75; m	76.2
6''	3.12-3.75; m	66.4
1'''	4.56; d; 1.1	100.7
2'''	3.12-3.75; m	70.9
3'''	3.12-3.75; m	71.5
4'''	3.12-3.75; m	72.4
5'''	3.12-3.75; m	68.2
6'''	1.06; d; 6.0	18.2
OMe(C-6)	3.76; s	60.8
OMe(C-4')	3.85; s	56.0

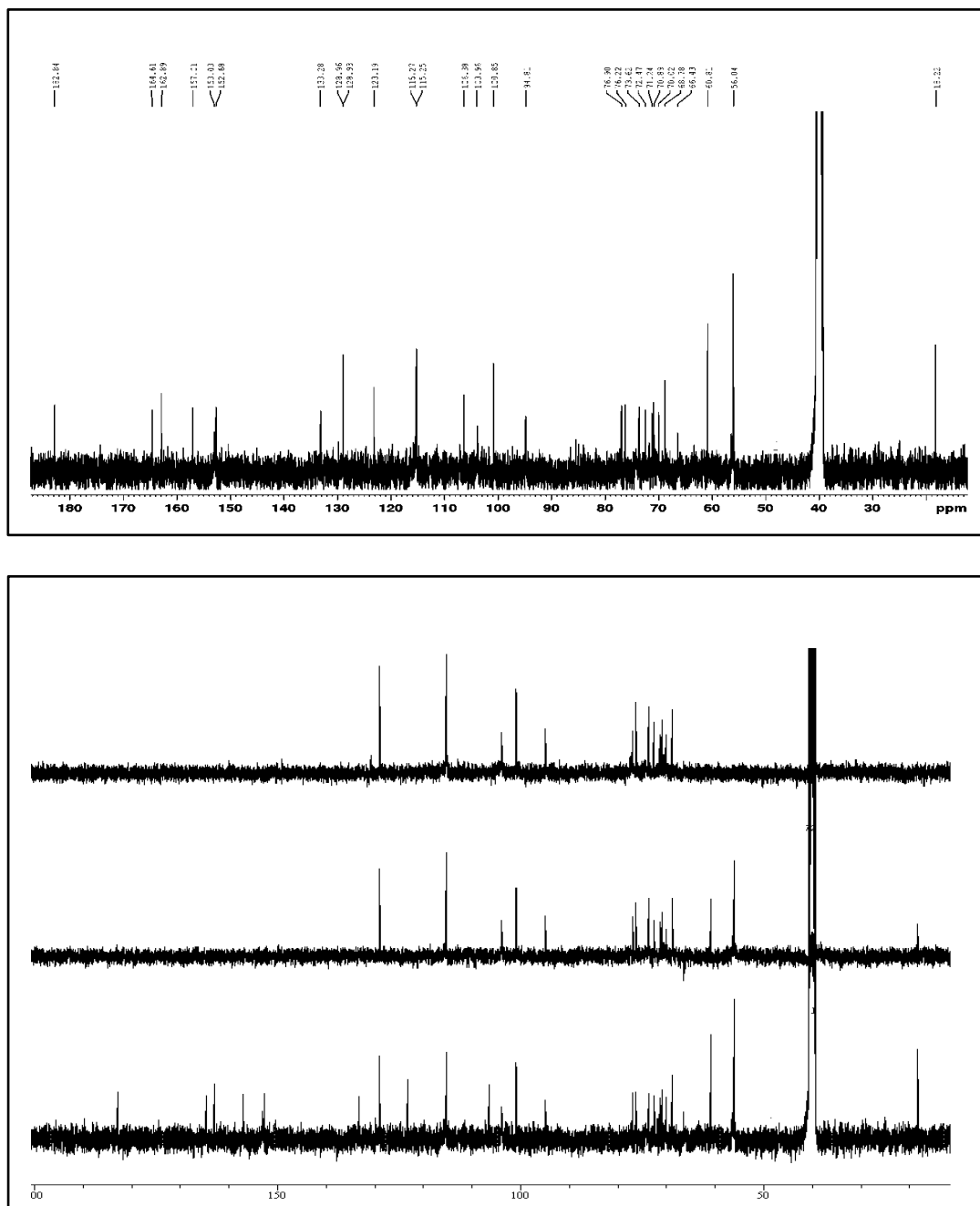


Figure 4.33 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of pectolarin (**6**) in DMSO-d_6 .

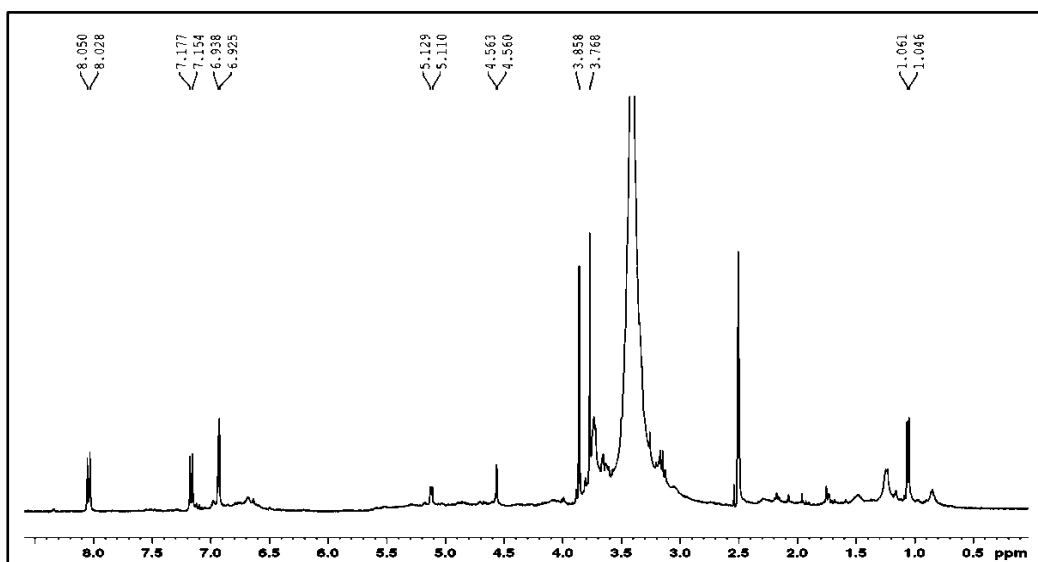


Figure 4.34 $^1\text{H-NMR}$ spectrum of pectolinarin (6) in DMSO-d_6 .

4.3.3 Benzoic acid derivatives

4.3.3.1 4-Hydroxy benzoic acid methyl ester (7):

The structure of compound **7** was elucidated by ^1H -, ^{13}C -NMR, DEPT-135 and DEPT-90 as 4-hydroxybenzoic acid methyl ester (**7**) (Fig 4.35).

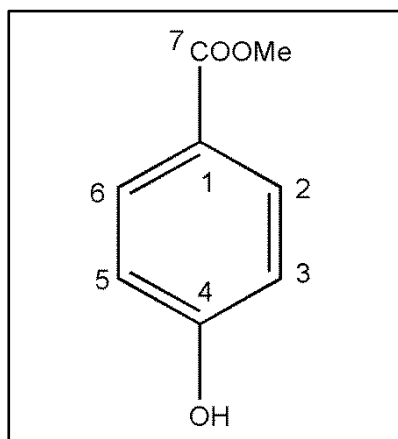


Figure 4.35. 4-hydroxy benzoic acid methyl ester (**7**)

Library search on the ^{13}C -NMR data (Table 4.8 and Fig 4.36) suggested this compound is a benzoic acid derivative. Based on its ^1H , ^{13}C -NMR and mass spectral data compound **7** was found to be 4-hydroxy benzoic acid methyl ester (Yoshioka *et al.*, 2004).

The ^1H -NMR (Table 4.8 and Fig 4.37) and ^{13}C -NMR (Table 4.8 and Fig 4.36) spectra of compound **7** are indicative of an aromatic ring system. The ^1H -NMR spectrum revealed a doublet at 7.88 ppm ($J=8.9$ Hz) that were assigned to H-2 and H-6, respectively. Similarly a doublet observed at 6.84 ppm ($J= 8.9$) was assigned to H-3 and H-5. Finally a methoxyl signal was observed at 3.92 ppm (s).

The ^{13}C -NMR spectrum indicated the presence of 8 carbons. The carbonyl group of the ester moiety presented a resonance at 168.6 ppm. The signal at 121.8 ppm and 132.5 ppm were assigned to C-1 and C-2/C-6, whereas the signals 116.0 and 163.6 ppm were attributed to C-3/C-5 and C-4, respectively. The methoxyl group appeared at 52.0 ppm.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 226, 287, 338 nm; ESI-MS (positive ion mode): m/z 153 [M+H]⁺ consistent with a molecular formula C₈H₈O₃.

4-Hydroxy-benzoic acid methyl ester has been found in many medicinal plant species for instance *Vitex rotundifolia* (Yoshioka *et al.*, 2004) and *Houttuynia cordata* (Jong and Jean, 1993). This is, however, the first report of its presence in *Kickxia ramosissima*.

Table 4.8 ¹H-NMR and ¹³C-NMR assignments for 4- hydroxy benzoic acid methyl ester (7) recorded in methanol-*d*₄.

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
1		121.8
2	7.88; d; <i>J</i> = 8.9	132.5
3	6.84; d; <i>J</i> = 8.9	116.0
4		163.6
5	6.84; d; <i>J</i> = 8.9	116.0
6	7.88; d; <i>J</i> = 8.9	132.5
7		168.6
OMe	3.86; s	52.0

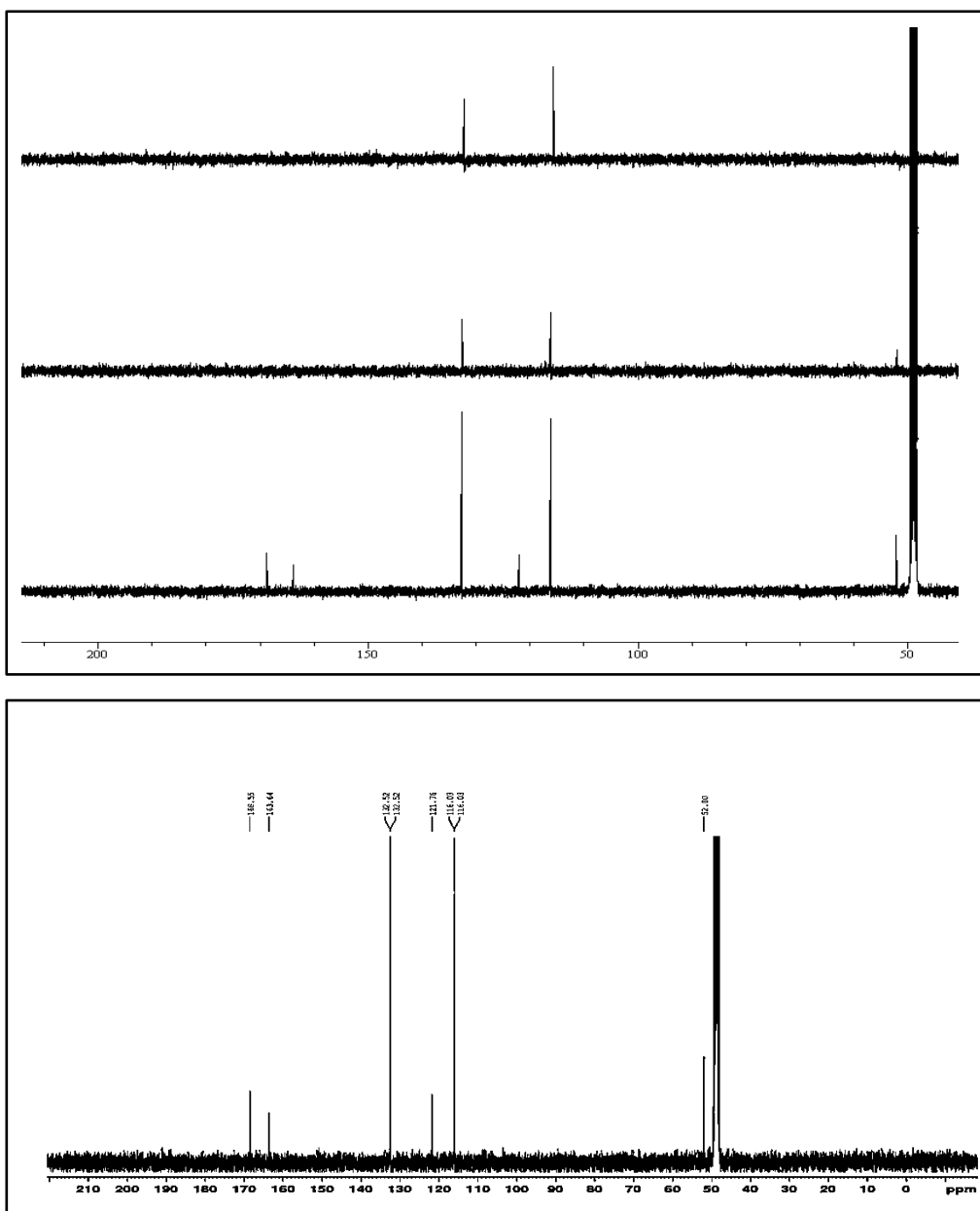


Figure 4.36 ^{13}C -NMR spectra including DEPT-135 and DEPT-90 of 4-hydroxy-benzoic acid methyl ester (7) in methanol- d_4 .

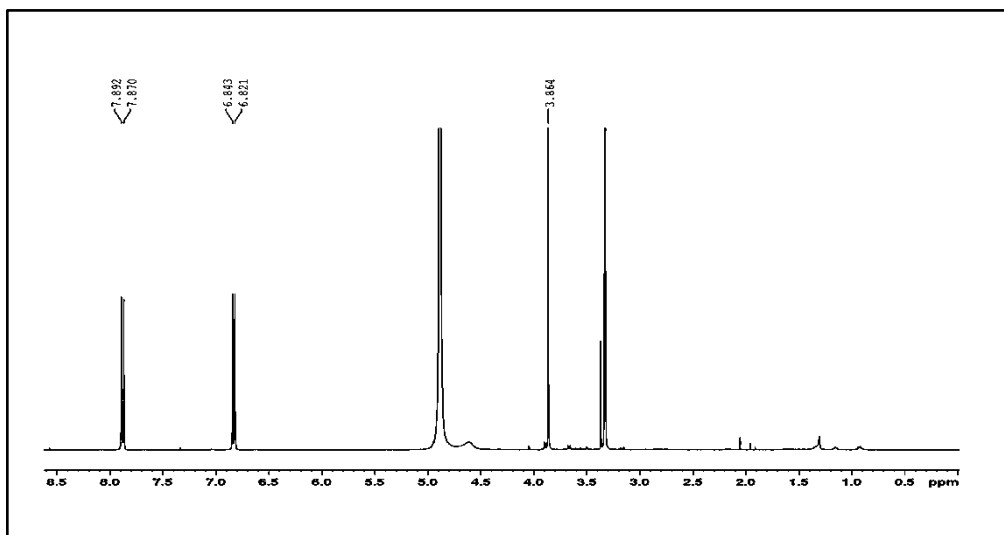


Figure 4.37 $^1\text{H-NMR}$ spectrum of 4-hydroxy-benzoic acid methyl ester (7) in methanol- d_4 .

4.4. BIOLOGICAL ACTIVITIES

4.4.1 Inhibition of AGEs formation by crude extracts

The different fractions were tested for potential antiglycation activity (Table 4.9). The ethyl acetate fraction was highly active (IC_{50} 88 $\mu\text{g/mL}$), whereas a moderate antiglycation activity was observed for the *n*-butanol (36% inhibition at 100 $\mu\text{g/mL}$) and methanol (32% inhibition) fractions. The chloroform fraction presented a mild inhibition of protein glycation (20% inhibition).

Table 4.9. Anti-glycation effects of *K. ramosissima* fractions.

Sample	Antiglycation	
	% Inhibition ^a	IC_{50} ($\mu\text{g/mL}$)
Total Extract	5	-
MeOH 90%	32	-
Chloroform	20	-
Ethyl acetate	64	88
<i>n</i> -Butanol	36	-
Aminoguanidine		19.9

^a at 100 $\mu\text{g/mL}$,

4.4.2 Inhibition of AGEs formation by isolated constituents

In the BSA-glucose assay the prevention of AGE formation was investigated by determining the fluorescence of the complex formed after incubation of a sugar (glucose) and protein (BSA) in the presence or absence of test compounds. In the case of iridoids (compounds **1-4**) only mild inhibition of protein glycation was noticed (Table 4.8). In particular compound **2** was most active (35% inhibition at the highest test concentration) followed by compound **3** (28% inhibition) and compound **1** (26% inhibition). Compound **4** did not present any activity. A nearly similar moderate trend was seen in the BSA-MGO assay. There has only been one previous report on weak inhibition of protein glycation by iridoids (West *et al.*, 2014).

Contrary to iridoids, the isolated flavonoids compound **5** (IC₅₀ 0.79 mM) and **6** (IC₅₀ 2.29 mM) were more active. The fact that the glycosylated compound **6** was less active than the aglycone **5** was indicative of the fact that glycosylation of flavonoids mainly decreases the activity (Matsuda *et al.*, 2003). This inhibition was mainly due to the non-oxidative mode of inhibition as obvious in the BAS- MGO assay, since the IC₅₀ values for compounds **5** (IC₅₀ 0.19 mM) and **6** (IC₅₀ 0.13 mM) were lower than in the glucose-BSA assay. Finally the benzoic acid derivative **7** did not present any inhibition in both models.

Table 4.10. Antiglycation (AGEs) potential of isolated compounds from *Kickxia ramosissima*

Compound	Protein glycation			
	BSA-Glucose		BSA-MGO	
	% inhibition ^a	IC ₅₀ mM	% inhibition ^b	IC ₅₀ mM
1	26		31	
2	35		23	
3	28		24	
4	no activity		9	
5		0.79		0.19
6		2.29		0.13
7	no activity		no activity	
Aminoguanidine		1.75		0.15
Quercetin		0.23		-

^a at 3 mM final concentration

^b at 1.3 mM final concentration

4.4.3 Inhibition of α -glucosidase by isolated constituents

In a sequel of testing the isolated compounds for their antidiabetic potential, the assay on inhibition of α -glucosidase was performed. These agents are able to prevent the fast breakdown of sugars by competitively inhibiting α -glycosidase activity and thus controlling the blood sugar levels. This category of oral hypoglycaemic agents is important in cases of post-prandial blood glucose elevation in diabetic patients. There are quite a few commercially available α -glucosidase inhibitors, but gastrointestinal side effect in patients mainly limit their use. It was therefore considered important to test the isolated compounds for α -glucosidase inhibition.

It was noticed that iridoids presented only mild levels of inhibition (Table 4.9). The highest inhibition (29% inhibition at the highest test concentration) was seen for compound **4** followed by compound **2** (15% inhibition) and compound **3** (9% inhibition). The newly isolated compound **1** did not present any activity. In general a number of reports suggested the potential of iridoids as α -glucosidase inhibitors, but mainly moderate inhibition was noticed (Hua *et al.*, 2014; Lin *et al.*, 2015). Keeping in mind the results of this investigation and the literature, we can conclude that as *K. ramosissima* contains a number of iridoids, the traditional claims about antidiabetic potential could be due in part to the combined effect of all iridoids of the plant.

The flavonoids however presented better activity compared to tested iridoids. Compound **5** showed the highest activity (IC_{50} 0.23 mM), whereas its glycosylated counterpart compound **6** presented moderate α -glucosidase inhibition (48% inhibition at the highest test concentration). The flavonoids are generally known for their multiple therapeutic effects including α -glucosidase inhibitory activity (Tadera *et al.*, 2006; Hong *et al.*, 2013). Also based on our findings, we could conclude that these flavonoids may contribute to the traditional use of *K. ramosissima* in diabetic conditions.

Finally moderate α -glucosidase inhibition (47% inhibition) was observed for the benzoic acid derivative **7**. In conclusion we could emphasise that the combined effect of all isolated compounds i.e iridoids, flavonoids and the benzoic acid derivative are contributing towards the α -glucosidase inhibitory potential of *Kickxia ramosissima*.

Table 4.11 α -Glucosidase inhibitory effect of isolated compounds from *Kickxia ramosissima*.

Compound	% inhibition ^a	IC ₅₀ (mM)
1	no activity	-
2	15	-
3	9	-
4	29	-
5	-	0.23
6	48	-
7	47	-
Acarbose		0.26

^a at 2 mM final concentration

4.4.4 Inhibition of 15-lipoxygenase by isolated constituents

During the current investigation some interesting results were observed for the iridoids as well as the flavonoids. The highest inhibition was seen for compound **2** (IC₅₀ 0.22 mM) followed by compound **3** (37% inhibition at the highest test concentration) and **1** (30% inhibition) (Table 4.12). Likewise in the case of flavonoids compound **6** (IC₅₀ 0.25 mM) presented the highest inhibition, whereas only a moderate activity (46% inhibition) was seen for compound **5**. Finally the benzoic acid derivative **7** presented only a moderate level of inhibition (30% inhibition). Based on these findings we could confirm the possible role of iridoids and flavonoids as anti-inflammatory agents in *Kickxia ramosissima*.

4.4.5 Inhibition of linoleic acid lipid peroxidation - AAPH Assay

During our investigation the isolated iridoids (**1-4**) were evaluated for inhibition of AAPH induced linoleic acid peroxidation. All iridoids exhibited weak or no activity (Table 4.10) compared to standard inhibitor Trolox (Fig 4.36 and Fig 4.37). Our findings were in agreement with earlier reports covering mussaenoside and mussaenosidic acid that also reported weak inhibition (Gousiadou *et al.*, 2013). Also for the flavonoids (**5-6**) a similar trend was observed as previously reported for a different type of flavonoids (Peyrat- Maillard *et al.*, 2003). Finally compound **7** did not present any activity. In conclusion we could say that the isolated compounds presented a weak inhibition of linoleic acid peroxidation that could play a limited role in inhibition of lipid peroxidation.

Table 4.12. 15-lipoxygenase inhibition and inhibition of linoleic acid (AAPH assay) by isolated compounds from *Kickxia ramosissima*.

Compound	15-LOX assay		AAPH assay
	% inhibition ^a	IC ₅₀ (mM)	AOP (min/ μ M) ^c
1	30	-	0.3
2	-	0.22	0.2
3	37	-	0.1
4	8	-	Not tested
5	46	-	0.2
6	-	0.25	0.3
7	30	-	Not active
Quercetin	-	0.14	-
Trolox	-	-	14.12 ^d

^a at 0.312 mM final concentration

^b at 0.06 mM final concentration

^c at 100 μ M

^d at 8 μ M

AOP antioxidant power

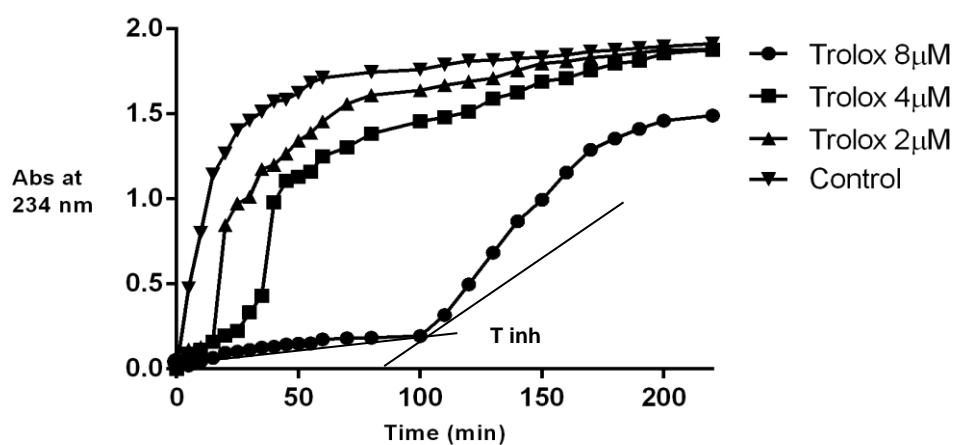


Figure 4.38 Effect of Trolox on AAPH induced linoleic acid oxidation

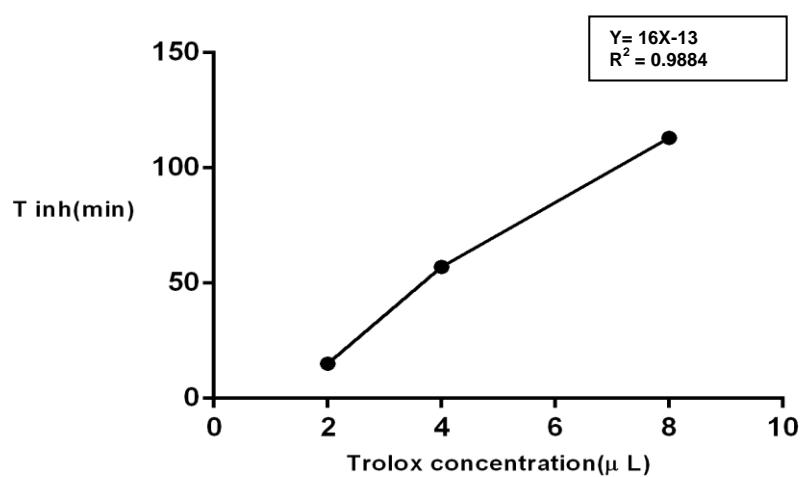


Figure 4.39 Time of inhibition (T_{inh}) as a measure of time.

4.5. Antimicrobial activity

4.5.1 Antimicrobial activity of fractions

The crude extracts were evaluated for antibacterial and antifungal activities. The highest levels of antimicrobial and antifungal activities were observed for n the *n*-hexane fraction, i.e. an IC₅₀ of 8.0 µg/mL against *S. aureus* and 24.4 µg/mL against *M. canis*. None of the other fractions were found active in the tested concentration range (Table 4.13).

Table 4.13 Antibacterial, antifungal and cytotoxic activities of *K. ramosissima* fractions.

Sample	IC ₅₀ µg/mL				
	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>M. canis</i>	MRC-5
MeOH 90%	>64	>64	>64	>64	>64
Chloroform	>64	>64	>64	>64	>64
Ethyl acetate	>64	>64	>64	>64	>64
<i>n</i> -Hexane	8.0	>64	>64	24.4	>64
<i>n</i> -Butanol	>64	>64	>64	>64	>64
Aqueous	>64	>64	>64	>64	>64

Reference: Tamoxifen(MRC-5) IC₅₀ 11.4 µg/mL; erythromycin (*S.aureus*)IC₅₀ 11.2 µg/mL, Trimethoprim (*E.coli*) IC₅₀ 0.25 µg/mL; Miconazole(*C. albicans*) IC₅₀ 5.99 µg/mL; Terbinafin (*M.canis*) IC₅₀ 0.11 µg/mL.

MRC-5: Human fetal lung fibroblasts; *S. aureus*: *Staphylococcus aureus*; *E. coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*.

4.5.2 Antimicrobial activity of isolated constituents

Compounds **2–7** were evaluated in an integrated screening panel for antimicrobial activity. Compound **1** was not tested due to the limited amount available. None of the isolated iridoids including the chlorinated iridoid (**4**) presented antimicrobial activity in the test range. Although few iridoids are well characterized for their antimicrobial activities (Ishiguro *et al.*, 1983; Davini *et al.*, 1986), such activity is surely related to certain structural variations. A number of investigations have reported antimicrobial activity of iridoid rich fractions of medicinal plants, for instance *Morinda citrifolia* (noni) fruits (West *et al.*, 2012). As stated before for *Nymphoides indica*, iridoids can be considered as prodrugs that may be deglycosylated and activated after oral administration.

The flavonoids, however, are well known for their antimicrobial activities (Xie *et al.*, 2015; Mohanty *et al.*, 2015, da Rocha *et al.*, 2015). Also during the current investigation, pectolinarigenin (**5**) was found active against *Staphylococcus aureus* (IC₅₀ 49.8 µM). This compound also showed weak antiprotozoal activities against *Plasmodium falciparum* K1 (IC₅₀ 41.8 µM) and *Trypanosoma cruzi* (IC₅₀ 32.0 µM), accompanied however by PMM cytotoxicity. Moreover it was not cytotoxic against MRC-5 cells (Table 4.11). However, the glucoside of compound **5**, i.e pectolinarine (**6**) did not present any activity in the antimicrobial screening panel.

Table 4.14. Antimicrobial, antifungal cytotoxic and antiprotozoal activities of isolated compounds from *Kickxia ramosissima* (IC₅₀, μM)

Compound	IC ₅₀ (μM)									
	<i>MRC5</i>	<i>Pf-K1</i>	<i>T. bruc</i>	<i>T. cruz</i>	<i>L. inf</i>	<i>PMM</i>	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>A. fumigatus</i>
1	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT
2	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
3	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
4	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
5	> 64	41.8	> 64	32.0	> 64	> 64	49.8	> 64	> 64	> 64
6	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
7	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64

Reference: Tamoxifen (*MRC-5*), IC₅₀ 10.48 μM; Suramine (*T. bruc*), IC₅₀ 0.03 μM, Chloroquine (*Pf-K1*), IC₅₀ 0.08 μM, Miltefosine (*L. inf*), IC₅₀ 9.02 μM; Benznidazol (*T.cruz*), IC₅₀ 2.13 μM; Erythromycin (*S. aureus*) IC₅₀ 11.30 μM; Chloramphenicol (*E.coli*), IC₅₀ 2.42 μM; Miconazole (*C. albicans*), IC₅₀ 4.70 μM; Terbinafine (*A. fumigatus*), IC₅₀ 1.38 μM.

MRC-5: human fetal lung fibroblasts; *Pf-K1*; *Plasmodium falciparum*; *T.bruc*: *Trypanosoma brucei*; *T. cruz*: *Trypanosoma cruzi*; *L.inf*: *Leishmania infantum*; *PMM*: peritoneal murine macrophages; *S.aureus*: *Staphylococcus aureus*; *E.coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*; *A. fumigatus*: *Aspergillus fumigatus*

4.6. SUMMARY AND CONCLUSION

Kickxia ramosissima (Wall.) Janch is a well-known medicinal plant in Pakistan that is traditionally used to treat diabetes mellitus (DM). Indeed quite a few studies have been carried out and some flavonoids, iridoids and sugars were isolated, but the detailed phytochemical and biological analysis of isolated constituents still needed to be performed. Therefore we evaluated various fractions of whole plant material and a series of isolated constituents. The compounds were further analysed for antidiabetic properties (inhibition of α -glucosidase activity; inhibition of Advanced Glycation Endproducts-AGEs), antioxidant activity (AAPH induced linoleic acid peroxidation), inhibition of 15-LOX (anti-inflammatory activity) and antimicrobial activities, as part of running programmes in the host laboratory. Phytochemical analysis of the extracts and fractions led to isolation and identification of 7 compounds, including iridoids (**1-4**) with a new compound, kickxiasine (**1**), mussaenosidic acid (**2**), mussaenoside (**3**), linarioside (**4**); the flavonoids, pectolarigenin (**5**) and pectolarin (**6**); and the benzoic acid derivative 4-hydroxy-benzoic acid methyl ester (**7**) respectively.

In the antidiabetic assays, the ethyl acetate fraction presented highest activity (IC_{50} 88.0 μ g/mL), whereas all other fractions were only moderately active. The isolated iridoids (**1-4**) presented weak inhibition in the BSA-glucose assay, and comparable results in the BSA-MGO assay. The flavonoids, however, showed interesting results as pectolarigenin (**5**) was highly active (IC_{50} 0.79 mM) followed by pectolarigenin (**6**) with IC_{50} 2.29 mM. Also in the BSA-MGO assay both flavonoids show high levels of inhibition i.e pectolarigenin (**5**) IC_{50} 0.19 mM and pectolarigenin (**6**) IC_{50} 0.13 mM. This could clearly indicate the inhibition through non-oxidative pathways. Compound 7 was inactive in all AGEs experiments. In the α -glucosidase inhibition assay, moderate inhibition was recorded for the iridoids. However, with regard to the flavonoids, pectolarigenin showed interesting activity (IC_{50} 0.23 mM), followed by pectolarin (**6**) which showed moderate inhibition (48% inhibition at a concentration of 2 mM). Finally the benzoic acid derivative (**7**) presented moderate inhibition (47% at a concentration of 2 mM).

In the 15-LOX experiment, moderate inhibition was recorded for most compounds, and pectolarigenin (**5**) showed the highest inhibition (46% inhibition at a concentration of 0.312 mM) followed by mussaenoside (**3**) (37% inhibition) and compound **1** (30% inhibition). Compounds **2** and **6** were found inactive. In the AAPH assay, weak or no inhibition was recorded for all compounds. Finally in the antimicrobial screening panel among the fractions, only the *n*-hexane fraction presented moderate activity against *Staphylococcus aureus* (IC₅₀ 8.0 µg/mL), *Micrococcus canis* (IC₅₀ 24.0 µg/mL). With regard to the isolated constituents, only pectolarigenin (**5**) showed moderate activity against *Plasmodium falciparum* K1 (IC₅₀ 41.8 µM), *Staphylococcus aureus* (IC₅₀ 49.8 µM) and *Trypanosoma cruzi* (IC₅₀ 30.0 µM) with no cytotoxicity. Based on our findings we could demonstrate the moderate antidiabetic, anti-inflammatory and antimicrobial activity of some constituents of *Kickxia ramosissima*.

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CHAPTER 5
PHYTOCHEMICAL AND BIOLOGICAL INVESTIGATIONS
ON *FERULA NARTHEX* EXUDATE

Submitted in part for publication in Food Chemistry:

Terpenyl Coumarins from *Ferula narthex* exudate with antiglycation and antimicrobial properties. Adnan Amin, Emmy Tuenter, Paul Cos, Louis Maes, Vassiliki Exarchou, Sandra Apers and Luc Pieters.

5.1 INTRODUCTION

The genus *Ferula* (Apiaceae) is quite diverse and includes around 180 species worldwide. It is commonly found at high altitudes in central Asia and the Middle East (Pimenov and Leonov, 2004). *Ferula narthex* Boiss. is a perennial herb growing at high altitudes (Fig. 5.1). In general its milky exudate is used as spice and folk medicine. Just before flowering in March/ April cuts are made in the upper part of the rhizome root and the milk is collected and dried. The genus *Ferula* is distributed throughout the world, especially in Afghanistan, Iran, India and Pakistan. (Indrayan, 2009). In Pakistan *F. narthex* has been reported mainly from Gilgit, Chitral, Kashmir and Balochistan (Shinwari and Gilani, 2003; Khan *et al.*, 2011; Kakari *et al.*, 2013, Alam *et al.*, 2016). In the Indian subcontinent several species, such as *Ferula gummosa*, *Ferula asafoetida*, *Ferula narthex*, *Ferula costata* and *Ferula latisecta* have been reported (Kapoor, 1990; Eigner and Scholz, 1999; Khan *et al.*, 2011; Kakari *et al.*, 2013).

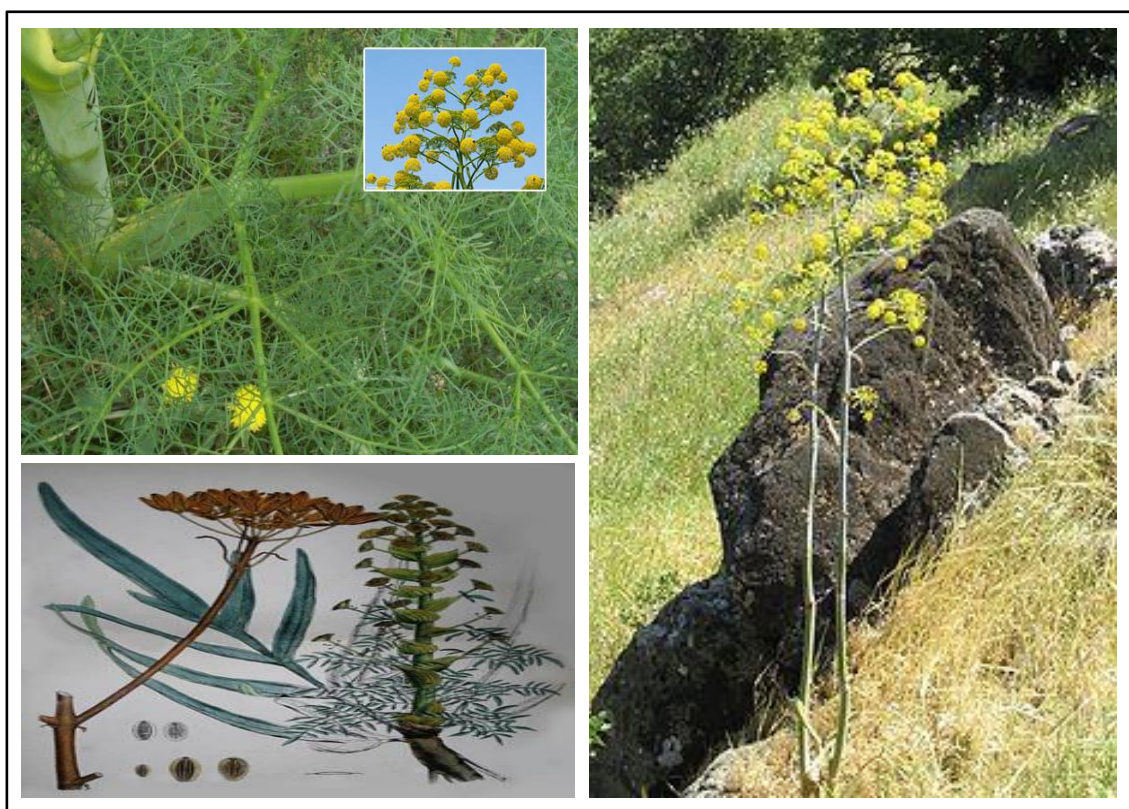


Figure 5.1 *Ferula narthex* in its natural habitat

Ferula narthex Boiss. is commonly called “Heeng”, “Hilteet” or “Raw”. It is used for treatment of a number of ailments including cough, asthma, toothache, gastric problems and also against constipation and angina pectoris (Shinwari and Gilani, 2003, Khan *et al.*, 2011, Srinivasan, 2005, Mahendra and Bisht, 2012). Most recently extracts of *Ferula narthex* have been reported as analgesic (Bashir *et al.*, 2013), insecticidal (Bashir *et al.*, 2013), antimicrobial (Bashir *et al.*, 2014a) and antidiabetic (Iranshahy and Iranshahi, 2011). It is also an important constituent of some traditional formulations (Achliya *et al.*, 2004)

Indeed the genus *Ferula* is mainly known for the occurrence of coumarins, sesquiterpenes and sesquiterpene coumarins (Fig. 5.2) (Iranshahi *et al.*, 2010a; Appendino *et al.*, 1994; Bandyopadhyay *et al.*, 2006,). Compounds isolated from members of the genus *Ferula* have been reported to possess a wide array of activities including antileishmanial, anti-inflammatory, cytotoxic, anticancer, antibacterial and antiviral (Lee *et al.*, 2009; Nazari and Iranshahi, 2011; Bashir *et al.*, 2014b). The extensive traditional usage of the exudate of *Ferula narthex* provoked us to investigate into detail its phytochemical and biological properties.

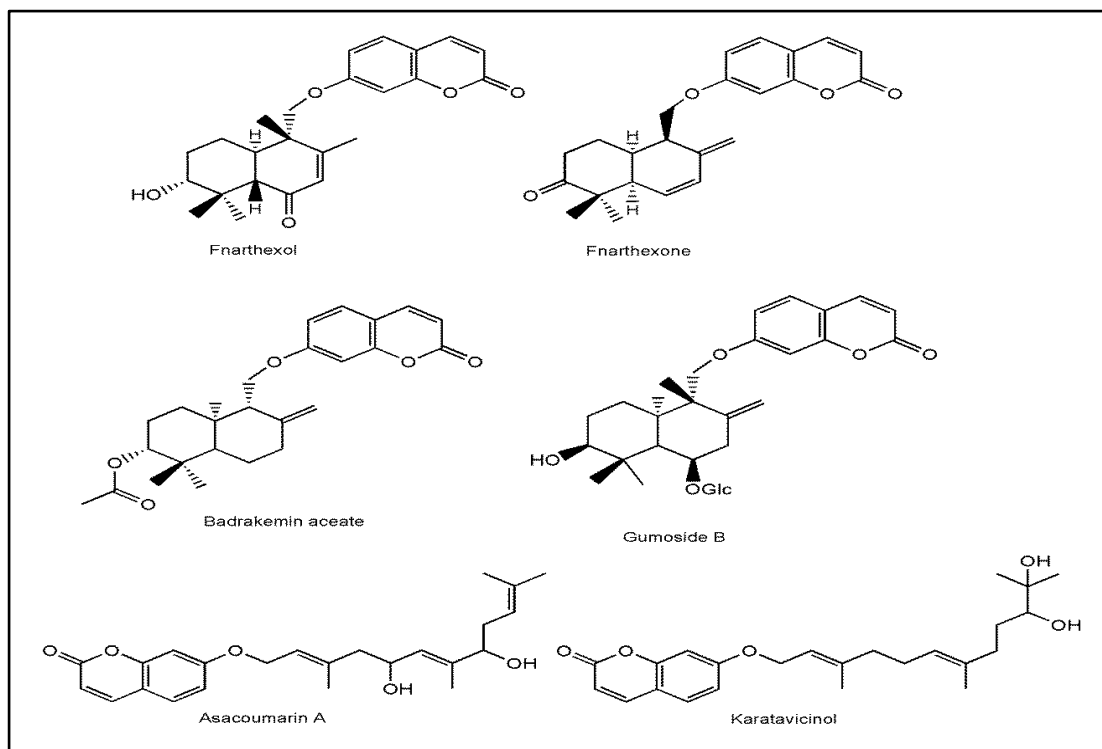


Figure 5.2 Compounds isolated from *Ferula* sp.

5.2 EXTRACTION AND ISOLATION

5.2.1 Plant material

Ferula narthex exudate was collected in September 2012 from the district Chitral (KPK), Pakistan. The samples were identified at the Islamabad Herbarium in the Taxonomy Department, Quaid-I-Azam University, Islamabad, Pakistan, where the voucher specimen was deposited (voucher No. 568 BC, accession no. ISL-72568). The semisolid exudate was stored below 20 °C till further use (Fig. 5.3).



Figure 5.3 *Ferula narthex* dried exudate.

5.2.2. Extraction and fractionation

Dried plant exudate (1 kg) was extracted with 90% (v/v) methanol by double cold maceration. The extract was filtered through Whatman No.1 filter paper using a vacuum pump. The collected filtrate was dried using a rotary evaporator under reduced pressure below 40 °C. The resultant semisolid material was lyophilised with a final yield of 87.1 g, and stored below 20 °C. The liquid–liquid partitioning was performed on the crude extract according to a standard extraction scheme (see chapter 2). After partitioning with different solvents as shown in the scheme, *n*-hexane (1.21g), methanol 90% (25.0 g), chloroform (1.96 g), ethyl acetate (0.2 g), *n*-butanol (4.87 g) and aqueous fractions (53.86 g) were obtained. All collected fractions were dried under reduced pressure at 40 °C, lyophilized and stored below 20 °C.

5.2.3 Phytochemical analysis

Phytochemical screening of extracts and fractions for various compound classes was carried out using published methods (Sofowora, 1993, Trease and Evans, 1989, Harborne, 1973; Egwaikhide and Gimba, 2007; Farnsworth, 1966) with slight modifications for sterols, triterpenes, carbohydrates and saponins flavonoids, alkaloids, and tannins (chapter 2). Phytochemical screening of extracts and fractions indicated that terpenes, alkaloids, tannins and saponins were present at different concentrations depending on their polarity (Table 5.1).

Table 5.1 Phytochemical analysis of *Ferula narthex* extracts and fractions

Extract / fraction	Phytochemicals				
	Alkaloids	Flavonoids	Tannins	Steroids/ triterpenoids	Saponins
MeOH 90%	+	++	-	+++	-
Chloroform	++	++	-	+++	-
Ethyl acetate	+	++	-	+	-
<i>n</i>-Hexane	-	-	-	-	-
<i>n</i>-Butanol	+	+++	+	++	-
Aqueous	-	-	+++	-	-

profoundly present: high degree of precipitation (dark coloration), moderately present: less degree of precipitation (medium colouration), slightly present: very low precipitation (very little colouration), absent: no change in colour (Senguttuvan et al., 2014)

Phytochemicals: ++++ profoundly present; ++ moderately present; + slightly present; - absent.

5.2.4. Thin layer chromatography

The NP (normal phase) TLC for all obtained fractions was executed using various solvent systems as mobile phase. The solvent systems used included CH₂Cl₂ / MeOH (65:35) with a few drops of NH₄OH for the methanolic fraction; CH₂Cl₂ / MeOH (72:28) for the chloroform fraction; *n*-hexane / CH₂Cl₂ (93:7) for the *n*-hexane fraction; CH₂Cl₂ / MeOH (40:60) for the *n*-butanol fraction and CH₂Cl₂ / MeOH (15:85) for the aqueous fraction. Developed TLC plates were examined under UV at 254 nm and 366 nm and after spraying with *p*-anisaldehyde reagent. For instance, TLCs for the methanolic and ethyl acetate fractions are shown in Fig 5.4.

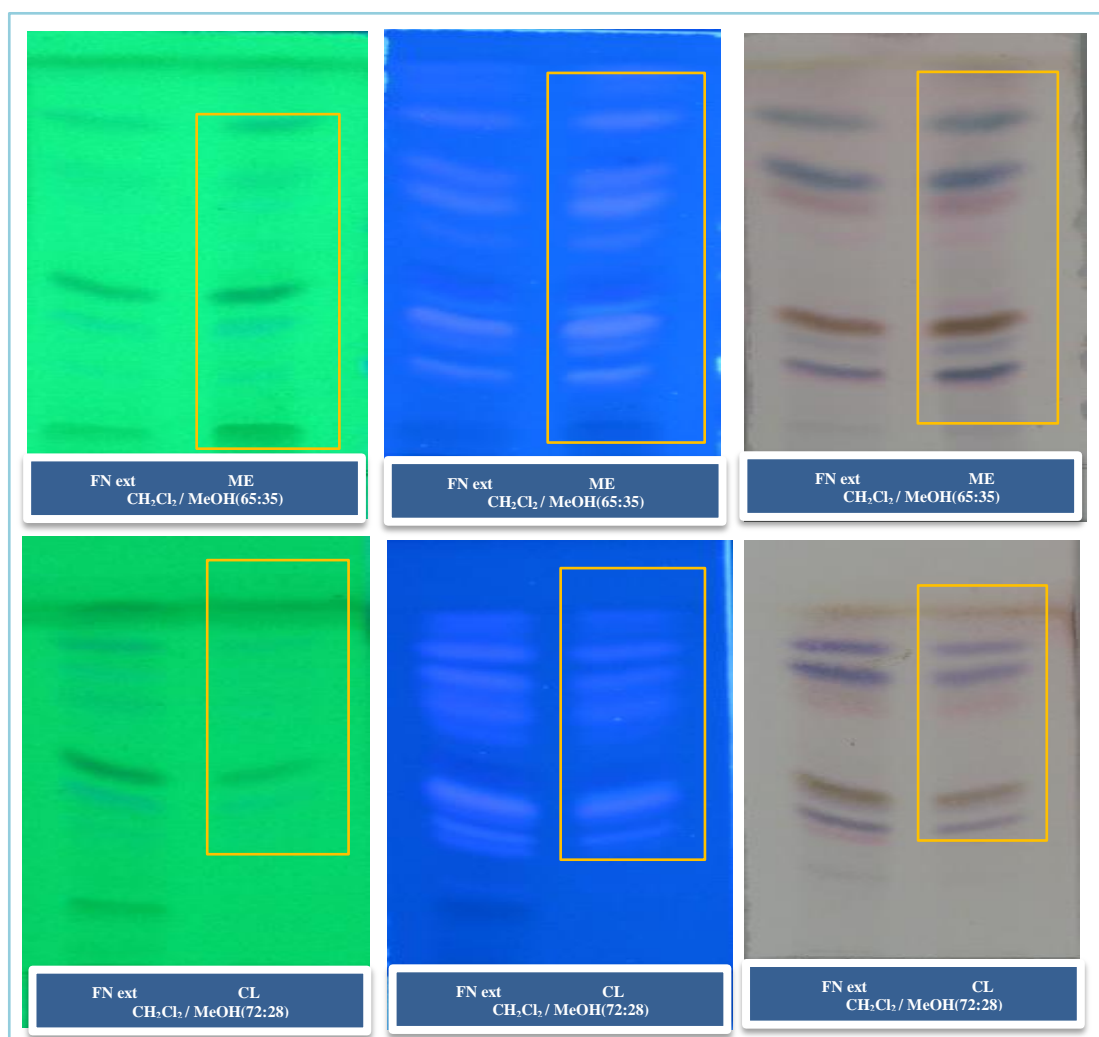


Figure 5.4. TLC profile of total extract (FN ext), methanolic (ME) and chloroform (CL) fractions on silica TLC (NP) plates at 254 and 366 nm after spraying with *p*-anisaldehyde reagent.

5.2.5. Flash chromatography

An aliquot of 0.8 g from the methanol 90% fraction was dissolved in 2 mL methanol and mixed with 1.1g silica; the mixture was dried with nitrogen gas. The dried extract was loaded on a prepacked Flash Grace Reveleris® silica cartridge of 80 g. The compounds were eluted using a gradient from methylene chloride to methanol. Based on UV and ELSD detection, multiple subfractions were collected (Fig. 5.5). All fractions were further analysed by TLC and similar fractions were combined. In this way 16 subfractions were obtained. Flash chromatography with similar conditions was repeated when necessary. Finally, based on TLC profiling, subfractions **FME1** (200 mg), **FME2** (150 mg), **FME4** (220 mg) and **FME5** (210 mg) were selected for further HPLC profiling.

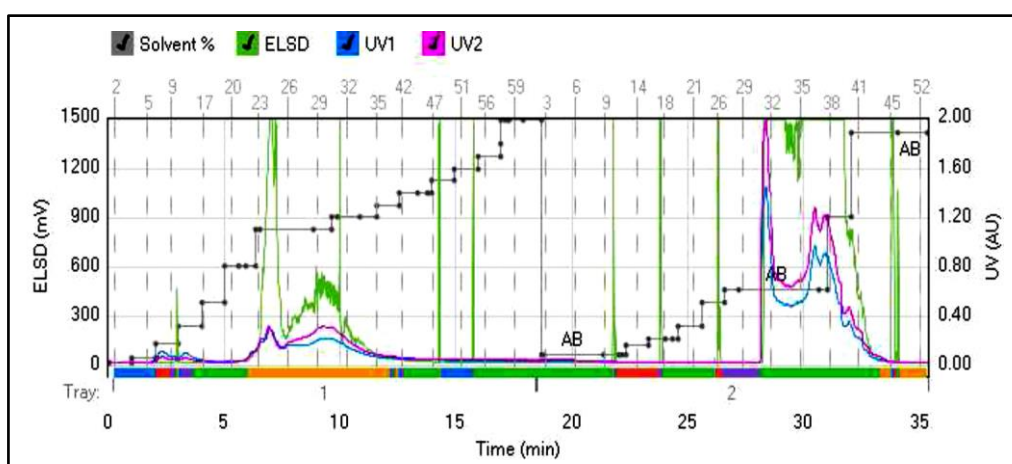


Figure 5.5 Flash chromatogram of the methanolic fraction.

Likewise an aliquot of 0.8 g from the chloroform fraction was loaded on a flash column as discussed above. The compounds were eluted using a gradient from methylene chloride to methanol. Based on UV and ELSD detection different subfractions were collected (Fig 5.6). All fractions were analysed by TLC and similar fractions were combined as described above; in this way 5 subfractions were obtained. In general the TLC of collected fractions revealed similarity with the methanolic fraction as obvious from Fig 5.4. Based on TLC profiling subfractions **FCL1** (120 mg), **FCL2** (140 mg), **FCL3** (180 mg), **FCL4** (100 mg) and **FCL5** (140 mg) were selected further HPLC analysis.

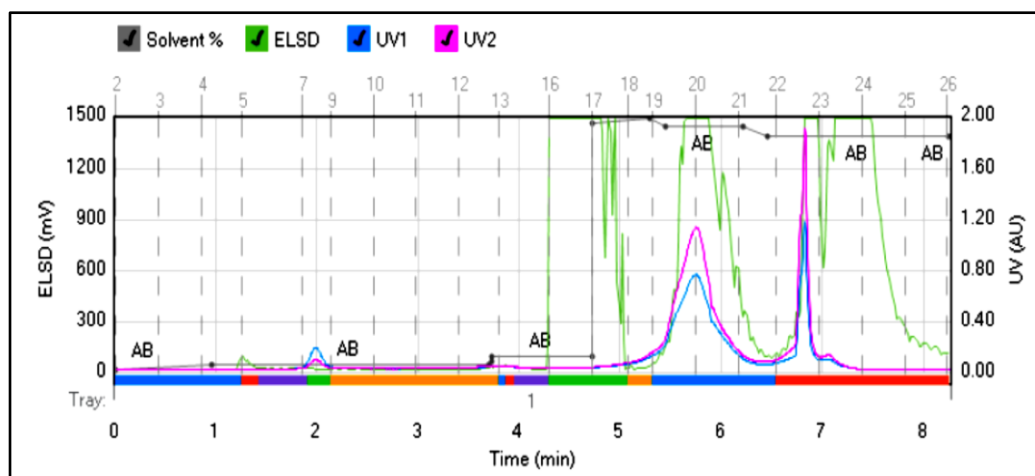


Figure 5.6 Flash chromatogram of the chloroform fraction.

Similarly an aliquot of 0.8 g from the *n*-butanol fraction was subjected to flash chromatography with a gradient from methylene chloride to methanol as previously explained. Finally 10 subfractions were obtained (Fig 5.7). As noticed for the chloroform fraction, the TLC profiling of the *n*-butanol fraction shared some common bands with the methanolic and chloroform fractions. Based on TLC analysis, subfractions **FB2** (140 mg), **FB4** (125 mg), **FB5** (154 mg), **FB6** (200 mg) and **FB7** (116 mg) were selected for HPLC analysis.

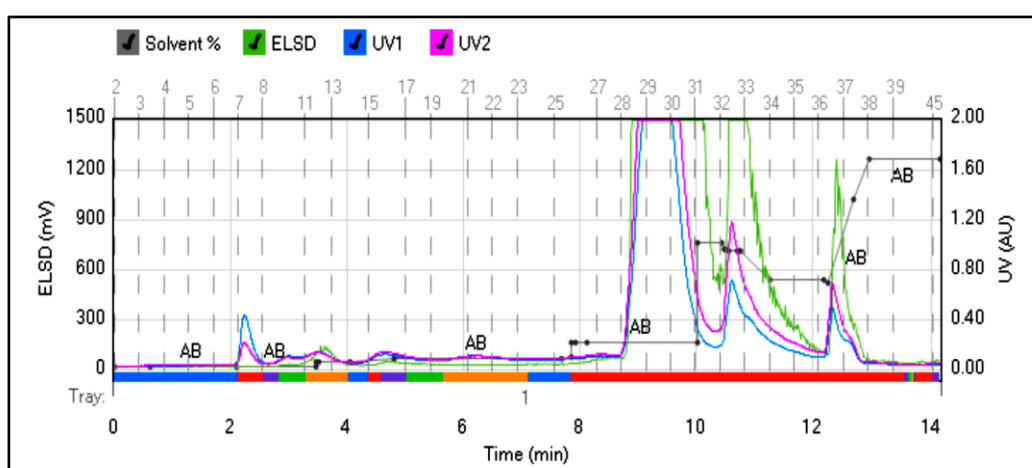


Figure 5.7 Flash chromatogram of the *n*-butanol fraction.

5.2.6. HPLC analysis and isolation of compounds

The methanolic fraction and all obtained subfractions **FME1**, **FME2**, **FME4** and **FME5** were analysed by HPLC using an optimized methanol/H₂O + 0.1% formic acid gradient, ranging from 20% acetonitrile to 80% in 60 min at a flow rate of 1 mL/min. Samples were prepared in a concentration range from 1-10 mg / mL in methanol (Fig 5.8). The isolation of pure compounds was performed by semi-preparative HPLC-DAD-MS using the same gradient at 3 mL/min, yielding compounds **1** (5.5 mg), **2** (4.6 mg) and **3** (6.2 mg).

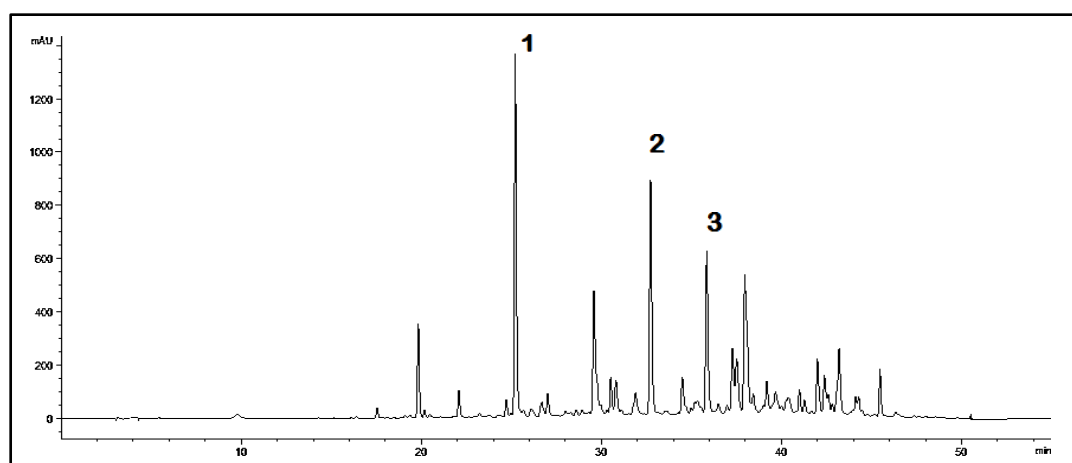


Figure 5.8 HPLC chromatogram of the chloroform fraction **FME2** at 254 nm.

Similarly the crude chloroform fraction and subfractions **FCL1**, **FCL2**, **FCL3**, **FCL4** and **FCL5** were analysed by HPLC using an optimized acetonitrile / H₂O + 0.1% formic acid gradient ranging from 35% acetonitrile to 80% in 45 min at a flow rate of 1 mL/min (Fig. 5.9). The compounds were isolated by semi-preparative HPLC-DAD-MS using the same gradient at a flow rate of 3 mL/min. Finally compounds **2** (2.5 mg), **3** (4.2 mg) and **5** (4.3 mg) were isolated.

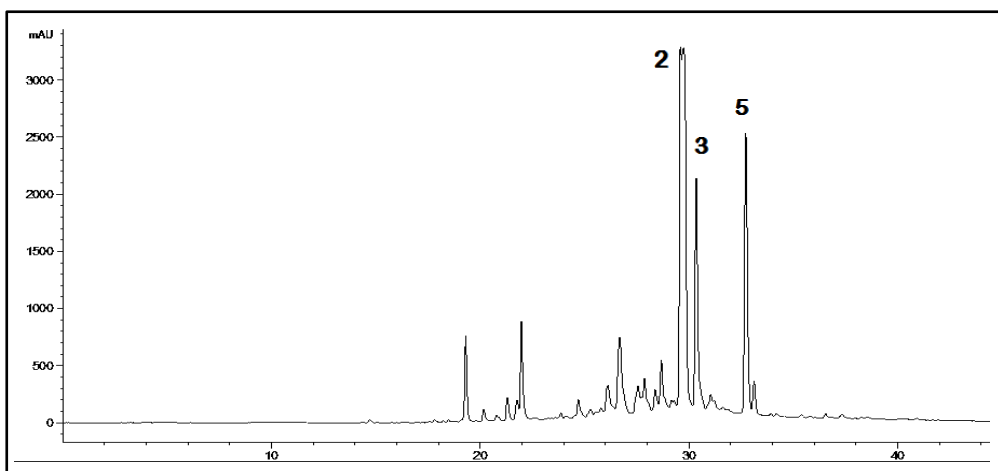


Figure 5.9 HPLC chromatogram of the methanol fraction **FCL 5** at 254 nm

The crude *n*-butanol fraction and subfractions **FB2**, **FB4**, **FB5**, **FB6** and **FB7** were analysed by HPLC using an optimized acetonitrile / H₂O + 0.1% formic acid gradient ranging from 40% acetonitrile to 65% in 50 min at a flow rate of 1 mL/min (Fig. 5.10). Compound isolation was performed by semi-preparative HPLC-DAD-MS using the same solvent gradient as HPLC at a flow rate of 3 mL/min. Compounds **2** (3.6 mg), **5** (4.2 mg) and **6** (4.8 mg) were finally isolated.

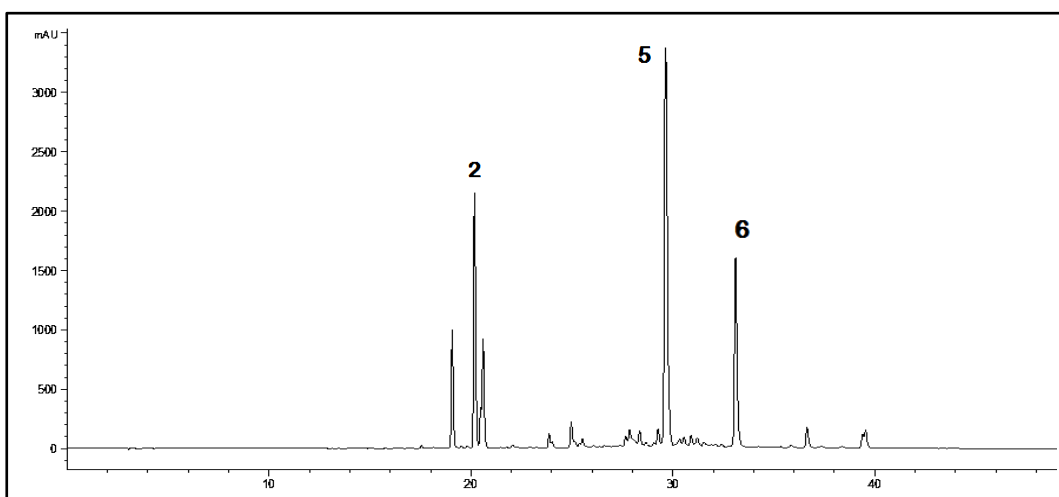


Figure 5.10 HPLC chromatogram of the *n*-butanol fraction **FB4** at 254 nm

5.3. STRUCTURE ELUCIDATION

Structures of the isolated compounds from *Ferula narthex* were elucidated using ^1H - and ^{13}C - NMR (including DEPT-135 and DEPT-90) as well as 2D-NMR (COSY, HSQC and HMBC) spectroscopy. The molecular ion was derived from the mass spectra obtained with the semi-preparative HPLC-DAD-MS system.

5.3.1 Fnarthexol (1)

The structure of compound **1** was elucidated by ^1H -NMR, ^{13}C -NMR (DEPT-135 and DEPT-90) and 2D-NMR (COSY and HSQC) as fnarthexol (Fig. 5.10).

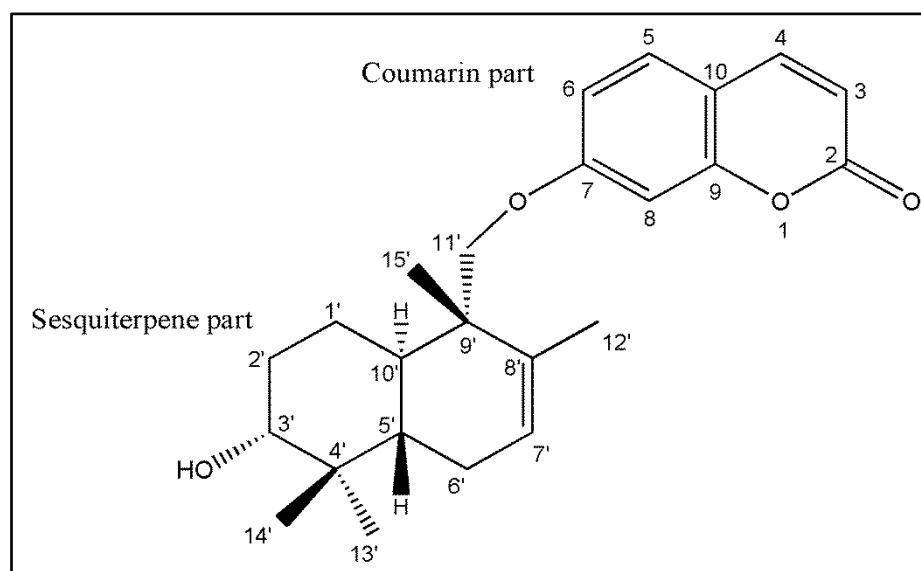


Figure 5.10 Structure of fnarthexol (1)

Library search on the ^{13}C -NMR data (Table 5.2 and Fig. 5.11) using the NMR Predict database and ^1H -NMR (Table 4.2 and Fig. 5.12) suggested that this compound was a sesquiterpene-coumarin. The spectral data were compared with NMR data from literature (Bashir *et al.*, 2014_b), which indicated that compound **1** was fnarthexol. Analysis of the COSY (Fig. 5.13) and HSQC (Fig. 5.14) correlations confirmed the structure as shown.

In ^{13}C -NMR (Table 5.2, Fig. 5.14) the sesquiterpene ring carbons C-1' and C-2' showed resonance signals at 24.3 ppm and 28.0 ppm, respectively. The signals at 79.6 ppm (indicative of oxygen substitution), 39.9 ppm and 50.9 ppm were assignable to C-3', C-4' and C-5', respectively. Similarly C-6' was attributed to 39.0 ppm. In the same ring C-7', C-8' and C-9' were assigned to the signals at 124.5 ppm, 133.9 ppm and 37.0 ppm respectively. The resonance at C-10 was attributed to the signal at 55.4 ppm whereas the peak at 68.2 ppm was assignable to C-11', i.e. a methylene group substituted by oxygen. Four methyl signals appearing at 21.8 ppm, 15.3 ppm, 28.7 ppm and 15.9 ppm were assigned to C-12', C-13', C-14' and C-15', respectively.

In the coumarin ring the carbonyl group (C-2) appeared at 163.4 ppm. The resonance signals at 113.6 ppm and 145.9 ppm were assigned to the double bond carbons C-3 and C-4. Similarly the CH-groups C-5, 6 and C-8 were attributed to the peaks at 130.5 ppm, 114.4 ppm and 102.2 ppm, respectively. Finally the signals at 163.7 ppm, 157.2 ppm and 114.1 ppm were assigned to the quaternary carbons C-7, C-9 and C-10, respectively.

In the ^1H -NMR spectrum (Fig 4.15 and Fig 4.16) of the sesquiterpene part, methylene signals at 2.07 ppm (m) and 1.63 / 0.92 ppm (m) and 2.0 ppm (m) were assigned to H-1', H-2' and H-6', respectively. An oxygenated methylene signal at 3.70 ppm (d, $J=8.5$ Hz) and 3.99 ppm (d, $J= 8.5$ Hz) was assigned to H-11'. The oxygenated methane signal observed at 3.22 ppm (dd) was assignable to H-3'. A multiplet appearing at 1.98 ppm was assigned to H-5', whereas the signal at 2.20 ppm was assigned to H-10'. A broad singlet appearing at 5.53 ppm was assigned to the olefinic proton H-7'. Finally four methyl signals at 1.69 ppm, 0.87 ppm, 0.98 ppm and 0.93 ppm were assigned to H-12', H-13', H-14' and H-15', respectively.

In the coumarin ring protons H-3 and H-4 (i.e. the double bond) corresponded to signals at 6.23 ppm and 7.90 ppm, respectively, whereas protons H-5 and H-6 resonated 4.55 ppm (d) and 6.95 ppm (dd), respectively. Finally a broad singlet appearing at 6.93 ppm was assigned to H-8.

Yellow powder; UV (acetonitrile / H₂O) λ_{max} 243, 325 nm; ESI-MS (positive ion mode): m/z 405 [M+Na]⁺ consistent with a molecular formula C₂₄H₃₀O₄.

Fnarthexol has only been isolated recently from *Ferula narthex* (Bashir *et al.*, 2014_b).

Table 5.2 ¹H and ¹³C NMR assignments for compound (1) recorded in MeOH-d₄

Position	δ_H (ppm); multiplicity; J (Hz)	δ_C (ppm)
1		
2		163.4
3	6.23; d; $J= 9.5$	113.6
4	7.90; d; $J= 9.5$	145.9
5	7.54; d; $J= 8.5$	130.5
6	6.93; br d; $J= 8.5$	114.4
7		163.7
8	6.93; br s	102.2
9		157.2
10		114.1
1'	2.07; m	24.3
2'	1.63; 0.92; m	28.0
3'	3.23; dd; $J=10.9; 4.5$	79.6
4'		39.9
5'	1.26; m	50.9
6'	2.0; m	39.0
7'	5.53; br s	124.5
8'		133.9
9'		37.0
10'	2.20; br s	55.4
11'	3.70; d; $J= 8.5$ 3.99; d; $J= 8.5$	68.2
12'	1.69; s	21.8
13'	0.87; s	15.3
14'	0.98; s	28.7
15'	0.93; s	15.9

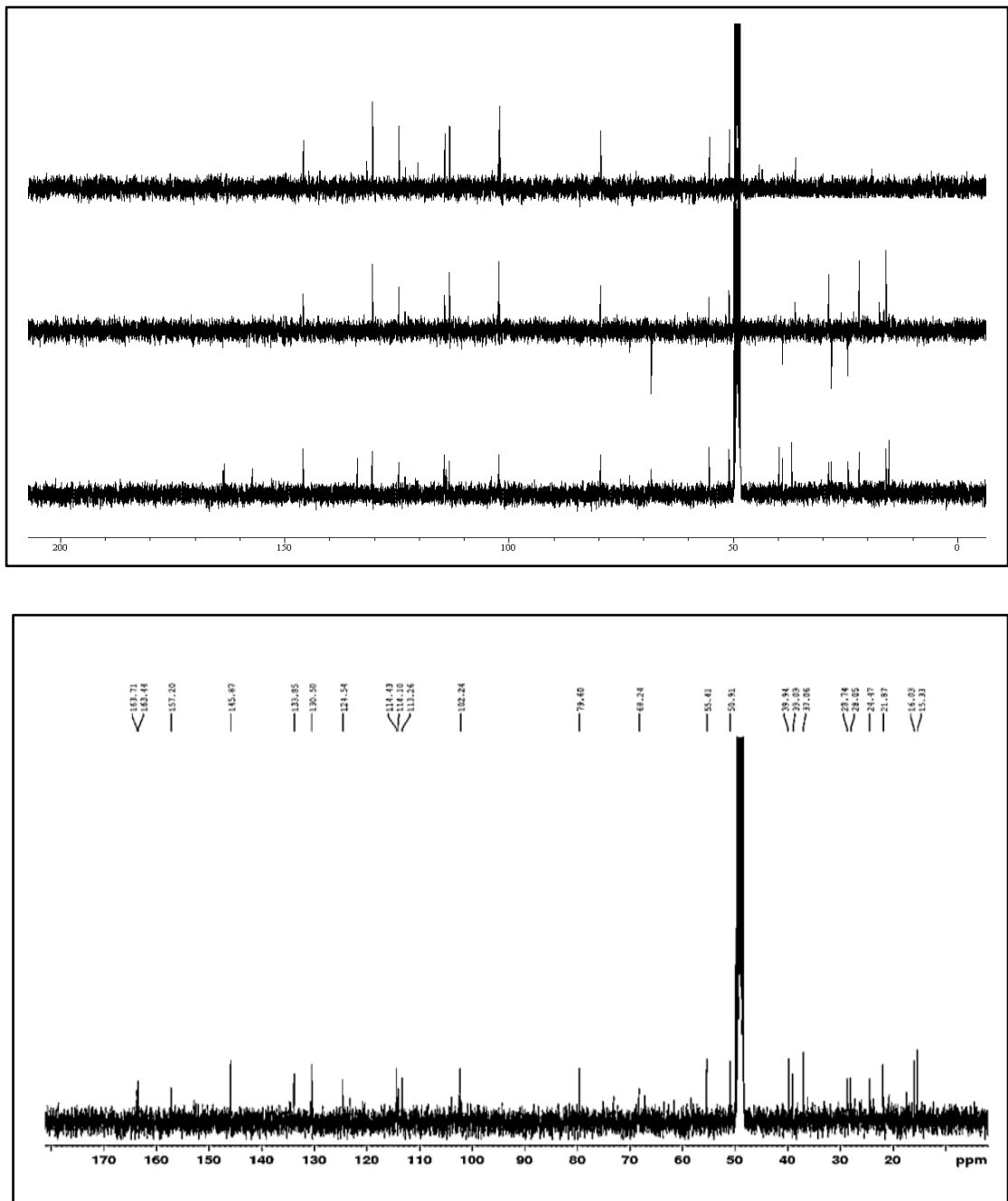


Figure 5.11 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of compound **1** in methanol- d_4 .

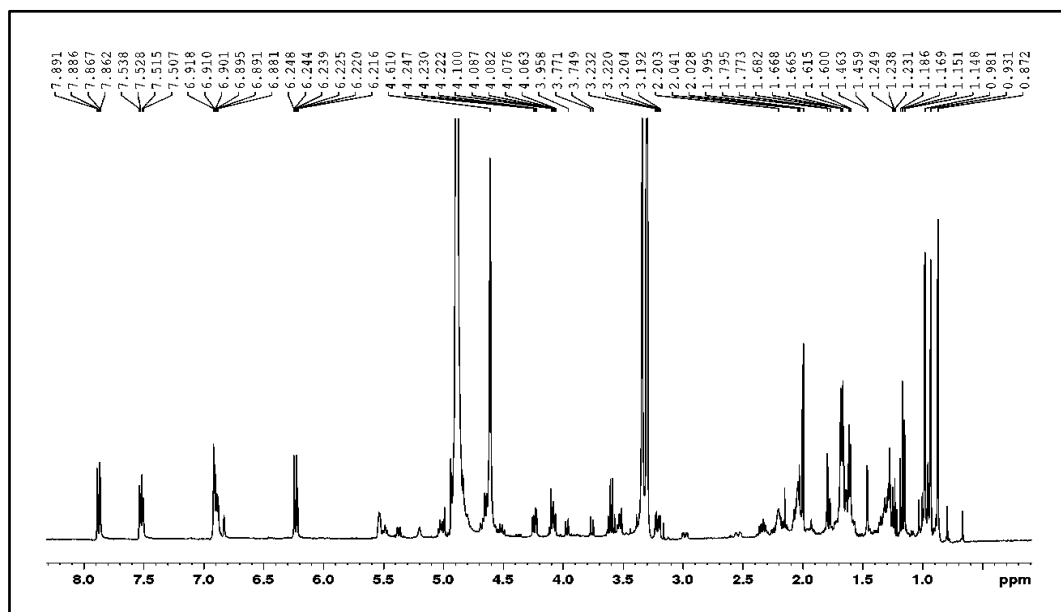


Figure 5.12 ¹H-NMR spectrum of compound 1 in methanol-d₄

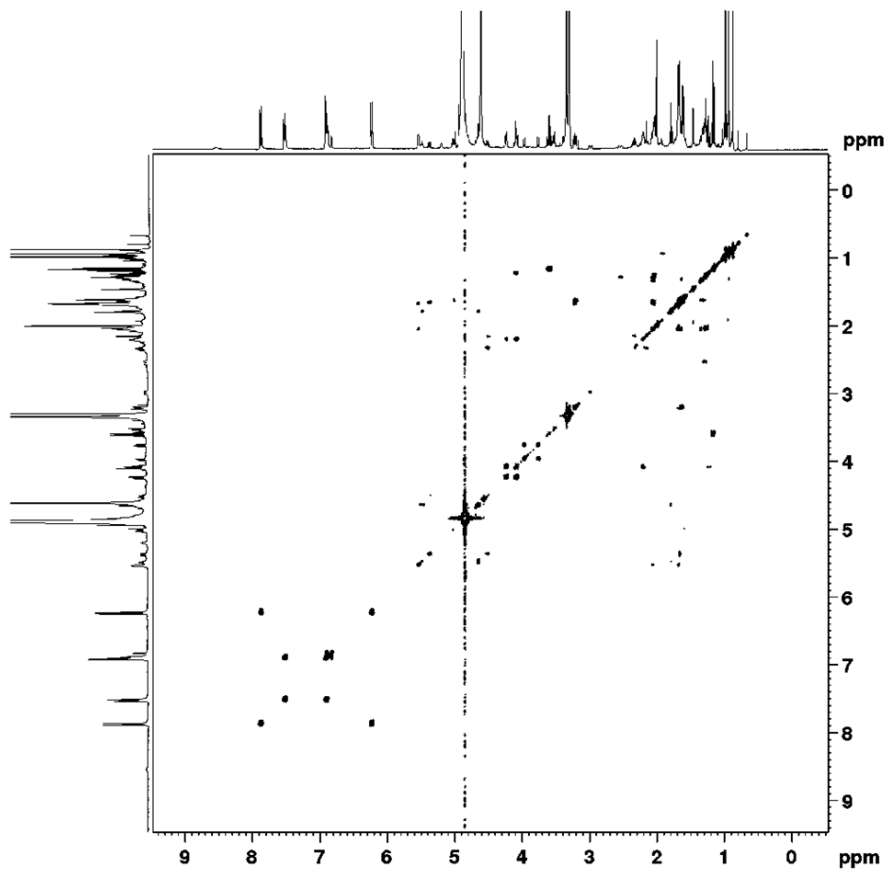


Figure 5.13 COSY spectrum of compound 1

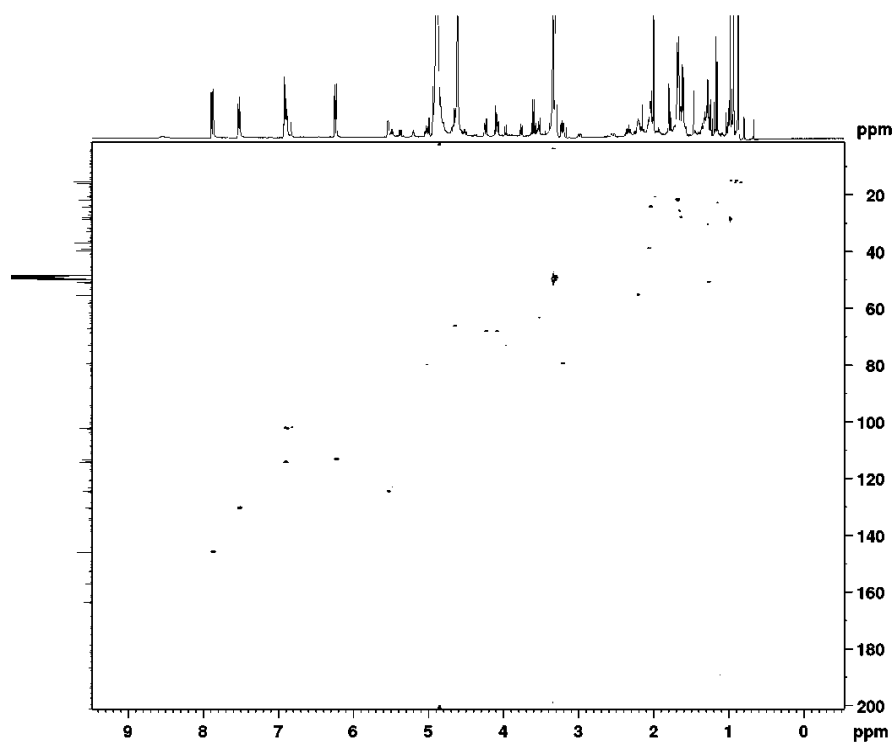


Figure 5.14 HSQC spectrum of compound **1**

5.3.2 Ligupersin A (2)

The structure of compound **2** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) and 2D-NMR (COSY, HSQC and HMBC) as **ligupersin A** (Fig. 5.15).

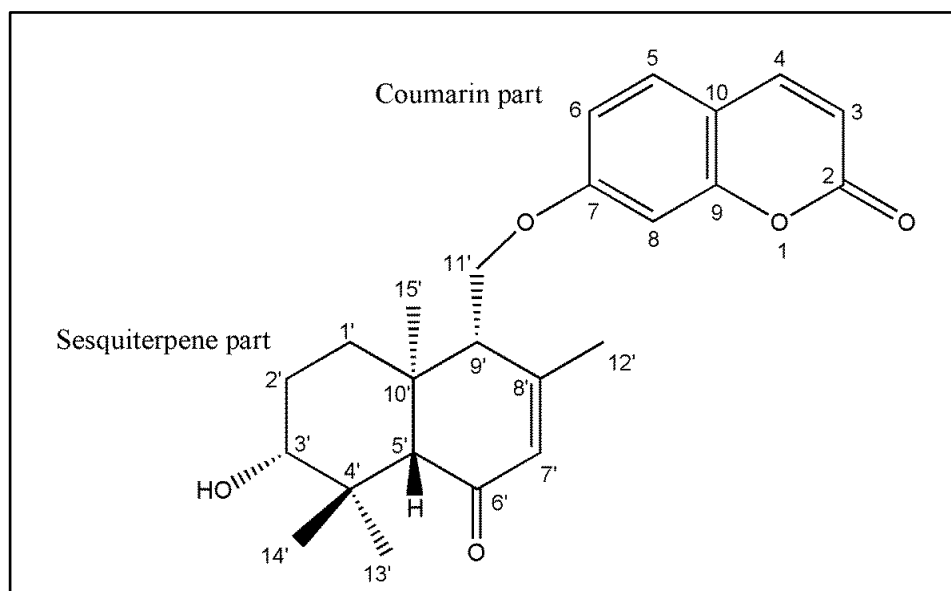


Figure 5.15 Structure of ligupersin A (2)

Library search on the $^{13}\text{C-NMR}$ data (Table 5.3 and Fig 5.16) using the NMR Predict database and $^1\text{H-NMR}$ (Table 5.3 and Fig. 5.17) suggested that this compound was a sesquiterpene-coumarin. The spectral data were compared with NMR data from literature (Marco *et al.*, 1991) which suggested that the compound was ligupersin A. Analysis of the COSY (Fig. 5.18), HSQC (Fig. 5.19) and HMBC (Fig. 5.20) correlations confirmed the structure as shown.

In ^{13}C NMR (Table 5.3, Fig. 5.18) the sesquiterpene ring carbons C-1' and C-2' (both methylene groups) showed resonance signals at 38.5 ppm and 27.6 ppm, respectively. The signals at 79.4 ppm (indicative of oxygen substitution), 39.1 ppm and 63.7 were assignable to C-3', C-4' and C-5', respectively. The carbonyl group at C-6 was attributed to the peak at 201.9 ppm. In the same ring C-7' and C-8' (double bond) were assigned to the signals at 130.7 ppm and 157.4 ppm, respectively. In the same ring C-9' and C-10' were observed at 56.5 ppm and 43.6 ppm, respectively. The resonance at 67.1 ppm was assignable to the oxygen-substituted methylene group C-11'. Finally four methyl signals appearing at 22.0 ppm, 16.0 ppm, 29.1 ppm and 16.7 ppm were assigned to C-12', C-13', C-14' and C-15', respectively.

In the coumarin ring the carbonyl group (C-2) appeared at 163.2 ppm. The resonance signals at 114.0 ppm and 145.9 ppm (double bond) were assigned to C-3 and C-4, respectively. C-5, 6 and C-8 were attributed to signals at 130.7 ppm, 114.5 ppm and 102.6 ppm, respectively. Finally signals at 163.5 ppm, 159.6 ppm and 114.4 ppm were assigned to C-7, C-9 and C-10, respectively.

In the ¹H-NMR spectrum (Table 5.3, and Fig. 5.19) of the sesquiterpene part, multiplets appearing at 1.67 ppm and 2.06 ppm were assigned to coupled protons at H-1', whereas multiplets at 1.65 ppm and 1.72 ppm were attributed to H-2'. The resonance at 3.20 ppm (dd, *J*= 4.3, 11.0 Hz) was assignable to H-3'. Indeed, the chemical shift indicated the presence of a hydroxyl group with α-orientation, as the *J* values were in agreement with published data (Marco *et al.*, 1991). A singlet appearing at 2.33 ppm was assigned to H-5'. The resonance signals appearing at 5.84 ppm (s) and 2.81 ppm (m) were assigned to H-7' and H-9', respectively. Likewise the resonances at 4.32 ppm (dd) and 4.40 ppm (dd) were attributed to H-11'. Four methyl signals were recorded at 2.0 ppm, 1.16 ppm, 1.26 ppm and 1.06 ppm were assigned to H-12', H-13', H-14' and H-15', respectively. This typical pattern of methyl groups suggested a drimane skeleton in the sesquiterpene part of the compound.

In the coumarin ring, protons H-3 and H-4 (double bond) corresponded to signals at 6.24 ppm and 7.88 ppm, respectively, whereas H-5 and H-6 were observed at 5.84 ppm (d) and 6.95 ppm (dd), respectively. Finally a doublet (*J*=2.5) appearing at 6.98 ppm was assigned to H-8.

Yellow powder; UV (acetonitrile / H₂O) λ_{max} 218, 236, 296, 325 nm; ESI-MS (positive ion mode); *m/z* 419 [M+Na]⁺ consistent with a molecular formula C₂₄H₂₈O₅.

Ligupersin A has been isolated from various medicinal plants including *Ligularia persica* (Marco *et al.*, 1991), *Ferula gummosa* (Iranshahi *et al.*, 2010_a), *Ferula flabelliloba* (Iranshahi *et al.*, 2010_b). However this is the first report in *Ferula narthex*.

Table 5.3 ¹H and ¹³C NMR assignments for compound (2) recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; J (Hz)	δ_{C} (ppm)
2		163.3
3	6.24; dd; $J= 9.5$	114.0
4	7.88; d; $J= 9.5$	145.9
5	7.84; d; $J= 8.5$	130.7
6	6.95; $J= 8.5; 2.5$	114.5
7		163.5
8	6.98; d; $J= 2.5$	102.6
9		159.6
10		114.4
1'	1.67; 2.06; m	38.5
2'	1.65; 1.72; m	27.6
3'	3.20; $J= \text{dd}; 4.3; 11.0$	79.4
4'		39.1
5'	2.33; s	63.7
6'		201.9
7'	5.84; br s	130.1
8'		157.4
9'	2.81; m	56.5
10'		43.6
11'	4.32; dd; $J= 10.5; 5.2$ 4.40; dd; $J= 10.5; 3.0$	67.1
12'	2.0; s	22.0
13'	1.16; s	16.0
14'	1.26; s	29.1
15'	1.06; s	16.7

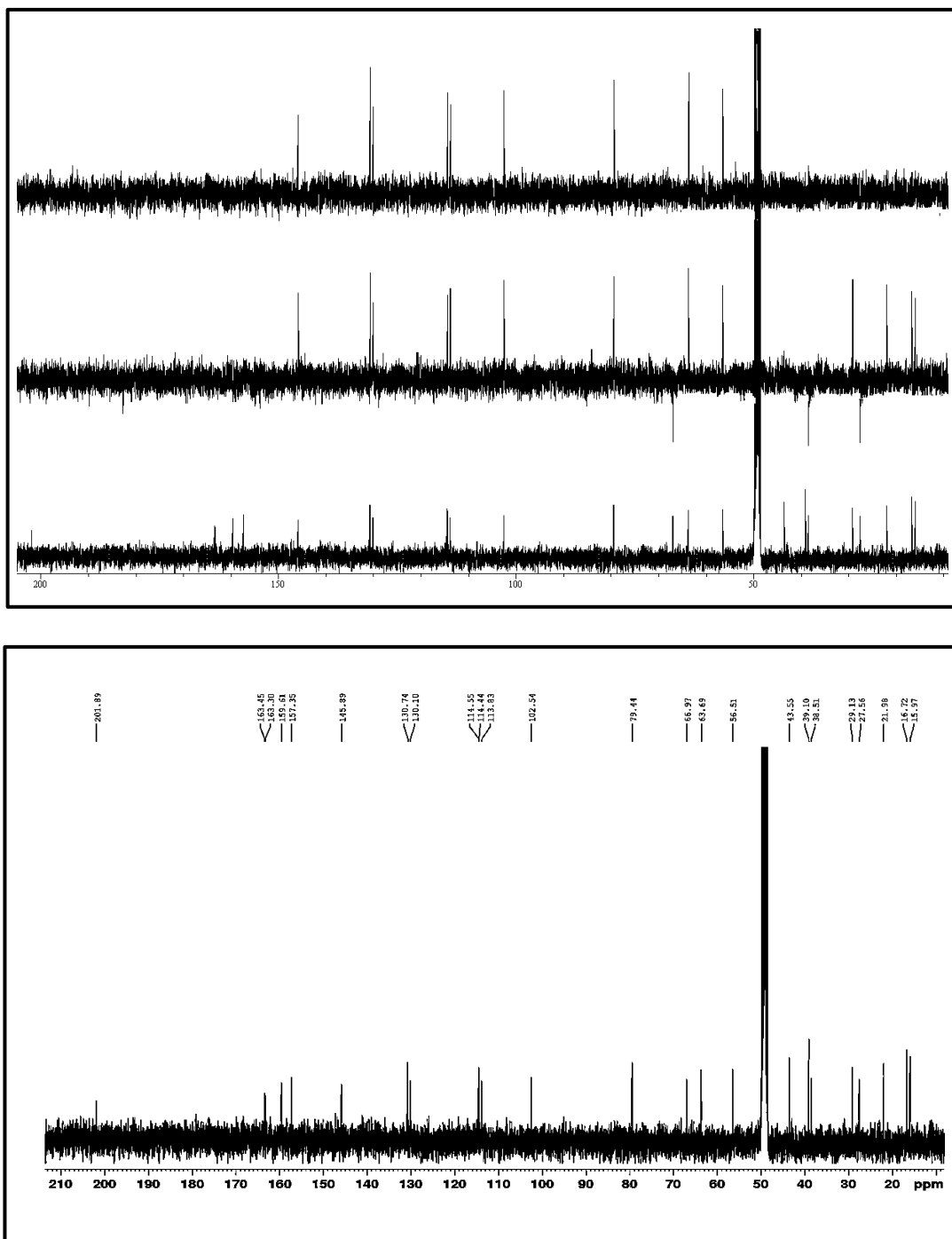


Figure 5.16 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of compound **2** in methanol- d_4 .

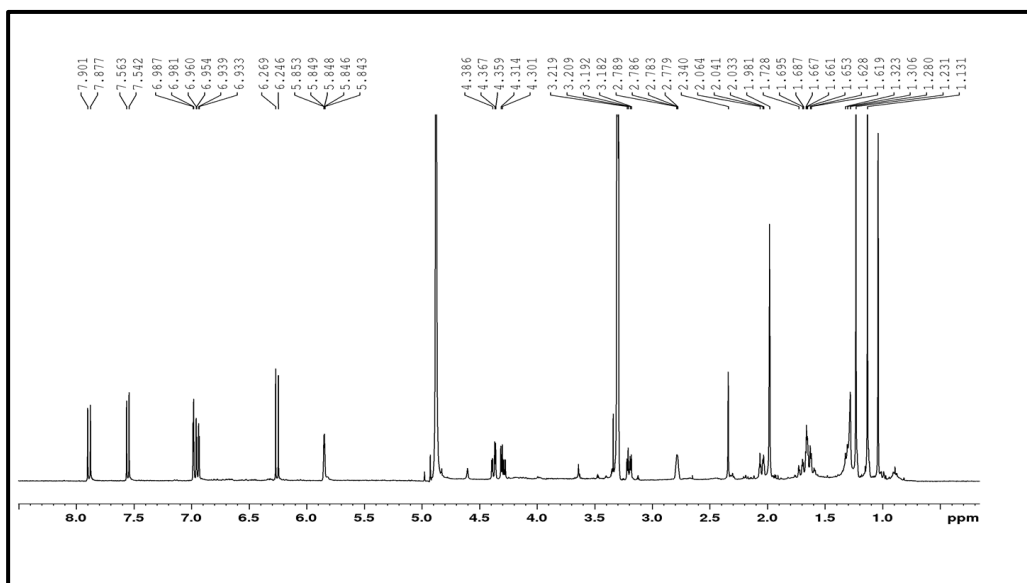


Figure 5.17 $^1\text{H-NMR}$ spectrum of compound **2** in methanol- d_4

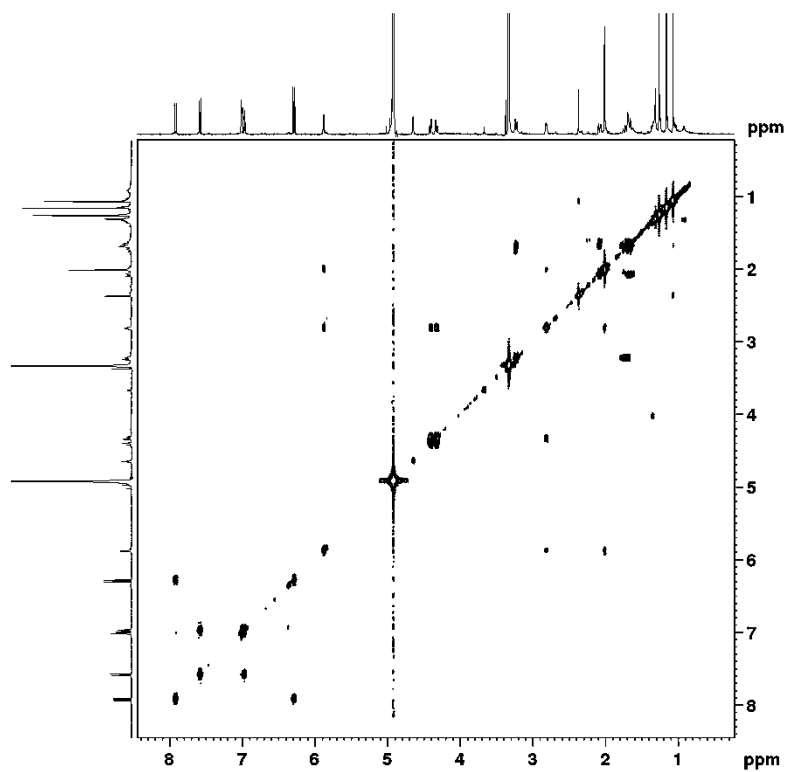


Figure 5.18 COSY spectrum of compound 2

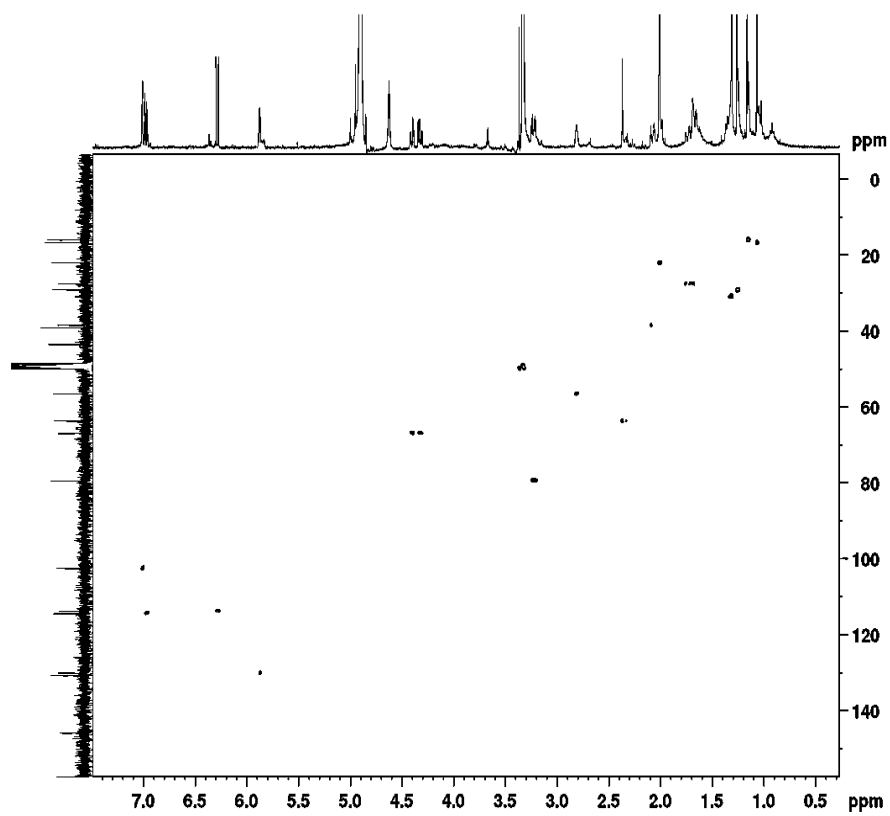


Figure 5.19 HSQC spectrum of compound 2

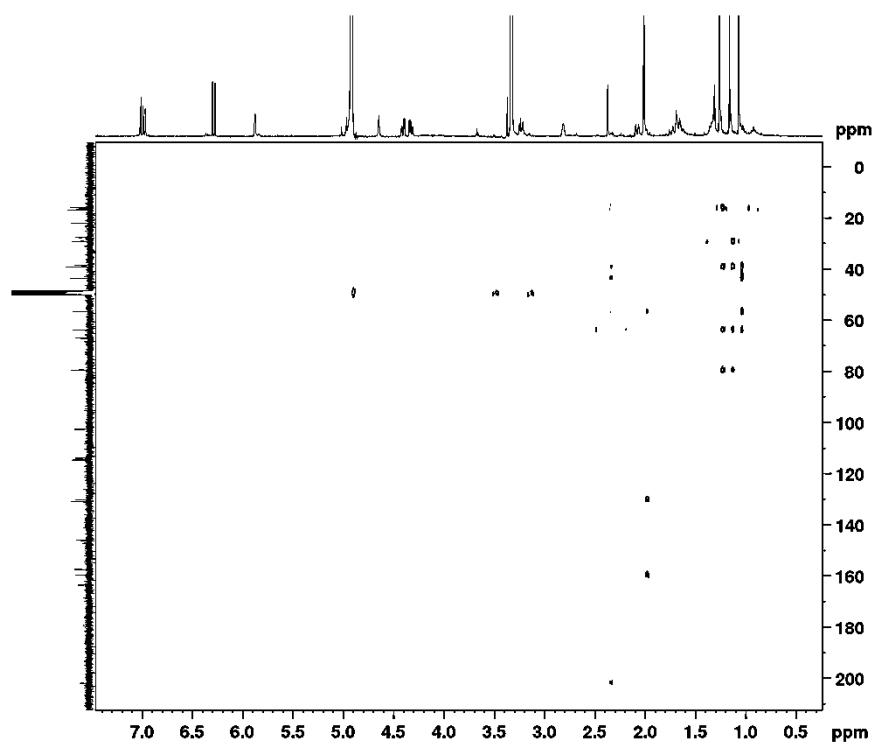


Figure 5.20 HMBC spectrum of compound 2

5.3.3 8'-Acetoxy-5'-hydroxyumbelliprenin (3)

The structure of compound **3** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) as **8'-acetoxy-5'-hydroxyumbelliprenin** (Fig. 5.21).

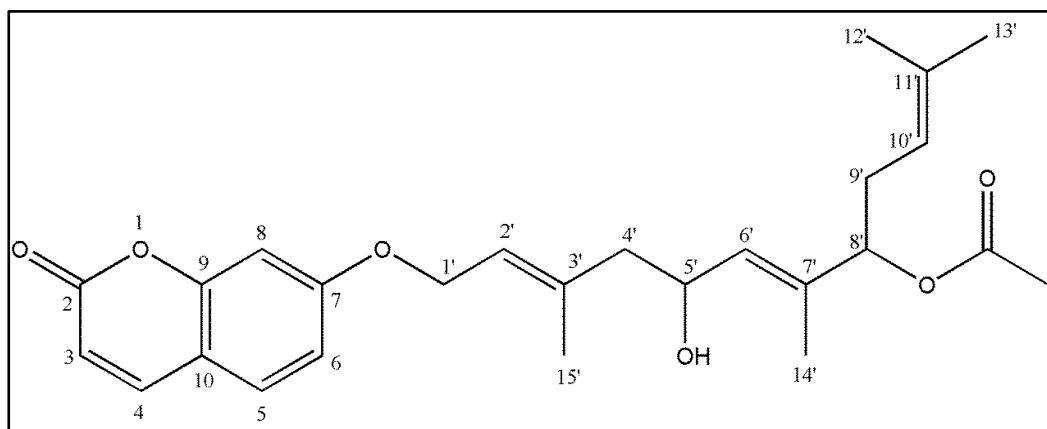


Figure 5.21 Structure of 8'-acetoxy-5'-hydroxyumbelliprenin (**3**)

Library search on the $^{13}\text{C-NMR}$ data (Table 5.4 and Fig 5.22) using the NMR Predict database and $^1\text{H-NMR}$ (Table 5.4 and Fig. 5.23) suggested that this compound was a substituted coumarin with a terpene side chain. The spectral data were compared with NMR data from literature (Appendino *et al.*, 1994), which indicated that the compound was 8'-acetoxy-5'-hydroxy umbelliprenin.

In $^{13}\text{C-NMR}$ the coumarin ring carbonyl group (C-2) showed a resonance at 163.4 ppm. The signals at 113.4 ppm and 145.8 ppm (double bond) were assigned to C-3 and C-4, respectively. C-5, 6 and C-8 were attributed to peaks at 130.5 ppm, 114.4 ppm and 102.6 ppm, respectively, and signals at 163.8 ppm, 157.1 ppm and 114.4 ppm to C-7, C-9 and C-10, respectively. In the aliphatic chain three methylene signals appeared at 66.7 ppm, 48.3 ppm and 32.9 ppm, and were assigned to C-1', C-4' and C-9', respectively. Resonance signals at 123.2 ppm, 139.3 ppm and 67.3 ppm were attributed to C-2', C-3' and C-5'. Carbon signals appearing at 131.9 ppm, 136.4 ppm and 80.0 were assignable to C-6', C-7' and C-8'. Finally C-10' and C-11' were assigned to the peaks at 120.3 ppm and 135.4 ppm.

Four methyl signals at 12.8 ppm, 18.0 ppm, 17.4 ppm and 26.0 ppm were attributed to C-12', C-13', C-14' and C-15', respectively. The acetoxy group appeared at 172.0 ppm (carbonyl) and 21.0 ppm (methyl).

In the ¹H-NMR spectrum (Fig 4.15 and Fig 4.16) of the coumarin part H-3 and H-4 (double bond) corresponded to signals at 6.21 ppm and 7.90 ppm, respectively, whereas H-5 and H-6 resonated at 7.51 ppm (d) and 6.90 ppm (dd), respectively. Finally a doublet ($J=2.5$ Hz) appearing at 6.84 ppm was assigned to H-8.

In the aliphatic chain signals appearing at 4.64 ppm and 5.50 ppm were attributed to H-1' and H-2', whereas multiplets at 2.34 ppm and 2.16 ppm were assigned to H-4'. The upfield chemical shift of the signal at 4.51 ppm, showed the presence of a hydroxyl group and this signal was assigned to H-5'. A broad doublet ($J=8.2$ Hz) at 5.40 ppm corresponded to H-6', whereas a triplet appearing 5.02 ppm was assigned to H-8'. The methylene protons at H-9' were assigned to signals at 2.33 ppm and 2.24 ppm (m), whereas H-10' was attributed to the signal at 4.83 ppm. The methyl groups H-12', H-13', H-14' and H-15' were assigned to signals at 1.60 ppm, 1.70 ppm, 1.80 ppm and 1.80 ppm, respectively. The singlet appearing at 2.0 ppm was assigned to the acetoxy group.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 205, 325 nm; ESI-MS (positive ion mode): m/z 463 consistent with a molecular formula C₂₆H₃₂O₆.

8'-Acetoxy-5'-hydroxyumbelliprenin has been reported previously been reported from asafetida, the gum resin obtained by incision of the roots of various plants from the genus *Ferula* (Appendino *et al.*, 1994; Lee *et al.*, 2009).

Table 5.4 ¹H and ¹³C NMR assignments for compound (3) recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; J (Hz)	δ_{C} (ppm)
2		163.4
3	6.21; d; $J= 8.5$	113.4
4	7.90; d; $J= 9.5$	145.8
5	7.51; d; $J= 8.5$	130.5
6	6.90; dd; $J= 8.0; 2.5$	114.4
7		163.8
8	6.84; d; $J= 2.5$	102.6
9		157.1
10		114.0
1'	4.64; br d; $J= 6.2$	66.7
2'	5.50; br t; $J=6.2$	123.2
3'		139.3
4'	2.34; m 2.16; m	48.3
5'	4.51; m;	67.3
6'	5.40; br d; $J= 8.2$	131.9
7'		136.4
8'	5.02; t; $J= 7.0$	80.0
9'	2.33; m 2.24; m	32.8
10'	4.83; overlapped with solvent peak	120.3
11'		135.4
12'	1.60; br s	12.8
13'	1.70; s	18.0
14'	1.68; s	17.4
15'	1.80; br s	26.0
Ac	2.0; s	21.0
		172.0

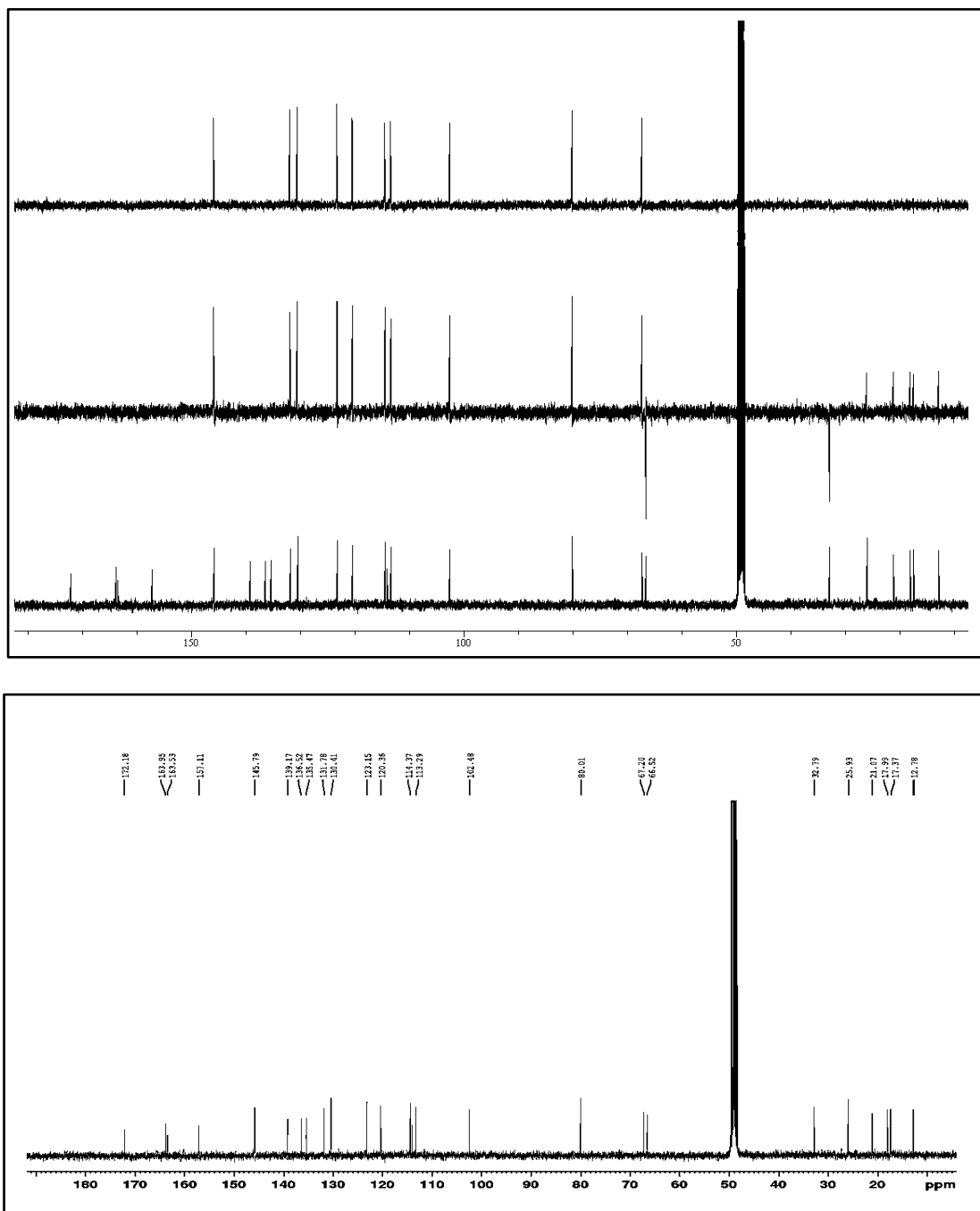


Figure 5.22 ¹³C-NMR, DEPT-135 and DEPT-90 spectra of compound 3 in methanol-d₄.

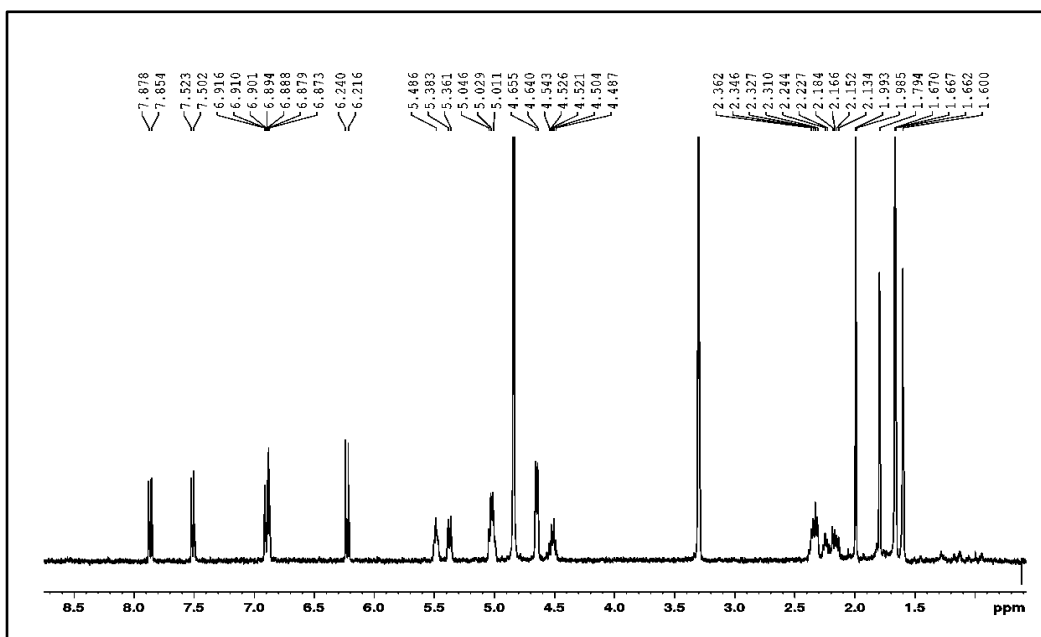


Figure 5.23 ¹H-NMR spectrum of compound 3 in methanol-d₄

5.3.4 Asacoumarin A (4)

The structure of compound **4** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) and 2D-NMR (COSY, HSQC, and HMBC) as **asacoumarin A** (Fig. 5.24).

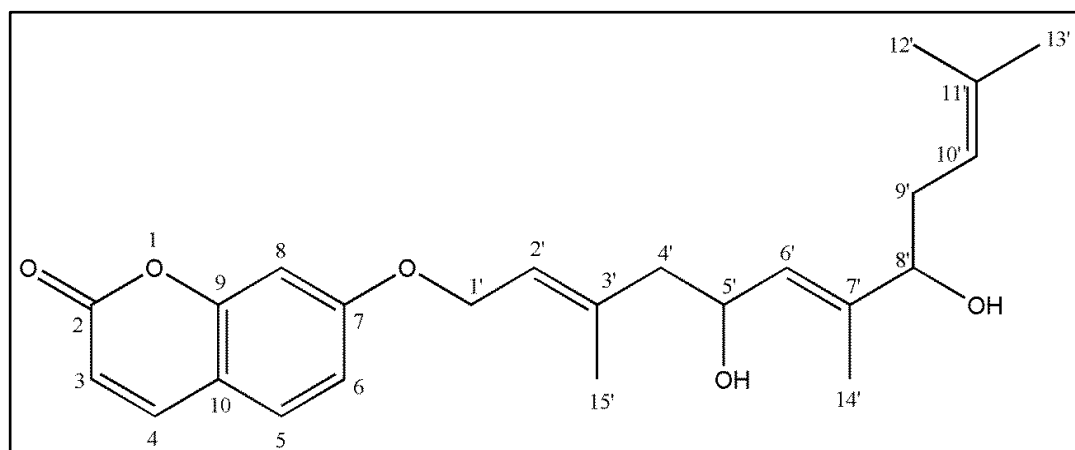


Figure 5.24 Structure of asacoumarin A (4)

Library search on the $^{13}\text{C-NMR}$ data (Table 4.5 and Fig 5.25) using the NMR Predict database and $^1\text{H-NMR}$ (Table 4.5 and Fig 5.26) suggested that this compound was a substituted coumarin. The spectral data were compared with NMR data from the literature (Abd El-Razek *et al.*, 2007) and analysis of the COSY (Fig. 5.27), HSQC (Fig. 5.28) and HMBC (Fig. 5.29) correlations confirmed the structure as asacoumarin A.

In $^{13}\text{C-NMR}$ 24 signals were recorded. The coumarin carbonyl (C-2) show a resonance at 163.4 ppm. The signals at 113.2 ppm and 145.8 ppm were assigned to C-3 and C-4. Similarly, the CH-groups C-5, 6 and C-8 were attributed to the signals at 130.3 ppm, 114.4 ppm and 102.2 ppm, respectively, whereas the at 163.8 ppm, , 157.1 ppm and 113.9 ppm were assigned to the quaternary carbons C-7, C-9 and C-10, respectively.

In the aliphatic chain three methylene signals appeared at 66.5 ppm, 48.3 ppm and 34.9 ppm, assigned to C-1', C-4' and C-9', respectively. Likewise the signals at 123.0 ppm, 140.1 ppm and 67.4 ppm were attributed to C-2', C-3' and C-5'. Carbon signals appearing at 130.4 ppm, 139.3 ppm and 78.5 were assignable to C-6', C-7' and C-8'. Finally C-10' and C-11' were assigned to 121.8 ppm and 133.9 ppm. Four methyl signals

at 18.0 ppm, 26.0 ppm, 17.3 ppm and 11.6 ppm were attributed to C-12', C-13', C-14' and C-15', respectively.

In the ¹H-NMR spectrum (Table 4.5 and Fig. 5.28) coumarin and terpene signals were recorded. In the coumarin part, protons H-3 and H-4 (double bond) corresponded to signals at 6.22 ppm and 7.88 ppm, respectively, whereas H-5 and H-6 resonated 7.50 ppm (d) and 6.88 ppm (dd), respectively. A doublet appearing at 6.85 ppm was assigned to H-8.

In the terpene part (aliphatic chain) signals appearing at 4.64 ppm and 5.50 ppm were attributed to H-1' and H-2', whereas multiplets observed at 2.34 ppm and 2.14 ppm were assigned to H-4'. The upfield chemical shift of the signal at 4.51 ppm, due to the presence of a hydroxyl group, was assigned to H-5'. Similarly a broad doublet at 5.30 ppm corresponded to H-6', whereas a triplet appearing at 3.89 ppm was assigned to H-8'. The methylene protons at H-9' were assigned to a multiplet at 2.11 ppm, whereas H-10' was attributed to the peak at 5.05 ppm. The methyl signals at H-12', H-13', H-14' and H-15' were assigned to signals at 1.60 ppm, 1.66 ppm, 1.80 ppm and 1.60 ppm, respectively.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 203, 325 nm; ESI-MS (positive ion mode): m/z 421 [M+Na]⁺ consistent with a molecular formula C₂₄H₃₀O₅.

Asacoumarine A has been isolated before from asafoetida (Appendino *et al.*, 1994; Kajimoto *et al.*, 1989; Abd El-Razek *et al.*, 2007), However, this is the first report in *Ferula narthex*.

Table 5.4 ¹H and ¹³C NMR assignments for compound (4) recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; J (Hz)	δ_{C} (ppm)
2		163.4
3	6.22; d; $J= 9.5$	113.2
4	7.88; d; $J= 9.5$	145.8
5	7.50; d; $J= 8.5$	130.3
6	6.88; dd; $J= 8.0; 2.5$	114.4
7		163.8
8	6.85; d; $J= 2.5$	102.2
9		157.1
10		113.9
1'	4.64; d; $J= 6.4$	66.5
2'	5.50; br t; $J=6.2$	123.0
3'		140.1
4'	2.34; m 2.14; m	48.3
5'	4.51; dd; $J=15.0; 6.4$	67.4
6'	5.3; d; $J= 8.2$	130.4
7'		139.3
8'	3.89; t; $J= 6.7$	78.5
9'	2.11; m	34.9
10'	5.05; t; $J=7.0$	121.8
11'		133.9
12'	1.60; br s	18.0
13'	1.66; s	26.0
14'	1.8; s	17.3
15'	1.60; br s	11.6

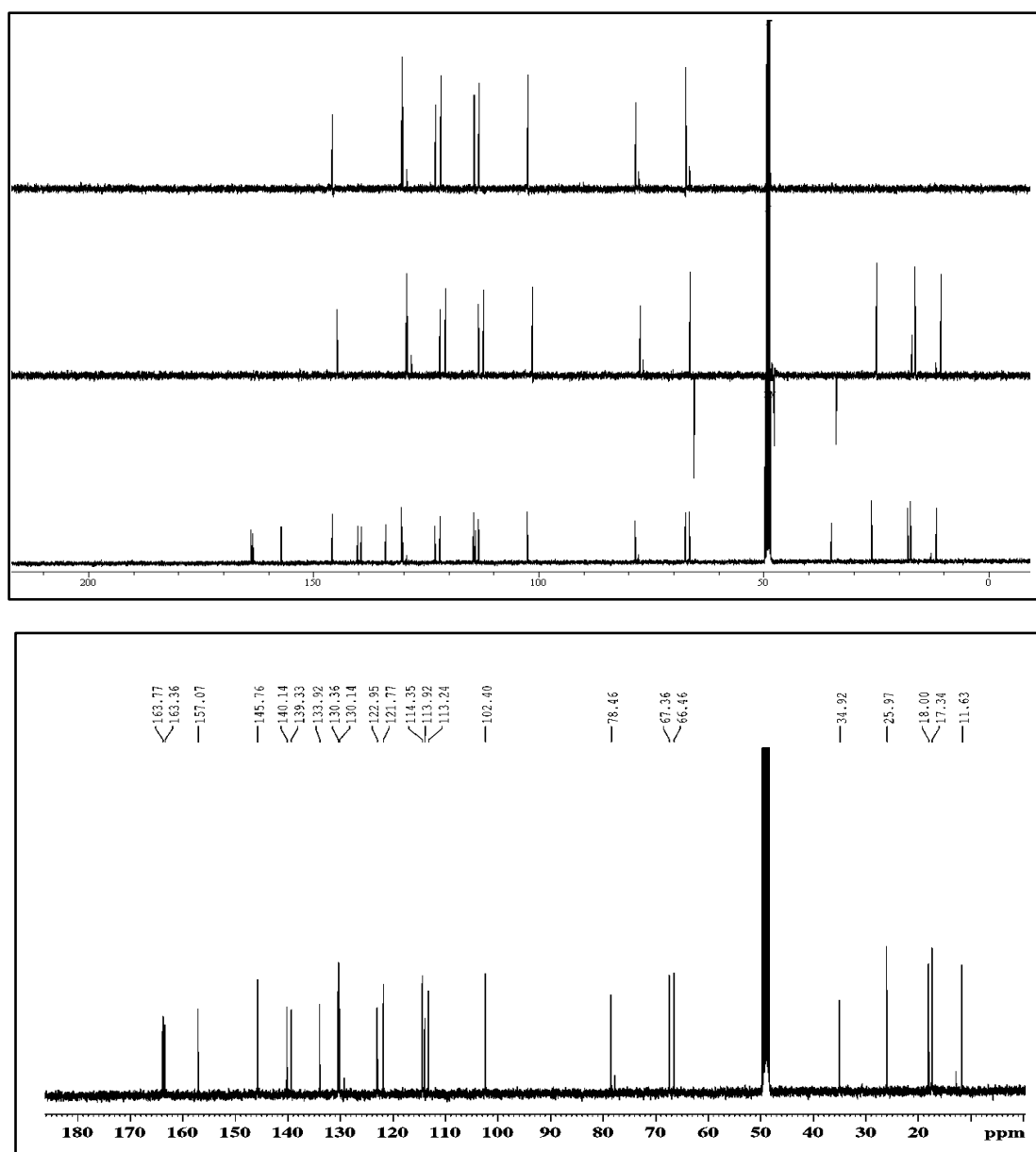


Figure 5.25 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of compound **4** in methanol- d_4 .

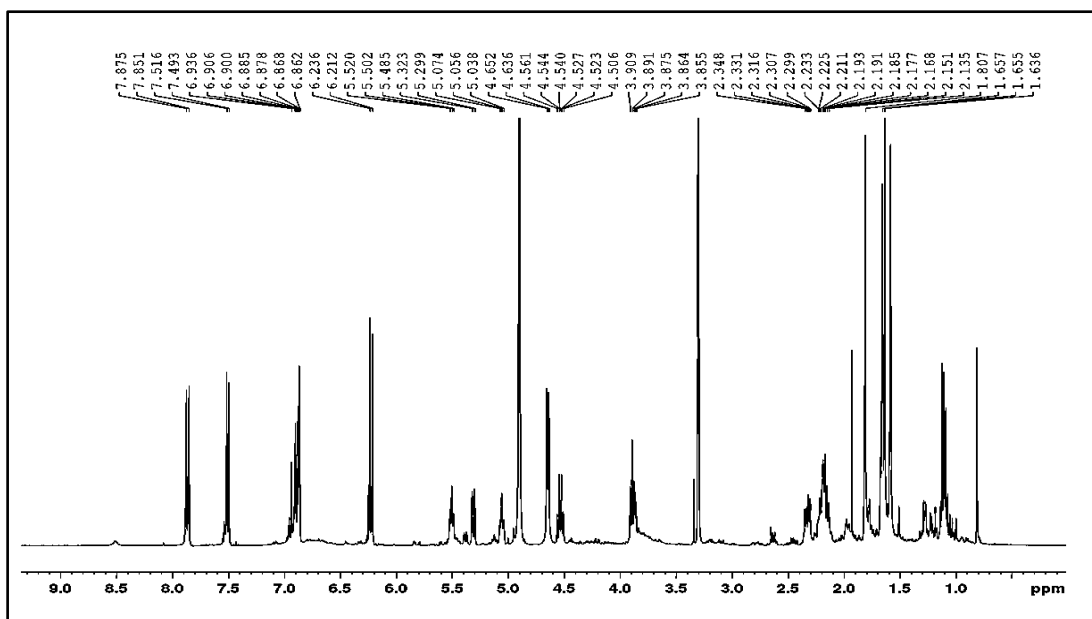


Figure 5.26 ¹H-NMR spectrum of compound **4** in methanol-d₄

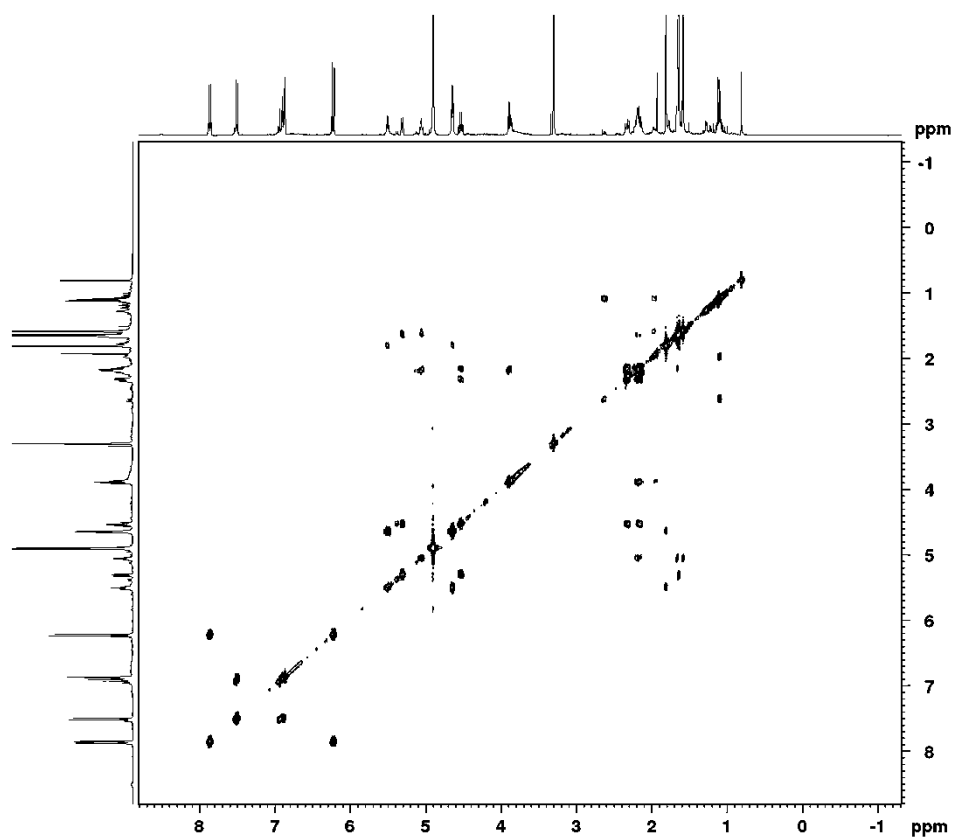


Figure 5.27 COSY spectrum of compound 4

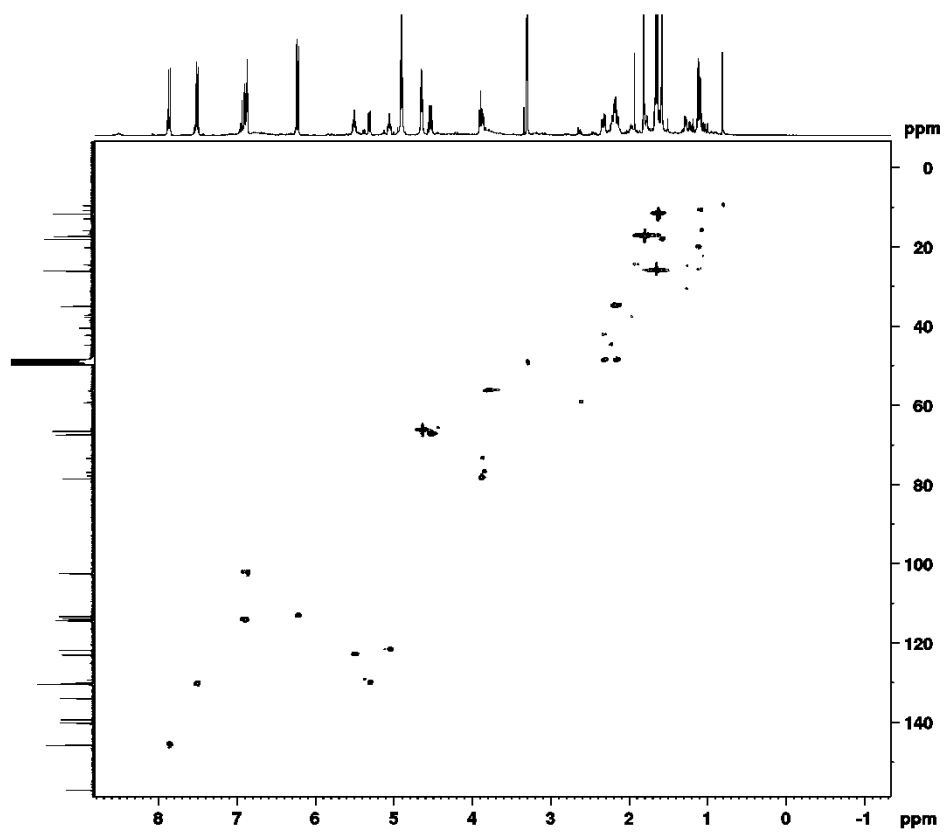


Figure 5.28 HSQC spectrum of compound 4

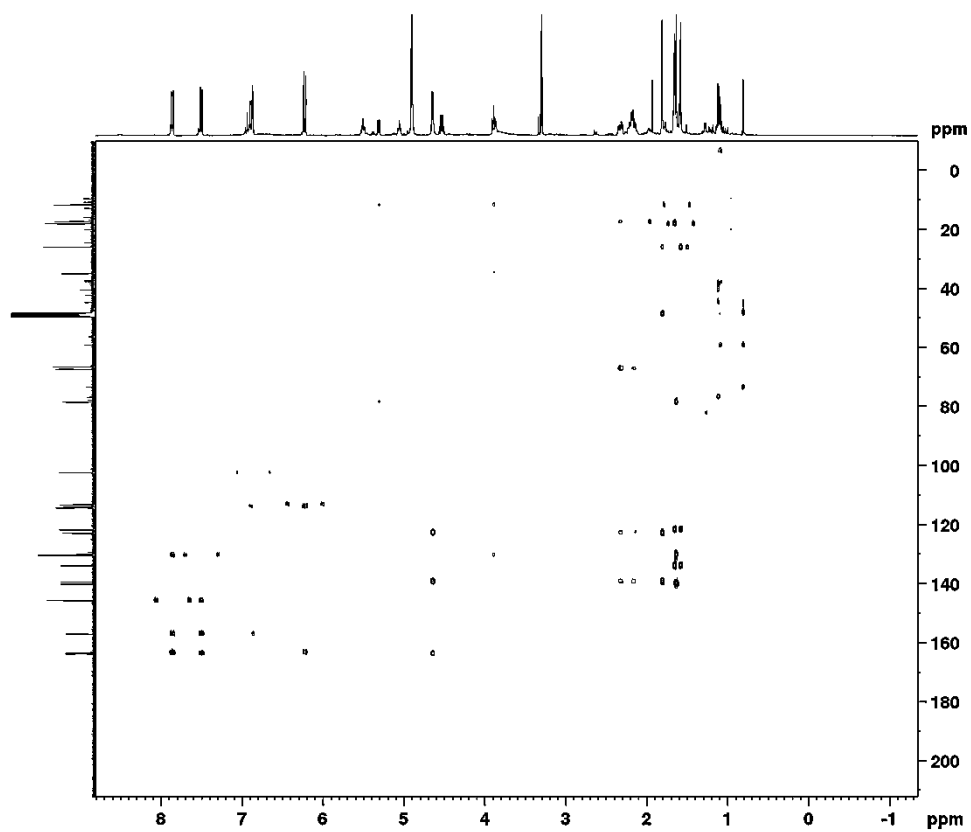


Figure 5.29 HMBC spectrum of compound 4

5.3.5 10'R-Karatavacinol (5)

The structure of compound **5** was elucidated by $^1\text{H-NMR}$, $^{13}\text{C-NMR}$ (DEPT-135 and DEPT-90) as 10'R-karatavacinol (Fig. 5.30).

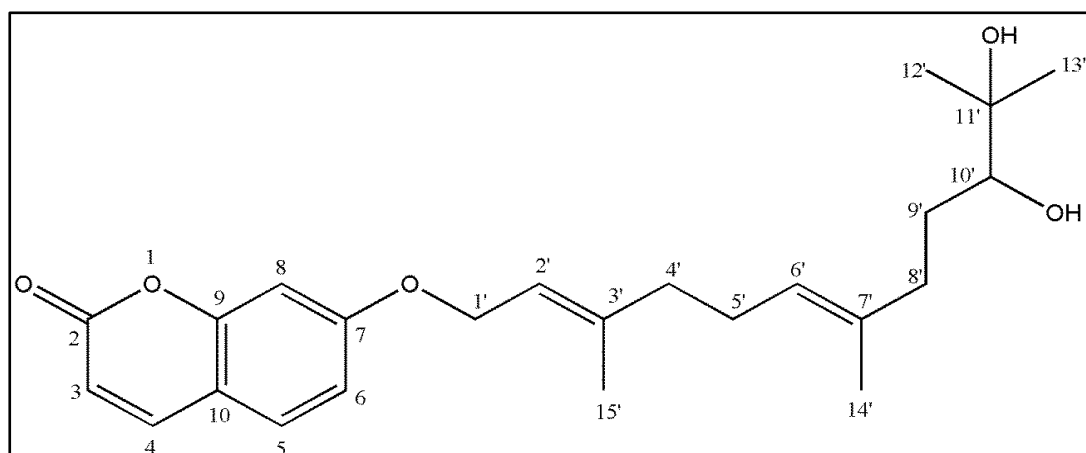


Figure 5.30 Structure of 10'R-karatavacinol (5)

Library search on the $^{13}\text{C-NMR}$ data (Table 5.6 and Fig. 5.31) using the NMR Predict database and $^1\text{H-NMR}$ (Table 5.6 and Fig. 5.32) suggested that this compound was a substituted coumarin with a terpene side chain. The spectral data were compared with NMR data from literature (Abd El-Razek *et al.*, 2007; Ahmad, 1999), which indicated that compound **5** was 10'R-karatavacinol.

In $^{13}\text{C-NMR}$ 24 signals were recorded. Indeed karatavacinol shared structural similarities with compounds **3** and **4**. However, five methylene signals were recorded in case of karatavacinol at C-1', C-5', C-6', C-8' and C-9' that corresponded to signals at 66.6 ppm, 30.8 ppm, 37.8 ppm, 40.5 ppm and 27.2 ppm, respectively. The carbonyl group appearing at 163.8 ppm was assigned to C-2 (carbonyl) of the coumarin moiety.

Similarly in the $^1\text{H-NMR}$ spectrum (Table 5.6 and Fig. 5.35) methylene signal appearing at 4.65 ppm (d; $J = 6.0$) was assigned to H-1'; whereas the multiplets appearing at 2.17 ppm, 1.20 ppm, 1.32 ppm and 1.70 ppm were assigned to H-4', H-5, H-8', H-9' in the

aliphatic chain, respectively. The upfield chemical shift of the signal at 3.21 ppm was indicative of an oxygenated CH at position 10'.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 211, 318 nm; ESI-MS (positive ion mode): m/z 423 [M+Na]⁺ consistent with a molecular formula C₂₄H₃₂O₅.

Karatavicinol has been isolated before from a number of *Ferula* species including *Ferula karatavica* (Kiryalov *et al.*, 1969), *Ferula sinica* (Ahmad, 1999), *Ferula foetida* (Abd EL-Razek *et al.*, 2007) and *Ferula sinkiangensis* (Teng *et al.*, 2013). This is however, the first time it is reported in *Ferula narthex*.

Table 5.6 ¹H and ¹³C NMR assignments for compound (5) recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
2		163.8
3	6.22; d; <i>J</i> = 9.5	113.8
4	7.90; d; <i>J</i> = 9.5	145.8
5	7.51; d; <i>J</i> = 8.5	130.3
6	6.91; dd; <i>J</i> = 8.0; 2.5	114.4
7		163.4
8	6.90; d; <i>J</i> = 2.5	102.5
9		157.0
10		113.2
1'	4.65; d; <i>J</i> = 6.0	66.6
2'	5.46; t; <i>J</i> =6.0	120.3
3'		142.8
4'	2.17; m	37.8
5'	1.20; m	30.8
6'	5.16; br t; <i>J</i> =6.0	125.0
7'		136.5
8'	1.32; m	40.5
9'	1.70; m	27.2
10'	3.21; dd; <i>J</i> =10.0; 2.0	77.7
11'		73.7
12'	1.10; s	25.0
13'	1.13; s	16.7
14'	1.61; s	25.6
15'	1.77; s	16.2

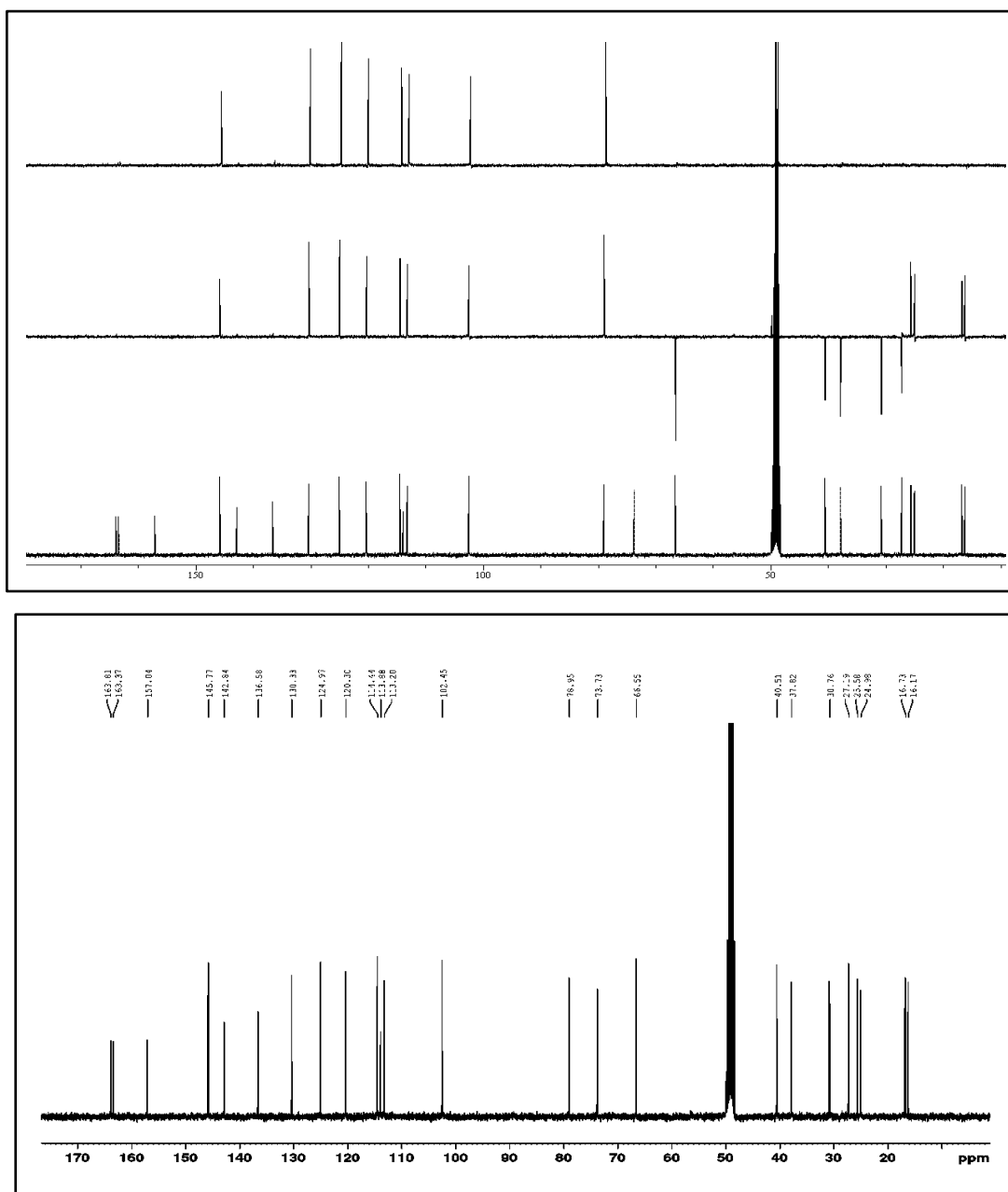


Figure 5.31 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of compound 5 in methanol- d_4 .

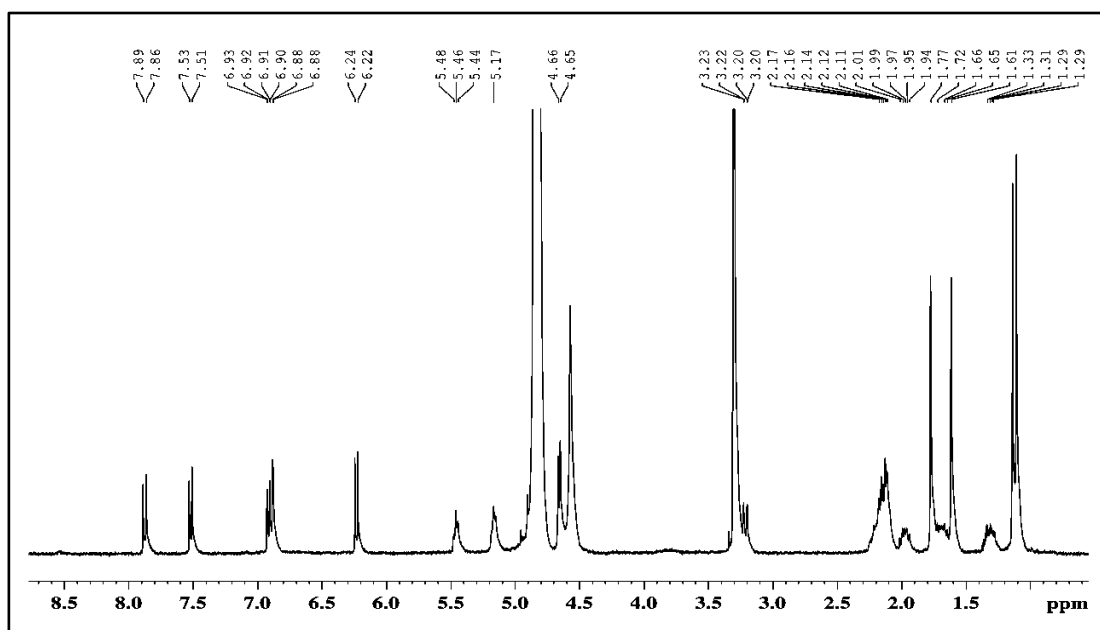


Figure 5.32 $^1\text{H-NMR}$ spectrum of compound **5** in methanol- d_4

5.3.6 10'*R*-Acetoxy-11'-hydroxyumbelliprenin (6)

The structure of compound **6** was elucidated by ¹H-NMR, ¹³C-NMR (DEPT-135 and DEPT-90) as 10'*R*-acetoxy-11'-hydroxyumbelliprenin (Fig. 5.33).

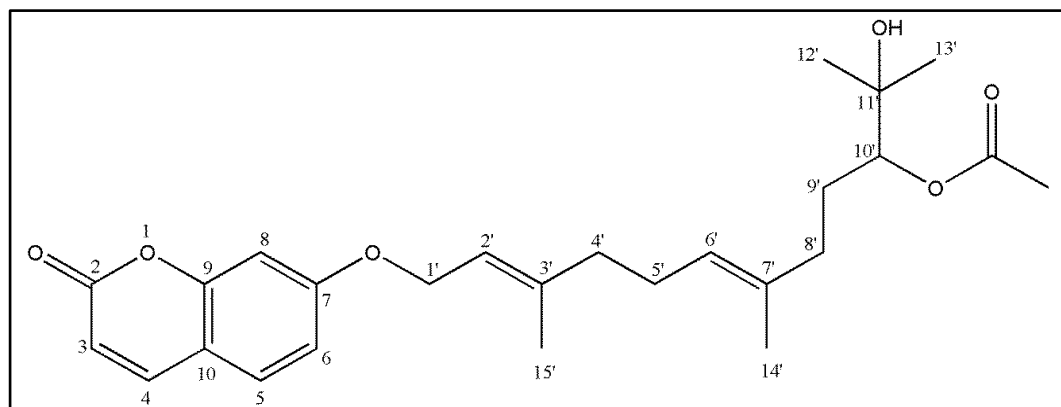


Figure 5.33 Structure of 10'*R*-acetoxy-11'-hydroxyumbelliprenin (**6**)

Library search on the ¹³C-NMR data (Table 5.7 and Fig. 5.34) using the NMR Predict database and ¹H-NMR (Table 5.7 and Fig. 5.35) suggested that this compound was a terpene-substituted coumarin. The spectral data were compared with NMR data from literature (Lee *et al.*, 2009), which indicated that compound **6** was 10'*R*-acetoxy-11' hydroxy-umbelliprenin.

In ¹³C-NMR 24 signals were recorded. Compound **6** showed structural resemblance with karatavacinol (**5**). Five methylene signals were recorded for C-1', C-4', C-5', C-8' and C-9' that could be assigned to the signals at 66.6 ppm, 40.4 ppm, 27.2 ppm, 37.4 ppm and 29.0 ppm, respectively. The carbonyl group appearing at 164.0 ppm was assigned to C-2 of the coumarin moiety. The signal at 172.8 ppm was assigned to the acetoxy group (carbonyl). Compound **6** is in fact the acetylated analogue of 10'*R*-karatavacinol, i.e 10'*R*-acetoxy-11' hydroxy-umbelliprenin.

In the ¹H-NMR spectrum, signals for the two main moieties i.e a coumarin and a sesquiterpene as in compounds **4** and **5**, were recorded. The sesquiterpene moiety showed signals for four methyl groups at 1.12 ppm, 1.13 ppm, 1.60 ppm and 1.77 ppm,

assigned to H-12', H-13', H-14' and H-15', respectively. The methylene signals observed at 2.05-2.16 (m) were assigned to H-4' and H-5', whereas multiplet signals at τ 1.92 ppm and 1.67 ppm were assigned to H-8' and H-9', respectively. The oxygenated methylene signal at 4.99 ppm was attributed to H-1'. The olefinic proton at H-2' was assignable to the signal at 5.50 ppm.

Yellow powder; UV (acetonitrile / H₂O) λ_{\max} 213, 321 nm; ESI-MS (positive ion mode): m/z 465 [M+Na]⁺ consistent with a molecular formula C₂₆H₃₄O₆.

10'-Acetoxy-11'-hydroxyumbelliprenin has been previously isolated from *asafoetida* (Lee *et al.*, 2009). This is, however, the first time it is reported in *Ferula narthex*.

Table 5.7 ¹H- and ¹³C-NMR assignments for compound **6** recorded in MeOH-d₄

Position	δ_{H} (ppm); multiplicity; <i>J</i> (Hz)	δ_{C} (ppm)
2		164.0
3	6.23; d; <i>J</i> = 9.5	113.2
4	7.90; d; <i>J</i> = 9.5	145.9
5	7.51; d; <i>J</i> = 8.5	130.4
6	6.91; dd; <i>J</i> = 8.0; 2.5	114.5
7		163.5
8	6.90; d; <i>J</i> = 2.5	102.5
9		157.1
10		113.9
1'	4.66; d; <i>J</i> = 6.0	66.6
2'	5.50; t; <i>J</i> =6.0	120.5
3'		143.0
4'	2.16-2.05; m	40.4
5'	2.16-2.05; m	27.2
6'	5.16; br t; <i>J</i> =6.0	125.5
7'		135.7
8'	1.92; m	37.4
9'	1.67; m	29.0
10'	4.75; dd; <i>J</i> =10.5; 2.0	80.6
11'		72.8
12'	1.12; s	25.7
13'	1.13; s	25.9
14'	1.60; s	16.0
15'	1.77; s	16.7
OAc (C-10')	2.0; s	21.1
		172.8

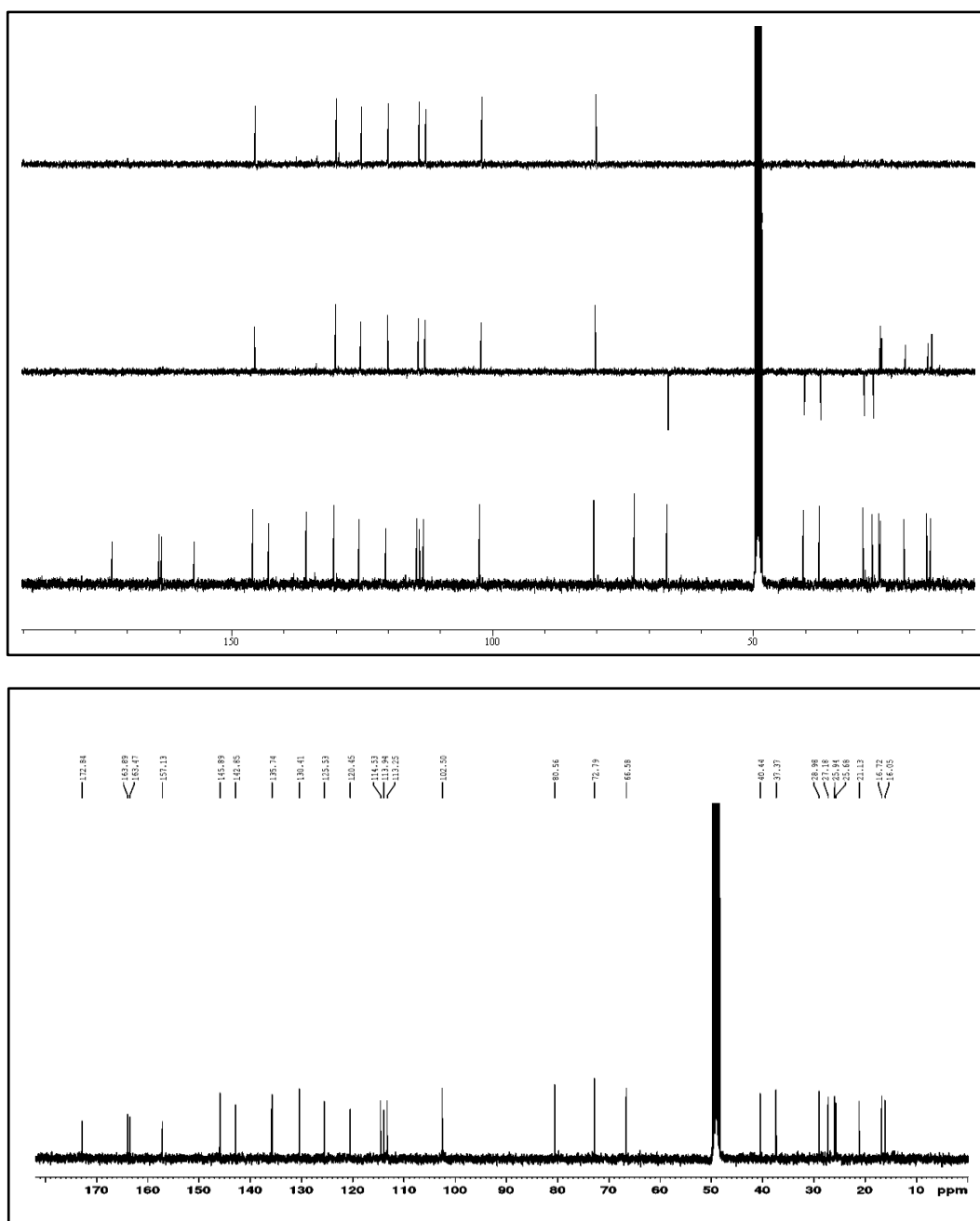


Figure 5.34 ^{13}C -NMR, DEPT-135 and DEPT-90 spectra of compound 6 in methanol- d_4 .

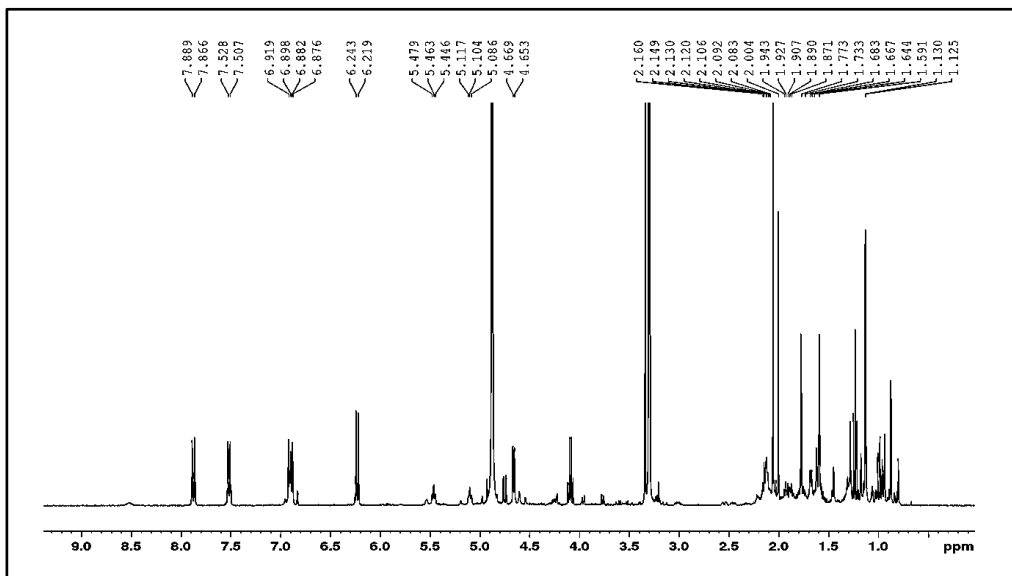


Figure 5.35 $^1\text{H-NMR}$ spectrum of compound **6** in methanol- d_4

5.4. BIOLOGICAL ACTIVITIES

5.4.1. Inhibition of AGEs formation

All fractions and crude extracts of *Ferula narthex* were evaluated for AGEs inhibition; however, due to practical issues of solubility and autofluorescence, it was not possible to obtain conclusive data.

All isolated compounds from *Ferula narthex* (**1-6**) were tested for their antiglycation potential in the BSA-Glucose and BSA-MGO models. Primarily in the BSA-glucose test, compound **2** displayed the highest activity (IC₅₀ 0.414 mM), being more active than the positive control aminoguanidine, followed by compound **4** (IC₅₀ 1.83 mM), whereas compounds **1** (47 % inhibition at the highest test concentration of 2 mM), **3** (44 % inhibition), **5** (32 % inhibition) and **6** (36% inhibition) showed moderate inhibition (Table 5.9). On the other hand, in the BSA-MGO assay, the highest activity was shown by compound **2** (IC₅₀ 1.03 mM), followed by **1** (IC₅₀ 1.71 mM) and **3** (IC₅₀ 1.03 mM), whereas moderate activities were observed for compounds **2** (40% inhibition at the highest test concentration of 2 mM), **4** (35% inhibition) and **6** (20% inhibition). Therefore it could be concluded that the antiglycation potential of the isolated constituents was due to both oxidative and non-oxidative modes of inhibition.

Table 5.9. Antiglycation (inhibition of AGEs) potential of isolated compounds from *Ferula narthex*

Compound	Protein glycation			
	BSA-Glucose		BSA-MGO	
	% inhibition ^a	IC ₅₀ mM	% inhibition ^b	IC ₅₀ mM
1	47			1.71
2	-	0.414	40	-
3	44			1.03
4		1.83	35	-
5	32		-	1.86
6	36		20	
Aminoguanidine		1.75		0.15
Quercetin		0.23		-

^a at 0.25 mM final concentration

^b at 2 mM final concentration

5.4.2 Antimicrobial activity

The crude extract and fractions were evaluated for antibacterial and antifungal activities. Highest levels of antimicrobial and antifungal activities were observed for the methanolic fraction, i.e. an IC₅₀ of 19.9 µg/mL against *S. aureus* and 22.0 µg/mL against *M. canis*, whereas this fraction was also found to be cytotoxic towards MRC-5 cells (IC₅₀ 36.5 µg/mL). Similarly, the *n*-hexane fraction was active (IC₅₀ 8.3 µg/mL) against *M. canis*, whereas the chloroform fraction was cytotoxic (IC₅₀ 49.8 µg/mL) against MRC-5 cells (Table 5.8).

Table 5.8 Antibacterial, antifungal and cytotoxic activities of *Ferula narthex* fractions.

Sample	IC ₅₀ µg/mL				
	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>M. canis</i>	MRC-5
MeOH 90%	19.9	>64	>64	22.0	36.5
Chloroform	>64	>64	>64	>64	49.8
Ethyl acetate	>64	>64	>64	>64	>64
<i>n</i> -Hexane	>64	>64	>64	8.3	>64
<i>n</i> -Butanol	>64	>64	>64	>64	>64
Aqueous	>64	>64	>64	>64	>64

Reference: Tamoxifen(MRC-5) IC₅₀ 11.4 µg/mL; erythromycin (*S. aureus*)IC₅₀ 11.2 µg/mL, Trimethoprim (*E. coli*) IC₅₀ 0.25 µg/mL; Miconazole(*C. albicans*) IC₅₀ 5.99 µg/mL; Terbinafin (*M. canis*) IC₅₀ 0.11 µg/mL.

MRC-5: Human fetal lung fibroblasts; *S. aureus*: *Staphylococcus aureus*; *E. coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*; *M. canis*: *Microsporom canis*.

The isolated compounds **1**, **3**, **4** and **6** were evaluated in an integrated screening panel for antimicrobial activity (Table 5.10). Compounds **2** and **6** were not tested due to the limited amount available. Although the methanolic fraction showed activity in the preliminary screening (Table 5.8), however none of the tested compounds presented anti-bacterial (*Staphylococcus aureus*, *Escherichia coli*) or antifungal activities (*Candida albicans*, *Aspergillus fumigatus*) in the tested range. It could be assumed that the individual compounds may act synergistically in the crude fractions that were found active. A few investigations have reported anti-bacterial properties of sesquiterpenes (Al-Yahya *et al.*, 1998; Iric *et al.*, 2012; Dastan *et al.*,2016); however, such activity is

surely related to certain structural variations that may not be met here. Also, it could be emphasized that such compounds may act as prodrugs and therefore could get activated in gastrointestinal tract and/or after absorption.

Compound **1** (fnarthexol) was moderately active against *Plasmodium falciparum* K1 (IC₅₀ 22.4 μM), whereas equal levels of inhibition were observed for *Trypanosoma brucei* (IC₅₀ 8.1 μM), *Trypanosoma cruzi* (IC₅₀ 8.6 μM) and *Leishmania infantum* (IC₅₀ 6.8.0 μM), accompanied however by PMM cytotoxicity (IC₅₀ 8 μM).

Structurally similar compounds (**3**, **4**, and **5**) also were active in the antiprotozoal assays. Compound **4** was found most active against *Plasmodium falciparum* K1 (IC₅₀ 1.3 μM), while a moderate activity against *Trypanosoma brucei* (IC₅₀ 32.6 μM), *Trypanosoma cruzi* (IC₅₀ 10.5 μM) and *Leishmania infantum* (IC₅₀ 12.7 μM) that was accompanied however by PMM cytotoxicity (IC₅₀ 32.0 μM). Compound **3** also displayed interesting activity against *Plasmodium falciparum* K1 (IC₅₀ 7.4 μM), with a moderate activity against *Trypanosoma brucei* (IC₅₀ 32.4 μM), *Trypanosoma cruzi* (IC₅₀ 19.1 μM) and *Leishmania infantum* (IC₅₀ 12.7 μM) that was accompanied by PMM cytotoxicity (IC₅₀ 32.0 μM).

Similarly compound **5** presented activity against *Plasmodium falciparum* K1 (IC₅₀ 16.0 μM), while a moderate activity was observed against *Trypanosoma brucei* (IC₅₀ 32.4 μM), *Trypanosoma cruzi* (IC₅₀ 9.4 μM) and *Leishmania infantum* (IC₅₀ 32.4 μM) that was accompanied by PMM cytotoxicity (IC₅₀ 32.0 μM). All tested compounds demonstrated mild cytotoxicity against MRC-5 cells.

Table 5.10. Antimicrobial, antifungal cytotoxic and antiprotozoal activities of isolated compounds from *Ferula narthex* (IC₅₀, μM)

Compound	IC ₅₀ (μM)									
	MRC5	Pf-K1	<i>T. bruc</i>	<i>T. cruz</i>	<i>L. inf</i>	PMM	<i>S. aureus</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>A. fumigatus</i>
1	8.0	22.4	8.1	8.6	6.8	8	> 64	> 64	> 64	> 64
2	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT
3	31.7	7.4	32.4	19.1	12.7	32	> 64	> 64	> 64	> 64
4	11.7	1.3	32.6	10.5	12.7	32	> 64	> 64	> 64	> 64
5	20.4	16.0	32.4	9.4	32.4	32	> 64	> 64	> 64	> 64
6	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT

Reference: Tamoxifen (MRC-5), IC₅₀ 10.48 μM; Suramine (*T. bruc*), IC₅₀ 0.03 μM, Chloroquine (*Pf-K1*), IC₅₀ 0.08 μM, Miltefosine (*L. inf*), IC₅₀ 9.02 μM; Benznidazol (*T. cruz*), IC₅₀ 2.13 μM; Erythromycin (*S. aureus*) IC₅₀ 11.30 μM; Chloramphenicol (*E. coli*), IC₅₀ 2.42 μM; Miconazole (*C. albicans*), IC₅₀ 4.70 μM; Terbinafine (*A. fumigatus*), IC₅₀ 1.38 μM.

MRC-5: human fetal lung fibroblasts; Pf-K1; *Plasmodium falciparum*; *T. bruc*: *Trypanosoma brucei*; *T. cruz*: *Trypanosoma cruzi*; *L. inf*: *Leishmania infantum*; PMM: peritoneal murine macrophages; *S. aureus*: *Staphylococcus aureus*; *E. coli*: *Escherichia coli*; *C. albicans*: *Candida albicans*; *A. fumigatus*: *Aspergillus fumigatus*

5.5. SUMMARY AND CONCLUSION

Ferula narthex Boiss. is a well-recognised medicinal plant in Pakistan. Its milky exudate obtained from bark is commonly used as a spice in everyday food and also as part of certain folk medicines in the subcontinent, such as “Unmadnashak Ghrita”. Traditionally it is used to treat a number of ailments including antimicrobial infections and diabetes mellitus. Therefore we performed a detailed phytochemical and biological investigation on the exudate of *Ferula narthex* and have isolated a series of compounds. The isolated compounds were analysed for their antiglycation properties (inhibition of Advanced Glycation Endproducts-AGEs), and antimicrobial activities, as part of running programmes in the host laboratory.

A total of 6 compounds were isolated (sesquiterpene- substituted coumarins) including fnarthexol (**1**), ligupersin A (**2**), 8-acetoxy-5-hydroxyumbelliprenin (**3**), asacoumarin A (**4**), 10' R- karatavicinol (**5**) and 10'-acetoxy-11'-hydroxyumbelliprenin (**6**).

In the antiglycation assays (inhibition of AGEs formation) compound **2** displayed the highest activity (IC₅₀ 0.414 mM) in the BSA-glucose assay, followed by compound **4** (IC₅₀ 1.83 mM). Compounds **2-6** presented moderate inhibition in the BSA-glucose assay. In the BSA-MGO assay, compound **3** presented the highest activity (IC₅₀ 1.03 mM), followed by compounds **1** (IC₅₀ 1.71 mM) and **5** (IC₅₀ 1.86 mM), indicating their potential to inhibit non-oxidative pathways of AGEs formation. Compounds **2**, **3** and **6** presented moderate inhibition in the same assay. It was evident that both oxidative and non-oxidative pathways contributed towards the antiglycation activity of isolated compounds.

In the antimicrobial assay for the crude extracts and fractions the methanolic fraction was considered as highly active, i.e. an IC₅₀ 19.9 µg/mL against *Staphylococcus aureus* and 22.0 µg/mL against *Microsporium canis*. However, this fraction was found to be cytotoxic towards MRC-5 cells (IC₅₀ 36.5 µg/mL). Similarly, the *n*-hexane fraction was active (IC₅₀ 8.3 µg/mL) against *Microsporium canis*, whereas the chloroform fraction was cytotoxic (IC₅₀ 49.8 µg/mL) against MRC-5 cells.

Nevertheless in the antimicrobial screening of isolated constituents, none of the tested compounds demonstrated anti-bacterial (*Staphylococcus aureus*, *Escherichia coli*) or antifungal activities (*Candida albicans*, *Aspergillus fumigatus*) in the test range. However, all tested compounds (**1**, **3**, **4** and **5**) were found active in the antiprotozoal and antileishmanial assays. Compound **1** was active against *Plasmodium falciparum* K1 (IC₅₀ 22.4 μM), whereas equivalent inhibitions were recorded for *Trypanosoma brucei* (IC₅₀ 8.1 μM), *Trypanosoma cruzi* (IC₅₀ 8.6 μM) and *Leishmania infantum* (IC₅₀ 8.0 μM). Among compounds **3**, **4** and **6** that are structurally similar, compound **4** was the most active one against *Plasmodium falciparum* K1 (IC₅₀ 1.3 μM), while moderate activities were observed against other protozoal organisms.

Based on findings of the current investigation, it could be hypothesized that the isolated constituents may contribute in part to the traditional use of *Ferula narthex* in diabetic conditions and for antimicrobial applications.

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CHAPTER 6
GENERAL CONCLUSIONS AND FUTURE PERSPECTIVES

During the current project three medicinal plants from Pakistan were investigated for their phytochemical constituents and biological activities. These medicinal plants were selected based on the fact that they were used traditionally for the treatment of diabetes mellitus and microbial infections by local people. Also the traditional healers use them in formulations which are based on the “Ayurveda”, “Unani” or “Tib” medicinal systems of the Indian subcontinent.

From *Nymphoides indica* leaves extracts a series of seco-iridoids and flavonoids was isolated. All crude extracts, fractions and compounds were tested for a number of biological activities. We were able to conclude that the extracts possessed moderate antimicrobial and antidiabetic activities. It was emphasised that the phenomenon of synergism and of metabolic conversions of constituents after oral intake may explain its traditional medicinal use, in spite of rather moderate *in vitro* activities.

For *Kickxia ramosissima*, we were able to isolate some iridoids and flavonoids, which were tested for various biological activities. It could be concluded that the crude extracts, fractions and isolated compounds presented some promising activities including antidiabetic effects, for which the plant is known. Similarly to *N. indica*, synergistic effects and the occurrence of many glycosylated compounds that are metabolised after oral intake could possibly play a key role in the pharmacological activity of *Kickxia ramosissima*.

Finally some coumarin-sesquiterpenes were isolated from the *Ferula narthex* exudate. Based on our findings, it was hypothesized that the isolated constituents may contribute in part to the traditional use of *Ferula narthex* in diabetic conditions and for antimicrobial applications. As we were able to present some antiprotozoal and anti-AGEs compounds, further detailed investigations are suggested for *in vivo* blood glucose lowering effects and a broader screening for antimicrobial analysis.

Although no highly active constituents have been identified, this study has contributed to the phytochemical and biological knowledge of three medicinal plants widely used in Pakistan. The results of the present investigation may in part support their traditional use against microbial infections and/or in diabetic conditions. Especially for

Nymphoides indica and *Kickxia ramosissima*, the possibility that prodrugs such as iridoid- and flavonoid glycosides, showing weak in vitro activities, are deglycosylated and metabolically activated after oral administration, cannot be excluded, and should be investigated in more detail. As indeed *in vitro* assays cannot really mimic *in vivo* situations, we suggest further *in vivo* trials of isolated constituents and the crude extracts. Based on the constituents isolated in the course of this project, these extracts can now fully be characterised, which is a necessary condition for performing reliable and reproducible *in vivo* assays.

As the traditional medicinal therapy is mainly based on observations obtained on human beings by traditional healers, it would be very interesting if the selected plants extracts were further evaluated in controlled clinical trials.

As traditionally these plant species are mostly used in formulations, it would also be very interesting to explore the other plants of these formulations individually. The comparison of the *in vivo* evaluation of the individual plants of such formulation and that of the total formulation would give us a better understanding of the therapy and more comprehensive conclusions could be made.

SUMMARY

Traditional medicine (TM) is an essential part of the health care system in the Indian subcontinent, including Pakistan. The roots of this system are as old as human civilization. Various forms of TM are practiced in this region including Ayurveda, Rigveda, Siddha, Tib and Yunani. These systems are either based on religious traditions or believes of super natural “healing powers” that are associated with the “earth's natural systems” that include medicinal plants and animal species, ambient healthy air and spring water. Even modern scientific techniques have confirmed the concept of personalised medicine, as it has been used for centuries in the traditional medicinal system of India. Irrespective of its scientific basis, religious believes, easy availability and effectiveness are the major reasons for acceptance of TM in this region.

This project focused on some medicinal plants traditionally used in diabetic conditions, and for antimicrobial applications, as part of running research programmes in the host laboratory.

Diabetes mellitus is a common chronic disorder that has become a challenging disease these days because of its continuously increasing prevalence world-wide, especially in developing nations. This growing prevalence is primarily due to bad life style, progressive urbanization, health status and social set up of the population. The chronic nature of the disease could lead to increased risk of severe complications, for instance kidney failure, blindness, cardiovascular diseases and foot ulcers. Diabetes mellitus is a multifarious metabolic disorder involving carbohydrate, fat and protein metabolism, that results from either insufficiency or dysfunction of insulin, ending up in elevated fasting and postprandial blood sugar levels.

Microbial infections are considered as major health issues in developing countries including Pakistan that constitute a major part of total annual death count. Poor hygiene, lack of proper health facilities and lack of knowledge are the major reasons for this widespread health concern in developing countries. Moreover, excessive and over-usage of antibiotics, leading to development of resistance has become critical.

Likewise, protozoal infections also contribute significantly to the total health burden in Pakistan. Geographical location, poor economic conditions, lack of health facilities and

heavy uncontrolled migration of Afghan refugees are major reasons for the widespread prevalence of protozoal infections in Pakistan. Cutaneous leishmaniasis (*Leishmania major*, *Leishmania tropica*) and malaria (*Plasmodium vivax* for 64% and *Plasmodium falciparum* for 26%) are main protozoal infection in Pakistan. Also, malaria accounts for 16% of the total disease burden of the country.

Although a range of drugs are available for the treatment of all such diseases, the cost of therapy, side effects and development of resistance in the cases of antimicrobial drugs, generally limit their use. Local people look towards traditional medicine for treatment of diabetes mellitus, microbial and protozoal infections. Keeping in mind the effectiveness and acceptance of TM, we selected some traditional medicinal plants and investigated their potential against some aspects of diabetes mellitus, and microbial infections.

Phytochemical investigations of *Nymphoides indica* extracts and fractions led to the isolation and identification of 15 compounds including 5 lipophilic compounds, i.e. the dicarboxylic acids azelaic acid (nonanedioic acid) and 4-methyl-heptanedioic acid, the monocarboxylic acids hexadecanoic acid and stearic acid and the fatty alcohol hexadecanol; 3 seco-iridoids, i.e. 7-epiexaltoside, 6",7"-dihydro-7-epiexaltoside (reported for the first time from nature) and menthiafolin; 3 flavonoids, i.e. 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside, 3-*O*-methylquercetin-7-*O*- β -glucoside and 3,7-di-*O*-methylquercetin; scopoletin and ferulic acid; and the monoterpenoids foliamenthic acid and 6,7-dihydrofoliamenthic acid methyl ester.

In the antidiabetic *in vitro* assays various fractions were assayed for potential antiglycation and α -glucosidase inhibitory activity. All tested fractions presented moderate inhibition (24-36% at a concentration of 100 μ g/mL) of the formation of AGEs. Similarly mild α -glucosidase inhibitory activity was observed for the total extract at 834 μ g/mL (13% inhibition), whereas the highest inhibition was shown by the methanolic (31% inhibition) and *n*-butanol fractions (25% inhibition) at the same concentration. At higher test concentrations the inhibitory activity did not further increase, and the 50% inhibition level could not be reached. Among isolated constituents interesting antiglycation activities were shown by 6",7"-dihydro-7-

epiexaltoside (IC₅₀ 0.36 mM), 3-*O*-methylquercetin-7-*O*-β-glucoside (IC₅₀ 0.42 mM) and 6,7-dihydrofoliamenthoic acid methyl ester (IC₅₀ 0.61 mM). Finally mild to moderate α-glucosidase inhibition was shown by 3,7-di-*O*-methylquercetin-4'-*O*-β-glucoside, 3-*O*-methylquercetin-7-*O*-β-glucoside and 3,7-di-*O* methylquercetin; scopoletin, ferulic acid, foliamenthoic acid and 6,7-dihydrofoliamenthoic acid methyl ester.

In the antimicrobial assays of the different fractions, the highest levels of inhibition were observed for the *n*-hexane fraction, i.e. an IC₅₀ of 19.5 μg/mL against *S. aureus* and 32.0 μg/mL against *M. canis*. Similarly the methanol 90% fraction was active (IC₅₀ 36.4 μg/ml) against *S. aureus* and cytotoxic (IC₅₀ 38.9 μg /mL) against MRC-5 cells. Among the isolated constituents the lipophilic compounds showed moderate antimicrobial activities, whereas 3,7-di-*O*-methylquercetin-4'-*O*-β-glucoside presented mild antiprotozoal activities against *Trypanosoma brucei* (IC₅₀ 8.0 μM), *Leishmania infantum* (IC₅₀ 32.0 μM) and *Trypanosoma cruzi* (IC₅₀ 30.0 μM). It was concluded that *N. indica* leaf extracts possess mild to moderate antimicrobial, antiprotozoal, antioxidant and antidiabetic activities. The traditional use of this plant species can possibly be due to synergism between its constituents, and to metabolic conversions (deglycosylations) of iridoids and flavonoid glycosides after oral intake.

The study on *Kickxia ramosissima* involved phytochemical and biological investigation of crude fractions and isolated constituents, including assessment of *in vitro* antidiabetic properties (inhibition of α-glucosidase activity; inhibition of Advanced Glycation Endproducts-AGEs), antioxidant activity (AAPH induced linoleic acid peroxidation), inhibition of 15-LOX (anti-inflammatory activity) and antimicrobial activities.

A total of 7 compounds were isolated including iridoids, i.e a new compound kickxiasine, mussaenosidic acid, mussaenoside, linarioside, the flavonoids pectolarigenin and pectolarin, and the benzoic acid derivative 4-hydroxy-benzoic acid methyl ester.

In the antidiabetic assays of crude extracts and fractions, the highest inhibition was recorded for the ethyl acetate fraction (IC₅₀ 88.0 μg/mL) whereas moderate levels of

antiglycation effects were observed for all other fractions. Also the isolated iridoids showed weak inhibition in the BSA-glucose assay, and comparable results in the BSA-MGO assay. The flavonoids, however, showed interesting results as pectolinarigenin was highly active (IC₅₀ 0.79 mM) followed by pectolinarigenin with IC₅₀ 2.29 mM. Also in the BSA-MGO assay both flavonoids showed high levels of inhibition i.e pectolinarigenin IC₅₀ 0.19 mM and pectolinarigenin IC₅₀ 0.13 mM. This trend clearly indicated the inhibition through non-oxidative pathways. The benzoic acid derivative did not present any activity in all AGEs experiments. In the α -glucosidase inhibition assay the iridoids presented a weak inhibition. However, with regard to the flavonoids, pectolinarigenin showed an interesting activity (IC₅₀ 0.23 mM), whereas moderate inhibition was observed for pectolinarin and the benzoic acid derivative.

In the 15-LOX experiment, moderate inhibition was recorded for pectolinarigenin, mussaenoside and kickxiasine. In the AAPH assay, weak or no inhibition was recorded for all compounds.

In the antimicrobial screening panel the *n*-hexane fraction presented activity against *Staphylococcus aureus* (IC₅₀ 8.0 μ g/mL) and *Microsporium canis* (IC₅₀ 24.4 μ g/mL), whereas none of the other fractions presented any activity in the test range. Among isolated constituents, only pectolinarigenin showed moderate activity against *Plasmodium falciparum K1* (IC₅₀ 41.8 μ M), *Staphylococcus aureus* (IC₅₀ 49.8 μ M) and *Trypanosoma cruzi* (IC₅₀ 30.0 μ M) with no cytotoxicity. Based on our findings we could demonstrate the moderate antidiabetic, anti-inflammatory and antimicrobial activity of some constituents of *Kickxia ramosissima*.

As mentioned before for *N. indica*, the traditional use of this plant species can possibly be due to synergism between its constituents, and to metabolic conversions (deglycosylations) of iridoids and flavonoid glycosides after oral intake.

During the phytochemical analysis of crude extracts of *Ferula narthex* exudate, 6 compounds (sesquiterpene-substituted coumarins) were isolated, including fnarthexol, ligupersin A, 8'-acetoxy-5'-hydroxyumbelliprenin, asacoumarin A, 10'*R*-karatavicinol and 10'*R*-acetoxy-11'-hydroxy umbelliprenin.

In the antidiabetic assays ligupersin A displayed the highest activity (IC_{50} 0.414 mM) in the BSA-Glucose assay, followed by asacoumarin A (IC_{50} 1.83 mM). In the BSA-MGO assay, 8'-acetoxy-5'-hydroxyumbelliprenin presented the highest activity (IC_{50} 1.03 mM), followed by feselol (IC_{50} 1.71 mM) and 10'*R*-karatavicinol (IC_{50} 1.86 mM). It was evident that both oxidative and non-oxidative pathways contributed towards the antiglycation activity of the isolated compounds.

In the antimicrobial assays the methanolic fraction was considered as highly active, i.e. an IC_{50} 19.9 $\mu\text{g/mL}$ against *Staphylococcus aureus* and IC_{50} 22.0 $\mu\text{g/mL}$ against *Microsporum canis*. However, this fraction was found cytotoxic towards MRC-5 cells (IC_{50} 36.5 $\mu\text{g/mL}$). Similarly the *n*-hexane fraction was active (IC_{50} 8.3 $\mu\text{g/mL}$) against *Microsporum canis*, whereas the chloroform fraction was found cytotoxic (IC_{50} 49.8 $\mu\text{g/mL}$) against MRC-5 cells.

In the antimicrobial screening of the isolated constituents, none of the isolated compounds demonstrated anti-bacterial or antifungal activities. However, moderate antiprotozoal and antileishmanial activities were noticed for all tested compounds.

Fnarthexol was moderately active against *Plasmodium falciparum* K1 (IC_{50} 22.4 μM), whereas equivalent inhibitions were recorded for *Trypanosoma brucei* (IC_{50} 8.1 μM), *Trypanosoma cruzi* (IC_{50} 8.6 μM) and *Leishmania infantum* (IC_{50} 8.0 μM). Among the other compounds that are structurally similar, asacoumarin A was the most active one against *Plasmodium falciparum* K1 (IC_{50} 1.3 μM), while moderate activities were observed against other protozoal organisms. Also mild cytotoxicity against MRC-5 cells was observed for all tested compounds.

Based on findings of the current investigation, it could be hypothesized that the isolated constituents may contribute in part to the traditional use of *Ferula narthex* in diabetic conditions and for antimicrobial applications.

SAMENVATTING

Traditionele geneeskunde is zo oud als de menselijke beschaving. De traditionele geneeskunde maakt een essentieel deel uit van de gezondheidszorg in het subcontinent Indië, waaronder Pakistan. In deze regio worden verschillende vormen van traditionele geneeskunde beoefend: de Ayurveda, Rigveda, Siddha, Tib en Yunani. Deze systemen zijn gebaseerd op religieuze tradities en geloof in bovennatuurlijke krachten, die geassocieerd zijn met aardse natuurlijke systemen, zoals medicinale planten en dieren, gezonde lucht en bronwater. Het concept van gepersonaliseerde geneeskunde, dat nu ook ingang vindt in de Westerse geneeskunde, wordt reeds vele eeuwen in de traditionele geneeskunde van Indië toegepast. Los van de wetenschappelijke basis, zijn religieuze opvattingen, beschikbaarheid en efficaciteit de voornaamste redenen voor de populariteit van traditionele geneeskunde.

In dit project staan enkele traditionele medicinale planten centraal, gebruikt bij diabetes en infectieziekten, als onderdeel van lopende onderzoeksprogramma's in deze domeinen in het gastlaboratorium. Diabetes mellitus is een veel voorkomende chronische ziekte, waarvan de prevalentie wereldwijd toeneemt, ook in ontwikkelingslanden. Deze toename is in de eerste plaats te wijten aan slechte voedingsgewoonten, voortschrijdende verstedelijking, de algemene gezondheidsstatus en sociale structuur van de bevolking. De chronische aard van deze aandoening kan leiden tot een verhoogd risico op complicaties, zoals nierfalen, blindheid, cardiovasculaire problemen en voetzweren. Ook infectieziekten anderzijds worden beschouwd als grote gezondheidsproblemen in ontwikkelingslanden, ook in Pakistan, die verantwoordelijk zijn voor een hoge jaarlijkse dodentol. Dit is voornamelijk te wijten aan slechte hygiëne, gebrek aan goede faciliteiten in de gezondheidszorg, en gebrek aan kennis over deze ziekten.

Ook protozoale infecties dragen aanzienlijk bij aan de totale ziektelast in Pakistan. Geografische ligging, de slechte economische omstandigheden, gebrek aan medische voorzieningen en zware ongecontroleerde migratie van Afghaanse vluchtelingen zijn belangrijke redenen voor de wijdverbreide prevalentie van protozoaire infecties in Pakistan. Cutane leishmaniasis (*Leishmania major*, *Leishmania tropica*) en malaria (*Plasmodium vivax* voor 64% en *Plasmodium falciparum* voor 26%) zijn de belangrijkste protozoaire infecties in Pakistan. Maartdaarme 16% van de totale ziektelast van het land uit.

Alhoewel heel wat geneesmiddelen beschikbaar zijn, vormen de kostprijs, de beschikbaarheid, de neveneffecten en in het geval van infectieziekten het ontstaan van resistentie een aantal beperkende factoren, waardoor de plaatselijke bevolking zijn toevlucht neemt tot de traditionele geneeskunde. Om deze reden werden enkele traditionele medicinale planten geselecteerd, met als doel hun potentiële bruikbaarheid na te gaan bij diabetes en bij infectieziekten.

Fytochemisch onderzoek van extracten van de bladeren van *Nymphoides indica* leidde tot de isolatie en identificatie van 15 inhoudsstoffen, waaronder 5 lipofiele producten (de dicarbonsuren azelainezuur (nonaandicarbonzuur) en 4-methyl-heptaandicarbonzuur; de monocarbonsuren palmitinezuur en stearinezuur; en het vetalcohol hexadecanol); 3 seco-iridoiden (7-epiexaltoside; 6",7"-dihydro-7-epiexaltoside, voor de eerste keer gerapporteerd; en menthiafoline); 3 flavonoiden (3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside; 3-*O*-methylquercetin-7-*O*- β -glucoside; en 3,7-di-*O* methylquercetin); scopoletine en ferulazuur; en de monoterpenen foliamenthoin zuur en 6,7-dihydrofoliamenthoin zuur methyl ester.

In de antidiabetische *in vitro* testen vertoonden alle geëvalueerde subfracties een matige activiteit tegen proteïn-glycatie (24-36% inhibitie van de vorming van Advanced Glycation Endproducts (AGEs) bij een concentratie van 100 $\mu\text{g}/\text{mL}$), en als inhibitor van α -glucosidase activiteit (tot 31% inhibitie bij een concentratie van 834 $\mu\text{g}/\text{mL}$ voor het methanolisch extract). Wat betreft de geïsoleerde producten werden voor de anti-glycatie werking, de hoogste activiteiten gevonden voor 6",7"-dihydro-7-epiexaltoside (IC_{50} 0.36 mM), 3-*O*-methylquercetin-7-*O*-glucoside (IC_{50} 0.42 mM) en 6,7-dihydrofoliamenthoin zuur methyl ester (IC_{50} 0.61 mM); terwijl 3,7-di-*O*-methylquercetin-4'-*O*- β -glucoside; 3-*O*-methylquercetin-7-*O*- β -glucoside; 3,7-di-*O* methylquercetin; scopoletine; ferulazuur; foliamenthoin zuur en 6,7-dihydrofoliamenthoin zuur methyl ester meest actief waren als inhibitor van α -glucosidase.

In de antimicrobiële testen was de n-hexaan fractie het meest actief, met een IC_{50} van 19.5 $\mu\text{g}/\text{mL}$ tegen *Staphylococcus aureus* en 32.0 $\mu\text{g}/\text{mL}$ tegen *Microsporium canis*. De methanol 90% fractie was actief (IC_{50} 36.4 $\mu\text{g}/\text{mL}$) tegen *S. aureus* en cytotoxisch (IC_{50} 38.9 $\mu\text{g}/\text{mL}$) op MRC-5 cellen. Wat betreft de geïsoleerde producten, vertoonden de lipofiele verbindingen een matige activiteit. 3,7-Di-*O*-methylquercetin-4'-*O*- β -

glucoside was matig actief tegen *Trypanosoma brucei* (IC₅₀ 8.0 µM), *Leishmania infantum* (IC₅₀ 32.0 µM) en *Trypanosoma cruzi* (IC₅₀ 30.0 µM).

Extracten van de bladeren van *N. indica* bezitten dus matige antimicrobiële, anti-glycatie en α-glucosidase inhiberende werking. Het traditioneel gebruik van deze plant kan mogelijk verklaard worden door synergisme tussen de inhoudsstoffen, of door metabole omzettingen (deglycosylaties) en activatie van de seco-iridoiden en flavonoiden na orale inname.

Uit *Kickxia ramosissima* werden 7 inhoudsstoffen geïsoleerd, meer bepaald de iridoiden kickxiasine (een nieuw product), mussaenosidine zuur, mussaenoside en linarioside; de flavonoiden pectolinarigenine and pectolinarine; en 4-hydroxymethylbenzoaat.

De hoogste anti-glycatie activiteit werd vastgesteld voor de ethyl acetaat fractie (IC₅₀ 88.0 µg/mL); ook de geïsoleerde iridoiden vertoonden een matige activiteit. De flavonoiden pectolinarigenine (IC₅₀ 0.79 mM) en pectolinarine (IC₅₀ 2.29 mM) waren actiever. Ook in BSA-MGO test waren deze flavonoiden zeer actief (IC₅₀ 0.19 mM en 0.13 mM, respectievelijk), wat aangeeft dat ook inhibitie van de niet-oxidatieve wegen betrokken is bij de werking. In de α-glucosidase inhibitie test waren de iridoiden slechts zwak actief, in tegenstelling tot de flavonoiden, met een IC₅₀ waarde van 0.23 mM voor pectolinarigenine.

In de antimicrobiële screening was de *n*-hexaan fractie actief tegen *Staphylococcus aureus* (IC₅₀ 8.0 µg/mL) en *Microsporum canis* (IC₅₀ 24.4 µg/mL). Enkel pectolinarigenine was matig actief tegen *Trypanosoma cruzi* (IC₅₀ 30.0 µM), *Plasmodium falciparum* K1 (IC₅₀ 41.8 µM) en *Staphylococcus aureus* (IC₅₀ 49.8 µM), in de afwezigheid van cytotoxiciteit.

Zoals reeds vermeld voor *N. indica*, kan het traditioneel gebruik van *K. ramosissima* mogelijk verklaard worden door synergisme tussen de inhoudsstoffen, of door metabole omzettingen (deglycosylaties) en activatie van de iridoiden en flavonoiden na orale inname.

Uit het exudaat van *Ferula narthex* tenslotte werden 6 bestanddelen verkregen, behorend tot de sesquiterpeen-gesubstitueerde coumarines, namelijk fnarthexol, ligupersine A, 8'-acetoxy-5'-hydroxyumbelliprenine, asacoumarine A, 10'-R-karatavicinol

en 10'R-acetoxy-11'-hydroxy-umbelliprenine. Ligupersine A vertoonde de hoogste anti-glycatie activiteit (IC₅₀ 0.414 mM), gevolgd door asacoumarine A (IC₅₀ 1.83 mM). In de BSA-MGO test was 8'-acetoxy-5'-hydroxyumbelliprenine het meest actief (IC₅₀ 1.03 mM), gevolgd door fnarthexol (IC₅₀ 1.71 mM) en 10'R-karatavicinol (IC₅₀ 1.86 mM). Zowel inhibitie van de oxidatieve als de niet-oxidatieve weg droeg dus bij tot de activiteit. De methanolische fractie vertoonde een hoge antimicrobiële activiteit (IC₅₀ 19.9 µg/mL tegen *Staphylococcus aureus* en IC₅₀ 22.0 µg/mL tegen *Microsporum canis*), maar was ook cytotoxisch tegen MRC-5 cellen (IC₅₀ 36.5 µg/mL). The *n*-hexane fractie was actief (IC₅₀ 8.3 µg/mL) tegen *Microsporum canis*. *Fnardhexol* was matig actief tegen *Plasmodium falciparum* K1 (IC₅₀ 22.4 µM), *Trypanosoma brucei* (IC₅₀ 8.1 µM), *Trypanosoma cruzi* (IC₅₀ 8.6 µM) en *Leishmania infantum* (IC₅₀ 8.0 µM). Asacoumarine A echter was zeer actief tegen *Plasmodium falciparum* K1 (IC₅₀ 1.3 µM). De geïsoleerde producten kunnen dus ten minste voor een deel het traditioneel gebruik verklaren.

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