

Phytochemical and Antimicrobial Studies on Selected
Medicinal Plants from The Iraqi Flora: *Citrus grandis*,
Citrus sinensis and *Ruta chalepensis*

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Dedicated To

My greatest mother, the reason of what I have become today.

Thanks for your great support and continuous care.

My husband, Mohammed, who supports me to achieve my dream. Thank you for your love, wisdom and timeless support.

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“NEVER GIVE UP”

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Abstract

This study was founded on traditional medicinal information and local traditional uses of three Iraqi medicinal plants from the Rutaceae family, namely, *Citrus grandis* (leaves), *Citrus sinensis* (leaves and peel) and *Ruta chalepensis* (fruits, leaves, stems and roots). A bioassay-guided approach was adopted to study the phytochemicals and antimicrobial activity of these plants. The *n*-hexane, DCM and MeOH extracts of all plants and plant parts, obtained from successive Soxhlet extraction, were assessed for their antimicrobial properties using the microtitre resazurin assay involving two Gram-positive bacterial strains, *Micrococcus luteus* and *Staphylococcus aureus*, two Gram-negative strains, *Escherichia coli* and *Pseudomonas aeruginosa* and one fungal strain, *Candida albicans*. The DCM extracts of *C. grandis* (leaves) and *C. sinensis* (leaves) showed notable antimicrobial activity against the microbes used in this study with different MIC values. Both DCM and MeOH extracts of *C. sinensis* (peel) also revealed significant activity. All *R. chalepensis* extracts exhibited antimicrobial activity except for the root extract, which did not show any antimicrobial activity with the maximum concentration of 10 mg/mL. Thirty three compounds including two novel ones, mainly coumarins and flavonoids, were isolated from the active extracts/fractions using a combination of thin layer chromatography (TLC), solid-phase extraction and reversed-phase high-performance liquid chromatography (HPLC). 1D and 2D nuclear magnetic resonance (NMR) and mass spectrometric techniques were employed to identify and confirm the structures of the isolated compounds. Twenty of 33 isolated compounds were evaluated for their anti-MRSA activity against five methicillin-resistant *Staphylococcus aureus* strains (SA1199B, XU212, MRSA340702, EMRSA-15MRSA274819) and the standard strain (ATCC25923). The mechanism of action of six compounds were investigated by scanning electronic microscopic (SEM) method to detect any morphological changes or damage caused by the tested compounds.

This study generated the first phytochemical and antimicrobial report of *R. chalepensis* fruits and Iraqi *C. grandis* leaves. To study the mechanism of action of compounds selected from the resazurin assay, a SEM protocol was developed.

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List of abbreviations

δ	Chemical shift (ppm)
$^1\text{H-NMR}$	Proton nuclear magnetic resonance
$^{13}\text{C-NMR}$	Carbon nuclear magnetic resonance
1D	One dimensional
2D	Two dimensional
ACN	Acetonitrile
ATCC	American Type Cell Culture Collection
BC	Before Christ
CD_3OD	Methanol deuterated
CDCl_3	Deuterated chloroform
cfu	Colony-forming unit
cm	Centimeter
COSY	Correlation spectroscopy
d	Doublet
DCM	Dichloromethane
DEPT	Distortionless Enhancement by Polarisation Transfer
DMSO	Dimethyl sulfoxide
DMSO-d_6	Deuterated dimethyl sulfoxide
EtOAc	Ethyl acetate
FDA	Food and Drug Administration
g	Gram
h	Hour
HRESIMS	High-resolution electrospray ionisation mass spectrometry
HMBC	Heteronuclear multiple bond coherence
HPLC	High performance liquid chromatography
HSQC	Heteronuclear single quantum coherence
Hz	Hertz
IC_{50}	Concentration needed to produce 50% of cells inhibition

ICU	Intensive care unit
J	Coupling constant
km	Kilometre
m	Multiplet (for NMR spectrum)
m/z	Mass to charge ratio
MeOH	Methanol
mg	Milligram
MHz	Mega hertz
MIC	Minimum inhibitory concentration
min	Minute
mL	Milliliter
μL	Microliter
MS	Mass spectroscopy
MRSA	multidrug-resistant <i>Staphylococcus aureus</i>
MSSA	methicillin-susceptible <i>Staphylococcus aureus</i>
MHB	Mueller-Hinton broth
MTT	3-(4,5-dimethylthiazo-2-yl)-2,5-diphenyltetrazolium
N/A	No activity
NCTC	National Collection types of Culture
NMR	Nuclear magnetic resonance
ppm	Parts per million
prep-HPLC	Preparative high performance liquid chromatography
R_f	Retention factor
PTLC	Preparative thin layer chromatography
s	Singlet
SEM	Scanning electron microscope
Semi prep-HPLC	Semi preparative high performance liquid chromatography
SPE	Solid phase extraction
t	Triplet
t_R	Retention time

TFA	Trifluoro acetic acid
TLC	Thin layer chromatography
μg	Microgram
μL	Microliter
μm	Micromole
UV	Ultraviolet
UV-DAD	Ultraviolet-diode array detector
VLC	Vacuum liquid chromatography

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Chapter 1
Introduction

1.1. Microbial resistance to antibiotics

Antibiotic resistance is a global problem and is most common in developing countries (Livermore, 2003). According to the World Health Organisation (WHO), every year microbial resistance to antibiotics leads to the deaths of more than 60,000 people worldwide, out of which 77% are children (Moran, 2005).

Microorganisms, particularly bacteria, develop resistance to standard antimicrobial drugs mainly because of clinical, cellular and molecular factors. Instances of misuse, over-prescription of antibiotics have become common practices in developing countries (Burroughs *et al.*, 2003). The unlicensed medicines suppliers, uncontrolled antibiotic sales, and availability of over the counter antibiotics without a prescription, has led to an exponential increase in drug resistance (Ayukekbong *et al.*, 2017). Hospitals are the place of emergence for resistant strains. In the 1930s, military hospitals had reported that *Streptococcus pyogenes* showed resistance to sulphonamide (Levy, 1982). Later in the 1940s, a civilian hospital in London found *Staphylococcus aureus* was resistant to penicillin (Levy and Marshall, 2004) and by 1950, this strain became widespread. In the late 1950s, a new type of microorganism *Shigella dysenteriae* appeared in Japan, which was resistant to antibiotics like chloramphenicol, streptomycin, tetracycline, and sulphonamides (Levy, 1982). In 1972 and 1973, another multidrug-resistant strain of the same bacterial species was identified in Brazil and Vietnam (Goldstein *et al.*, 1986). Generally, every bacterial species has different antibiotic resistance specificity.

Resistant strains increase mortality, morbidity, and cost of treatment (Cosgrove, 2006). In England, successful execution of approximately 4,000,000 operations solely depends on antibiotics, especially for caesarean sections and for women suffering from urinary tract infections (Llor and Bjerrum, 2014; Shallcross *et al.*, 2015). Furthermore, some women need antibiotics during their gestational period to prevent the passage of infection from her body to the newborn child (WHO, 2014). Antimicrobial resistance is causing

hindrance in treatment, and many antibiotics often come with various side effects. Therefore, there is an urgent need to develop new effective antibiotics with fewer side effects and at a lower cost.

1.2. Traditional uses of medicinal plants to treat infections

Medicinal plants have been used globally by different civilisations for many centuries to treat diseases, including microbial infections (Ramawat *et al.*, 2009). The Sumerians, from Mesopotamia (Iraq now), in 2600 BC, first mentioned the use of medicinal herbs by writing plant recipes on clay scrolls (Ramawat *et al.*, 2009; Sarker and Nahar, 2012). In most communities, plants play a vital role as food or medicine. Most societies have extensive traditional knowledge of plant usage. However, if these uses are not well documented, successive generations will lose these pieces of information (Hostettmann *et al.*, 2000; Houghton and Mukherjee, 2009). Approximately 400,000 flowering plants have been recorded globally, of which more than 50,000 have been used as medicinal plants (Verpoorte, 2009). These medicinal plants have been used to treat different types of health problems and some probably have an impact on protection from diseases. In addition, the high content of mineral nutrients may enhance the ability of plants to boost the immune system (Kubena and McMurray, 1996).

People of developing countries tend to be reported with more cases of infectious diseases due to their life style and often poor living conditions. Their suffering has been exacerbated by infections caused by drug-resistant pathogens and the high cost of treatment (Mondol and Shin, 2014). According to WHO (2008), more than 80% of people in Asia and Africa use medicinal plants as the first choice for their healthcare.

People in India use plants such as, *Achyranthes aspera*, *Allium cepa*, *Allium sativum*, *Aloe vera*, *Anacardium occidentale*, *Areca catechu*, *Azadirachta indica*, *Bauhinia variegata*, *Beta vulgaris*, *Brassica oleracea*, *Calendula officinalis*, *Calotropis gigantea*, *Camellia sinensis*, *Cannabis sativa*, *Cassia fistula*, *Cissampelos pareira*, *Crocus sativus*, *Cleome*

viscosa, *Costus speciosus*, *Curcuma longa*, *Daucus carota*, *Echinacea angustifolia*, *Eucalyptus globulus*, *Euphorbia spp.*, *Ficus benghalensis*, *Ficus racemosa*, *Ixora coccinea*, *Jatropha gossypifolia*, *Lavandula officinalis*, *Lawsonia inermis*, *Lycopersicon esculentum*, *Madhuca longifolia*, *Mangifera indica*, *Matricaria chamomilla*, *Mirabilis jalapa*, *Morinda pubescens*, *Morus alba*, *Opuntia dillenii*, *Plumbago zeylanica*, *Portulaca oleracea*, *Pongamia pinnata*, *Prunus persica*, *Rosmarinus officinalis*, *Saraca asoca*, *Scoparia dulcis*, *Terminalia bellirica*, *Thymus vulgaris* and *Vitex altissima*, to treat skin diseases, wounds, and skin cancers. Moreover, Turkish communities use *Rhus coriaria*, *Echium italicum*, *Juniperus oxycedrus*, and *Hypericum perforatum* for their anti-infective properties (Yeşilada *et al.*, 1993).

70% of the population of the second most-populous continent Africa still relies on their traditional medication system. They have a widespread use of *Prunus africana* and *Agathosma betulina* to treat urinary tract infections (Hostettmann *et al.*, 2000). Moreover, they use *Pelargonium sidoides* and *Glycyrrhiza glabra* as expectorants, and to treat upper respiratory tract infections. *Securidaca longepedunculata* is one of the common plants used to treat different infections, such as, skin infections, malaria, bilharzias and fever (Hostettmann *et al.*, 2000; Kamsu-Foguem and Foguem, 2014). In addition, *Catharanthus roseus* is utilised as a cancer chemotherapy (Kamsu-Foguem and Foguem, 2014). As a traditional herbal remedy, African people use *Hypoxis hemerocallidea* and *Sutherlandia spp.* as a primary treatment for HIV/AIDS (Mills *et al.*, 2005).

Egyptians use *Centaurium umbellatum Gilib*, *Azadirachta indica*, *Alternanthera philoxeroides*, *Castanospermum australe*, *Carica papaya*, *Euphorbia hirta*, *Kaempferia parviflora* and *Boesenbergia rotunda*, as antipyretics and garlic to treat intestinal worms (Petrovska, 2012; Pigili and Runja, 2014). Chinese plants such as *Artemisia annua*, *Lycoris radiata*, *Pyrrosia lingua*, *Isatis indigotica*, *Torreya nucifera*, and *Lindera aggregata* can inhibit the SARS-COV enzymes in coronavirus (Lin *et al.*, 2014). In addition, these plants

have been found to be successful in treating atopic eczema in children (Sheehan and Atherton, 1994).

1.3 Iraqi plants to treat infections

Iraq, situated in Western Asia, has a unique culture and diverse ecosystem. The country is bounded by Turkey to the north, Iran to the east, Kuwait to the southeast, Saudi Arabia to the south, Jordan to the southwest and Syria to the west. Iraq has distinctive geographical conditions due to the mountains in the north and northeast, and deserts in the southeast with low rainfall. The arid climate of Iraq with subtropical influence is favourable for the growth of particular types of flora, which has prevalent medicinal values. The medicinal plants mentioned in traditional Iraqi medicine is derived from knowledge of their historical usage. There is evidence that, 60,000 years ago, the Neanderthals lived in the equivalent of present-day Iraq. With the considerable advancement in civilisation starting from Sumerian, Babylonian, and Assyrian, the Iraqi people have transferred their folklore of herbal therapeutic information to successive generations (Chakravarty, 1976; Ghazanfar, 1994; Cowan, 1999; Al-Douri, 2014).

In spite of the continuous development of Iraq as a country, people are still dependent on the power of herbs as drinks or poultices for curing infections for their influential therapeutic benefits. Plants such as *Achillea vermicularis*, *Colutea cilicica*, *Gentiana olivieri*, *Potentilla supina* and *Polygonum aviculare* are used to treat helminth infections, dysentery and diarrhoea. *Aethionema grandiflorum*, *Agrimonia eupatoria*, *Allium sativum*, *Artemisia herba-alba*, *Cinnamomum zeylanicum*, *Citrullus colocynthis*, *Nigella sativa* and *Salvia officinalis* are commonly used to treat bacterial infections. For respiratory infections, *Anthemis nobilis*, *Allium sativum*, *Cassia occidentalis*, *Casuarina equisetifolia*, *Cordia myxa*, *Echium italicum*, *Glycyrrhiza glabra*, *Lavender angustifolia*, *Pimpinella anisum* and *Zingiber officinale* are used. In addition, for any gastrointestinal infections, *Ocimum basilicum* is a well-known remedy. Plants like, *Agropyron repens*, *Ammi majus*, *Herniaria*

glabra and *Jasminum officinale* are used for urinary tract infections. *Ammannia baccifera*, *Caccinia crassifolia*, *Citrus sinensis*, *Erodium cicutarium*, *Ficus carica* and *Fumaria parviflora* are used to treat skin infections. *Clerodendrum inerme* is claimed to be a good remedy for venereal infections, *Euphorbia hirta* for parasitic infections, and *Euphorbia tinctoria* for wounds (Molan *et al.*, 2012; Naqishbandi, 2014; Al-Snafi, 2018).

1.4 Natural products as antimicrobial agents

Natural products are chemical compounds, which have biological effects and are produced by living organisms in their natural environments (Sarker and Nahar, 2012; Obanla *et al.*, 2016). The potential use of highly purified natural products as antimicrobial agents dates back to 1929, when Alexander Fleming first reported the production of a bacteria-inhibiting substance in mould, *Penicillium*, later he termed this substance as “penicillin” (1). Unfortunately, he could not isolate the substance. Later in 1940, Florey and Chain were able to isolate penicillin and demonstrated its use as a therapeutic agent to treat a large number of bacterial diseases. The period between the 1950s and 1970s was the golden era of discovery of novel antibiotics that led to many new classes of antibiotics being discovered. Between 1981 and 1985, the *Journal of Antibiotics* reported approximately 50 effective compounds per year against pathogenic microorganisms were derived from natural products (Gootz, 1990). After that, 70 out of the 90 currently available antibiotics originated from natural products (Pelaez, 2006). During 2000-2008 more than 300 natural products compounds along with their antimicrobial properties were reported (Saleem *et al.*, 2010). The year 2015 perhaps started a new golden age for natural products research. In this year, Youyou Tu was awarded with the Nobel Prize in Physiology and Medicine for the discovery of artemisinin (2) and dihydroartemisinin (3), used to treat malaria (Shen, 2015). Antimicrobial agents can be obtained from different sources of natural products like, plants, animals, bacteria, algae and fungi (Gyawali and Ibrahim, 2014; Park, 2015). The number of

approved drugs since 1981 to 2014, derived from natural products in the period 1981 to 2014 is shown in Figure 1.1.

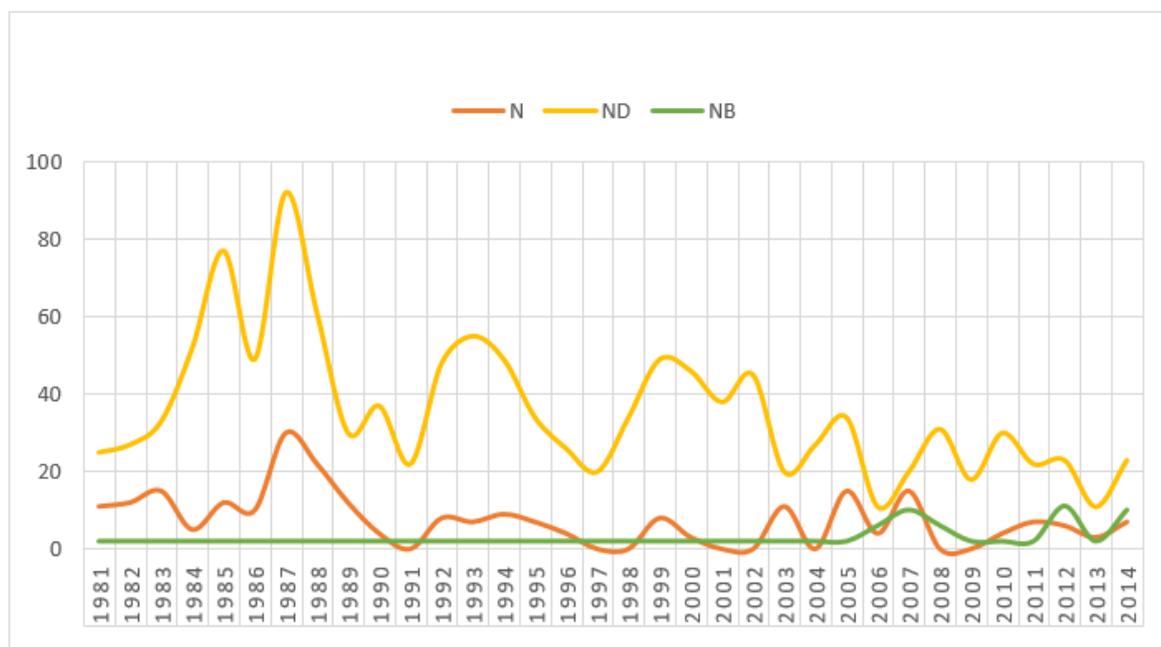


Figure 1.1: New approved drugs (1981-2014) derived from natural products

N: unmodified, semi or totally synthetic compounds, ND: semisynthetic modified natural compounds, NB: botanical drug

In 2015, 70% of novel drugs were from organic resources and out of them 64% were from natural products. The main source of 85% of new approved drugs in 2016 were organic compounds, with a half of them being natural products. In 2017, 87% of the novel drugs approved by the Food and Drug Administration (FDA) were obtained from organic sources, 50% of the active ingredients were natural products such as, ertugliflozin (4), deflazacort (5) and naldemedine (6) (Newman and Cragg, 2016; de la Torre and Albericio, 2018). Until September 2018, 75% of the new authorised drugs are from organic compounds and the active constituent for 50% of them derived from natural products such as cannabidiol (7) and sisomicin (8) (Figure 1.2).

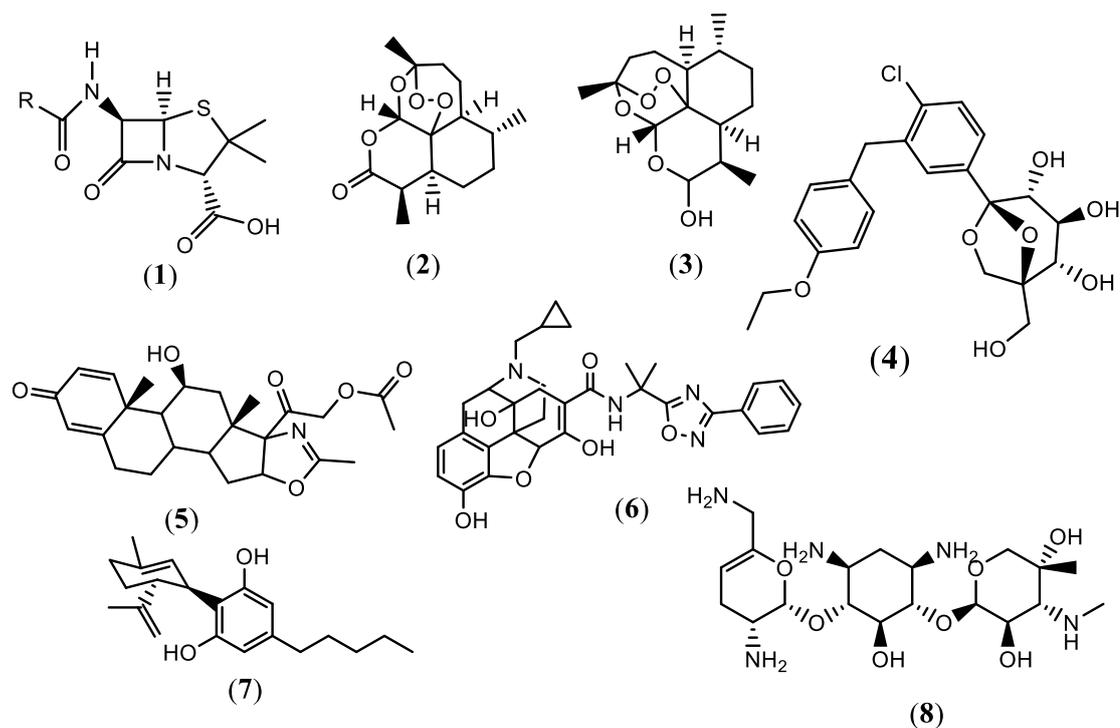


Figure 1.2: Structures of compounds 1-8

Penicillin (1), artemisinin (2), dihydroartemisinin (3), ertugliflozin (4), deflazacort (5), naldemedine (6), cannabidiol (7) and sisomicin (8)

1.4.1 Plants

Higher Plants produce a wide range of secondary metabolites such as coumarins, alkaloids, glycosides, flavonoids, steroids, terpenoids, sugars, saponins, organic acids, phenolics, aliphatic alcohols, aldehydes, ketones, and isoflavonoids etc., many of which can inhibit the growth of microorganisms, heal diseases or promote human health (Sarker and Nahar, 2012; Gyawali and Ibrahim, 2014; Obanla *et al.*, 2016).

The majority of researchers are working with the crude extract as phytopharmaceuticals (Iwu *et al.*, 1999). For example, the mustard crude extract showed antimicrobial activity against *E. coli* at 0.01% concentration (Chandual Ahire and khade, 2018) while, the methanolic extract of *Brassica oleracea* (red cabbage) inhibited bacteria *Staphylococcus aureus*, *Bacillus subtilis* and *Escherichia coli* at MIC 100 mg/mL (Hafidh *et al.*, 2018). According to Kumar *et al.* (2014) both of *Syzygium aromaticum* (Clove) and *Allium sativum* (Garlic) revealed antimicrobial effect against *Bacillus cereus*,

Staphylococcus aureus, *Salmonella typhi* and *Escherichia coli* at different concentrations (1000 ppm, 1500 ppm and 2000 ppm). Some studies focused on purification of the secondary metabolites and testing their antimicrobial properties (Gyawali and Ibrahim, 2014). According to Shan *et al.* (2007), 46 different plant extracts were found to contain phenolic constituents, which exhibited antibacterial activity against *Bacillus cereus*, *Listeria monocytogenes*, *Staphylococcus aureus*, *Escherichia coli*, and *Salmonella anatum* (Hafidh *et al.*, 2011).

The new trends of research are focussing on active compounds (Iwu *et al.*, 1999). Important pharmaceutically were isolated from plants include atropine (**9**) from *Atropa belladonna*, salicin (**10**) from *Salix spp.*, caffeine (**11**) from *Coffea arabica*, ephedrine (**12**) from *Ephedra spp.*, quinine (**13**) and quinidine (**14**) from *Cinchona spp.*, theobromine (**15**) from *Theobroma cacao*, theophylline (**16**) from *Camellia sinensis*, morphine (**17**) and codeine (**18**) from *Papaver somniferum*, digoxin (**19**) from *Digitalis spp.* and vinblastine (**20**) from *Catharanthus roseus* (Chinou, 2008) (Figure 1.3). The antimicrobial activity of pure compounds is dependent on chemical structure and concentration. Until now, over 30,000 antimicrobial compounds have been isolated from different plants (Gyawali and Ibrahim, 2014). Most of the expectations pointed to an increase in the use of pharmaceuticals derived from plants, even the WHO report has shown international interested in using these as drugs (Verma and Singh, 2008).

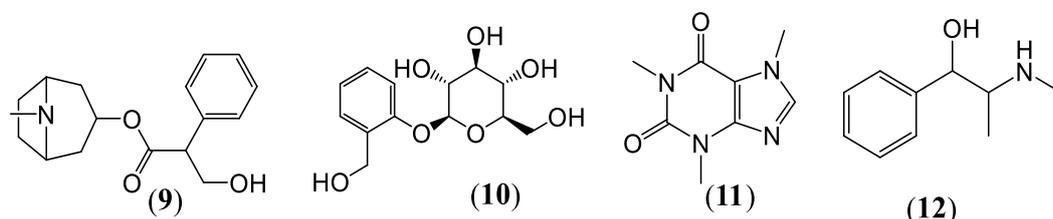


Figure 1.3: Structures of compounds from 9-20

Atropine (**9**), salicin (**10**), caffeine (**11**), ephedrine (**12**), quinine (**13**), quinidine (**14**), theobromine (**15**), theophylline (**16**), morphine (**17**), codeine (**18**), digoxin (**19**) and vinblastine (**20**)

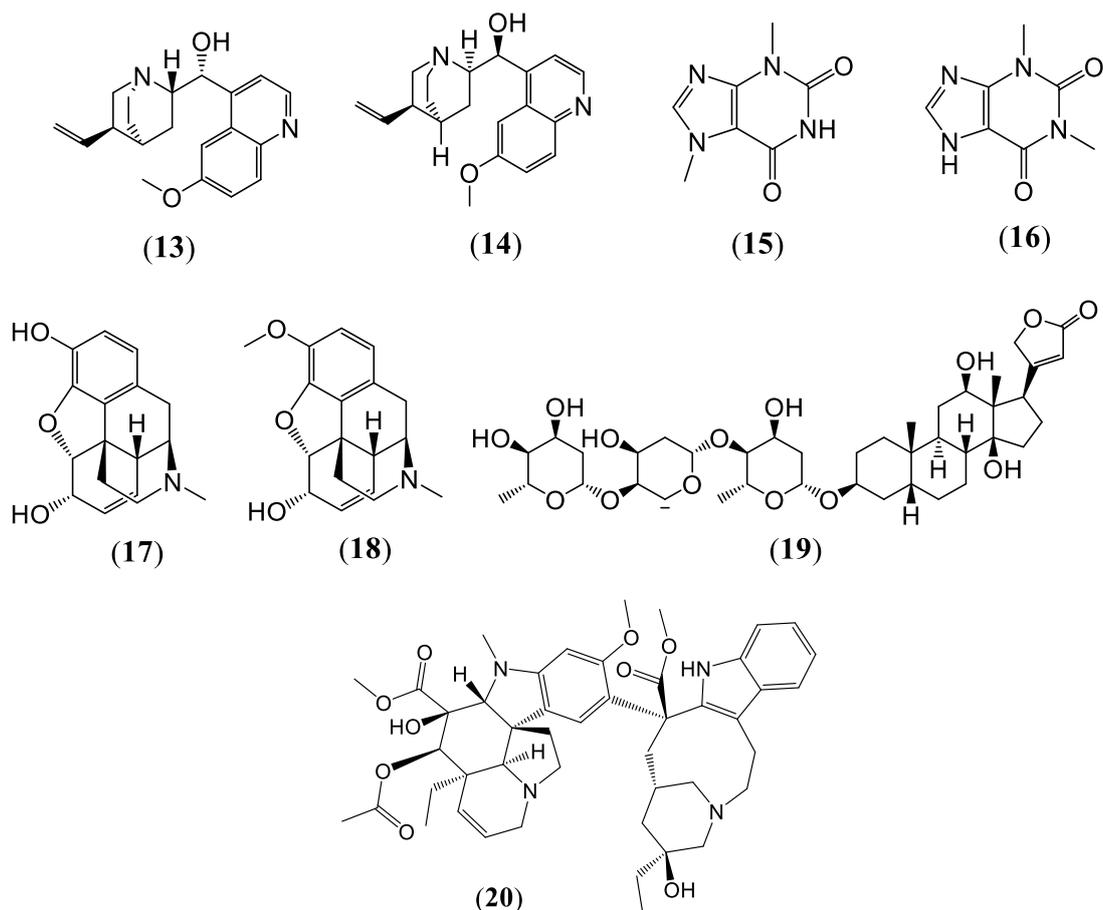


Figure 1.3(continued): Structures of compounds from 9-20

Atropine (9), salicin (10), caffeine (11), ephedrine (12), quinine (13), quinidine (14), theobromine (15), theophylline (16), morphine (17), codeine (18), digoxin (19) and vinblastine (20)

1.4.2 Microorganisms

Microorganisms, particularly bacteria and fungi, produce secondary metabolites during the stationary phase of growth. These metabolites are a rich source of valuable compounds for use in medicine industry and agriculture (Sultanbawa, 2011; Niu and Tan, 2013). Compounds such as penicillin (1), cephalosporin (21), tetracycline (22), chloramphenicol (23), aminoglycosides (24) and macrolides are well known antibiotics (Figure 1.4). The discovery of penicillin followed by innovation of numerable new antibiotics derived from microorganisms include rifamycin (25), erythromycin (26), chlortetracycline (27), streptomycin (28), cephalosporin C (29), lincomycin (30), vancomycin (31), nystatin (32), nalidixic acid (33), and daunorubicin (34) (Figure 1.4). All

of these compounds have a wide range of activities against Gram-positive and Gram-negative bacteria and fungi (Pelaez, 2006; Newman and Cragg, 2007; Dias *et al.*, 2012).

Fungi are a large source of secondary metabolites and novel compounds (Gunatilaka, 2006). These metabolites can have a broad-spectrum of activities against pathogens (Zhang *et al.*, 2006). Taxol (**35**), first discovered from *Taxus brevifolia*, has been isolated from *Pestalotiopsis microspore*, *Pestalotiopsis guepini*, *Seimatoantlerium tepuiense*, *Periconia* sp. and *Tubercularia* sp. (Strobel, 2003) (Figure 1.4).

Endophytic bacteria have a broad genetic diversity and are a rich source of natural products with a wide range of biological activities such as Streptomycetes, considered to be the source of up to 80% of the antibiotics (Strobel, 2003; Zhang *et al.*, 2006; De Lima Procópio, 2012). Many researchers have reported antibiotics, antioxidant agents, anticancer compounds, and other biologically active compounds from bacteria. For example celastramycin A (**36**) was purified from *Streptomyces* MaB-QuH-8 (Pullen *et al.*, 2002; Kikuchi *et al.*, 2009; De Lima Procópio, 2012) (Figure 1.4).

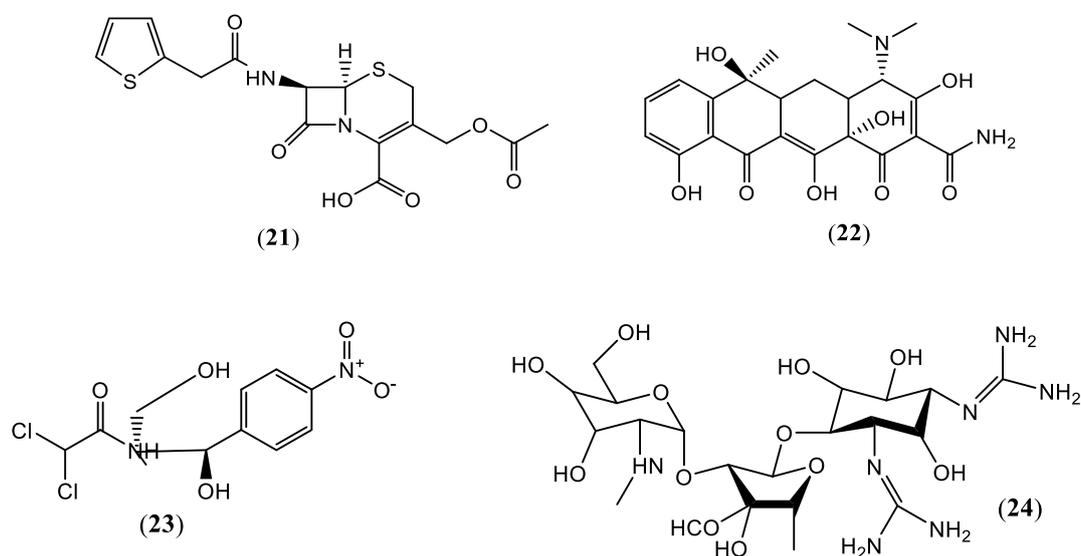


Figure 1.4: Structures of compounds from 21-36

Cephalosporin (**21**), tetracycline (**22**), chloramphenicol (**23**), aminoglycosides (**24**), rifamycin (**25**), erythromycin (**26**), chlortetracycline (**27**), streptomycin (**28**), cephalosporin C (**29**), lincomycin (**30**), vancomycin (**31**), nystatin (**32**), nalidixic acid (**33**), daunorubicin (**34**), taxol (**35**) and adenine (**36**)

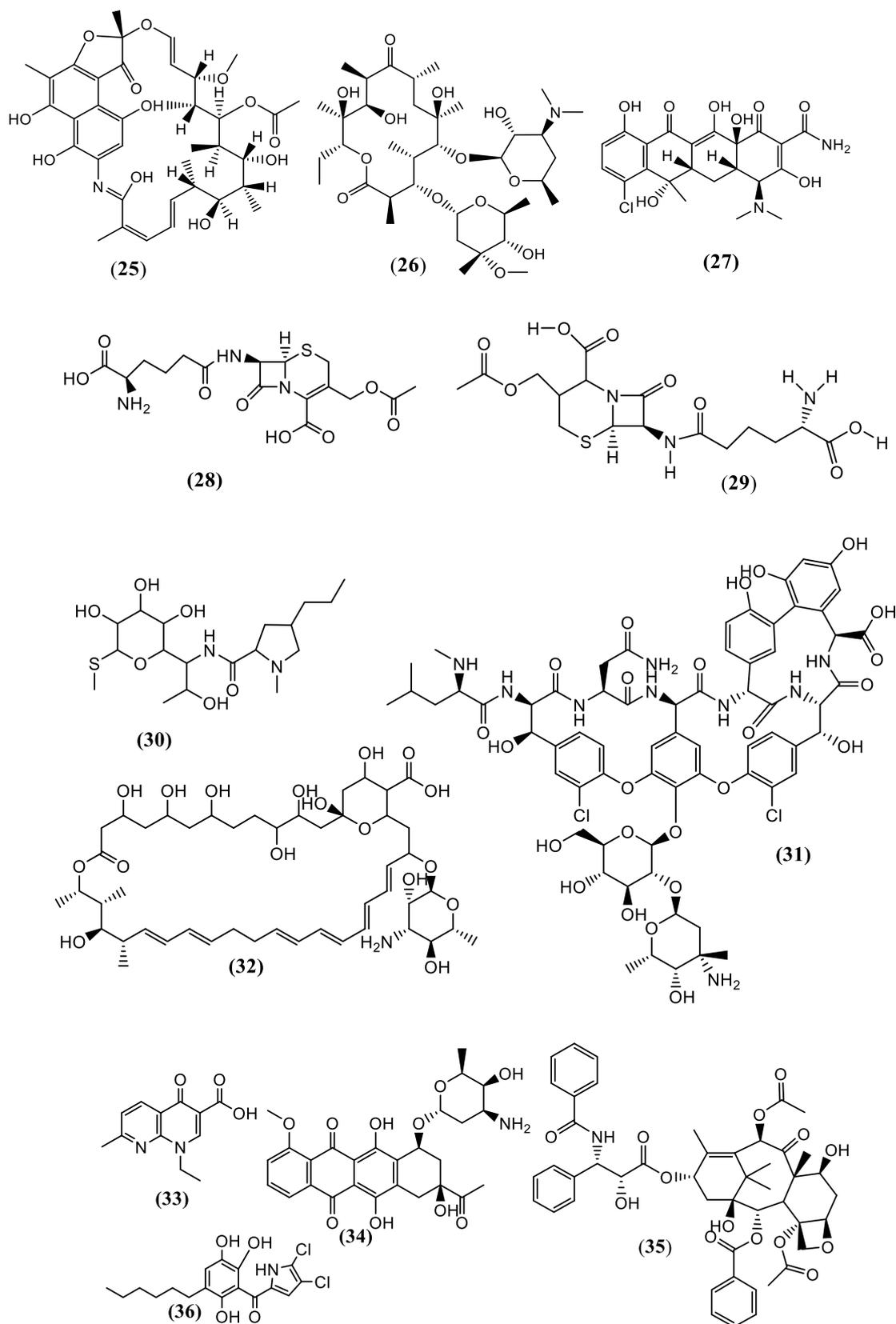


Figure 1.4 (Continued): Structures of compounds from 21-36

Cephalosporin (21), tetracycline (22), chloramphenicol (23), aminoglycosides (24), rifamycin (25), erythromycin (26), chlortetracycline (27), streptomycin (28), cephalosporin C (29), lincomycin (30), vancomycin (31), nystatin (32), nalidixic acid (33), daunorubicin (34), taxol (35) and adenine (36)

In addition, bacteria also produce enzymes and other proteins and enzymes, which have biological activities (Sahai and Manocha, 1993). *Bacillus subtilis* BS-2, for example, produces a thermostable and UV-tolerant antifungal protein (He *et al.*, 2003). It is well established that only a small fraction of microorganisms can be isolated from natural environments using current isolation techniques. Less than 1% of isolated bacteria and 5% of fungal species have ever been described (Patil *et al.*, 2016; Harvey, 2008). There is a need to explore alternative microbial habitats to isolate new microbes with chemical and functional diversity to obtain novel antibiotics (Patil *et al.*, 2016).

1.4.3 Marine sources

Discovery of bioactive compounds from marine organisms gained momentum in the middle of the 20th century and every year hundreds of novel compounds are now reported (Sawadogo *et al.*, 2013). Up to 2012, around 8,000 new compounds were reported, and most of them have claimed therapeutic properties for different diseases (Sarker and Nahar, 2012). Comparisons between the activity of terrestrial natural products and marine natural products show the great superiority of marine products which may be because marine organisms live in a strongly competitive environment for space and nutrition, and are therefore potential sources for detecting new biologically active compounds (Lu *et al.*, 2010; Rahman *et al.*, 2010; Mondol and Shin, 2014). The marine environment is considered particularly a valuable place for the variety of organisms (Sawadogo *et al.*, 2013).

Sponges, one of the oldest marine organisms belongs to metazoan phylum, considered to be a key source for pharmacologically active compounds. More than 5000 different compounds have already been discovered from sponges and 800 are documented as antimicrobial agents. Marine sponges are considered to be a major source of new compounds (Laport *et al.*, 2009; Skariyachan *et al.*, 2014). Marine organisms living at a depth of 11 km or more survive stressful conditions in terms of temperature, pressure and light (Sarker and Nahar, 2012). These conditions have contributed to the development of unique metabolites, which are different from those of other organisms. Marine

microorganisms can be a good source of new compounds, which may be a variety of chemical classes including alkaloids, peptides, terpenes, shikimates, polyketides and many of these are uncharacterized (Sarker and Nahar, 2012). Until now, only a few marine microorganisms have been examined for bioactive metabolites.

1.5 Mechanism of action of antimicrobial natural products

It is difficult to determine the exact mechanism of actions of natural antimicrobial agents, due to the concurrent occurrence of multiple interactions. Even though, different methods have been adapted to evaluate the mechanism of actions (Gyawali *et al.*, 2014). Studies on the mechanism of actions reveal the basis for the selective toxicity.

- Cell membrane: Some compounds have the ability to distribute membrane function and interact with membrane proteins, causing malformation in construction and functionality (Rimbara, 2012). For example, the Cranberry extract inhibits *S. aureus* by damaging the cell wall (Wu *et al.*, 2008). The Grape seeds and pomegranate polyphenols make a cell surface roughening and formation of blobs on the cell surface (Gyawali *et al.*, 2014)
- pH effects: an increase in proton concentration lowers pH of the substrate. The lower pH prevents intercellular acid molecules dissolving or creates a substrate transfer disorder by altering cell membrane permeability (Davidson *et al.*, 2013).
- Protein synthesis: The organic acids in plant extracts can have effects on the cytoplasmic membrane by interacting with membrane proteins (Gyawali *et al.*, 2014). In addition, the organic acids can be responsible for inhibiting NADH oxidation (Davidson *et al.*, 2013). Inhibition of plasmid conjugation and nucleic material can be caused by organic acids. This effect may not necessarily kill bacteria but could at least cause a decline in a bacterial population. Linoleic acid (**37**) from vegetables inhibits conjugal plasmid F and R388 (Fernandez-Lopez *et al.*, 2005; Jandacek, 2017) (Figure 1.5). Most leaves containing fatty acids and polyketides compounds were identified as specific conjugal inhibitors (Smith and Romesberg, 2007).

- Efflux pumps inhibitor: Proteinaceous transporters found in an organism's outer membrane, function to get rid of accumulated antibiotic molecules (Bay and Turner, 2016).
- The cytoplasmic membrane: Short chain organic acids in plant extracts can interfere with energy metabolism and affect membrane protein (Ricke, 2003). For example, Tea tree oil, which contain 100 terpenes and their related alcohols (Carson, 2002) can destroy the structure of the cytoplasmic membrane for *Escherichia coli* and *Staphylococcus aureus* (Cox *et al.*, 1998).

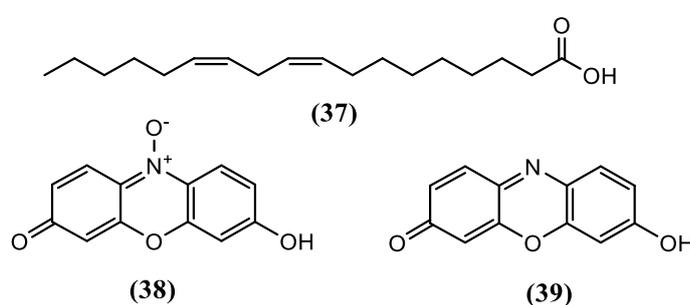


Figure 1.5: Structures of compounds 37-39

Linoleic acid (37), resazurin (38) and resorufin (39)

1.6 Methods for *in vitro* evaluation of antimicrobial activity

Different methods can be used to assess the *in vitro* antimicrobial activity for various materials including purified phytochemicals or crude natural products extracts and fraction. Often, one method may not be suitable for assessing the antimicrobial activity of different test samples. For example, for non-polar extracts, the diffusion method is not enough to confirm the antimicrobial activity. Many researchers have reported the solid dilution method as more accurate with non-polar extract. In addition, the amount of compound also dictates the researcher may choose a particular method (Houghton and Mukherjee, 2009). Among these methods, the most widely recognised are recorded below.

1.6.1 Diffusion method

Diffusion methods are widely used to evaluate the antimicrobial activity of plant extracts. In these methods, microorganisms are inoculated in solid media containing a

reservoir (hole, cylinder or disc) for material testing that means the plant extract will be in direct contact with the microbes through media. After incubation, the diameter of the inhibition zone around the reservoirs (Figure 1.6) is measured. In general, this method is good for primary antimicrobial screening of plant extracts by measuring the minimum inhibition concentration (MIC) (Ayukekbong *et al.*, 2017). When using this method, precautions should be taken not to compare the results between extracts obtained from different studies and to use the same solvent to dissolve the extracts. When using any of the three protocols, it is important to sterilise the sample and choose the suitable methods for extracts depending on solubility. If the extract remains undissolved in water, then the mixing in the whole plate will be the best choice for testing aqueous suspension (Hegggers *et al.*, 1987; Thomson *et al.*, 1989; Houghton and Mukherjee, 2009; Balouiri *et al.*, 2016).

1.6.2 Dilution method

In this method, a liquid culture medium is inoculated with the target organism and mixed with the plant extract. The suspension is incubated to measure the growth by turbidimetry in comparison with the control without plant extract. This method is most suitable for water-soluble extracts. The “microdilution method” can be used (Houghton and Mukherjee, 2009). This method uses 96-well microtitre plates (Figure 1.7) and the growth of bacteria is determined by turbidity (Kalemba and Kunicka, 2003). Only a small amount of plant extracts is needed for this method and one plate can be used to test many different concentrations. In addition, it is an easy method to determine the MIC (Houghton and Mukherjee, 2009).

Different methods of screening compounds obtained from plant extracts are discussed by Drummond and Waigh (2000), one of these methods is the resazurin assay (milk test), in which a blue compound is used as an indicator for oxidation in the reductase test of milk and as a pH indicator (pink at 3.8, violet at 6.5). Resazurin (**38**) is a metabolic indicator for living cells (O'brien *et al.*, 2000). So far, it has not been clear, where this

reaction occurs within the cell (Sánchez *et al.*, 2002). Researchers have suggested many possibilities on how Resazurin (38) is working inside living cells: such as , it can spread to the cell and is reduced to resorufin (39), possibly by the influence of several different reductase enzymes in mitochondria, cytosol and microsomes.

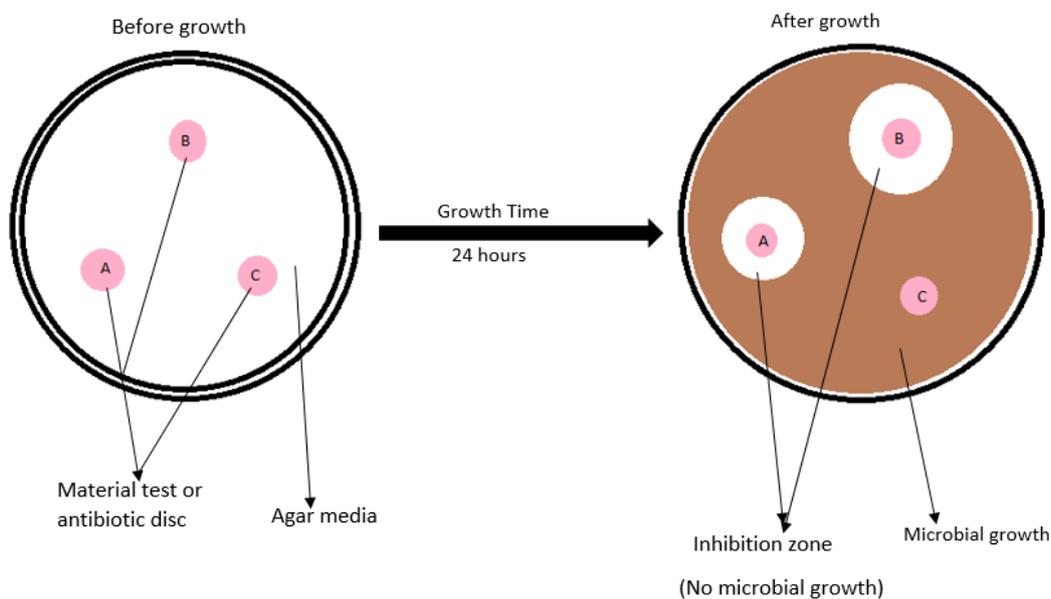


Figure 1.6: Disk diffusion method

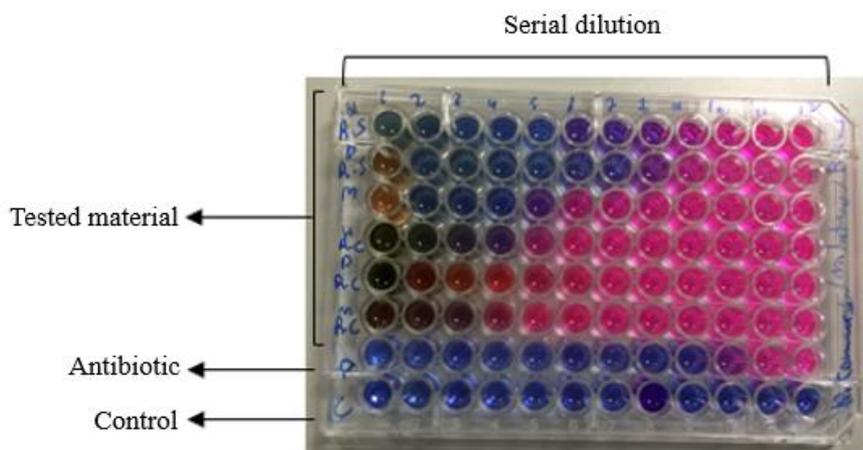


Figure 1.7: Dilution method

The fluorescent resorufin is then distributed from living cells to the surrounding medium (Wu, 2010). A second suggestion is that internal enzymes within cells such as NADH are involved in the interaction, which generates the co-reduction equivalents in this interaction (Riss and Moravec, 2006). Other studies have suggested that the reduction of

resazurin to resorufin may occur on the surface of the plasma membrane, or perhaps in the culture media (Quent *et al.*, 2010). Sarker *et al.* (2007) put forward a resazurin method with new modifications for better accuracy. The first method described by Drummond and Waigh (2000) used incubated microorganisms only, while Sarker *et al.* (2007) measured the incubated microorganisms by spectrophotometer and compared with the Macfarlane standard. This modified resazurin method is simple, sensitive, fast, strong, reliable, and can be successfully used to evaluate the antibacterial properties of natural products.

1.7 Plants used in this study

This study included three plants from the Rutaceae family from the Iraqi flora: *Citrus grandis* (leaves), *Citrus sinensis* (leaves and peel) and *Ruta chalepensis* (leaves, stems, flowers and roots). The Rutaceae family is commonly known as “the *Citrus* family”. The *Citrus* species are commercially important, and the Rutaceae is the biggest family in the Iraqi flora, comprising around 160 genera and 1700 species (Heywood *et al.*, 2007; Martínez-Pérez *et al.*, 2017). Different types of plants like shrubs, trees, herbs and sometimes climbing or crawling plants come under this family. Most of the Rutaceae members have essential oil in the visible glands of their leaves, flowers, fruits and seeds. Leaves are simple or compound and alternate or opposite; most of the plants in this family have beautiful flowers with bright colours and strong fragrance (Swingle, 1943; Groppo *et al.*, 2008; Tamokou *et al.*, 2017; Saduf *et al.*, 2018). The flowers have long corollas ranging from a few millimetres to several centimetres, and the symmetry pattern of a flower is actinomorphic or zygomorphic. Carpels and stamens are two to many; carpels are free or united and can contain two to many ovaries (Groppo *et al.*, 2008). Fruits can be dry like *Ruta chalepensis* or fleshy like *Citrus sinensis* (Groppo *et al.*, 2008). The species grow in all parts of the world especially in Asia, Australia, South Africa and Africa (Swingle, 1943). This plant family has been used in traditional medicine for treating snakebites, stomatitis, rheumatism, bronchitis and other diseases (Sandjo *et al.*, 2014).

The Rutaceae has been reported to contain 200 different coumarins (Gray and Waterman, 1978). The most economically important genus in the family is *Citrus* (Chase *et al.*, 1999). This family is a good source of alkaloids such as benzyloquinoline, furquinolines, quinolines, acridones, pyanoquinolines, indole derivatives, imidazoles and oxazoles (Waterman, 1975; Shobana *et al.*, 1989; Groppo *et al.*, 2008), furanocoumarins, furoquinoline alkaloids, phenolics and terpenes (Adamska-Szewczyk *et al.*, 2016), a big range of limonoids (Wang *et al.*, 2009) flavonoids (Koblovská *et al.*, 2008) coumarins and volatile oils. Lots of plants from this family have been used traditionally, for example, the genus *Dictamnus* has been used in traditional Chinese medicine to heal bleeding, rheumatism, itching, hepatitis disease and skin problems (Tang *et al.*, 1999; Wu *et al.*, 1999). Another member of the Rutaceae is *Toddalia asiatica*, in South Africa, it is used to treat malaria, indigestion problems, stomachache, snakebites, and is used in rituals, coughs, influenza, lung diseases, nasal problems, bronchial pains and rheumatism (Adjanohoun *et al.*, 1993; Maundu, 2001; Meyer, 2005; Duraipandiyar *et al.*, 2006; Orwa *et al.*, 2008; Kokwaro, 2009).

A number of scientific reports have been published on the potential medicinal uses of plants belonging to the Rutaceae family; such as the extract of the leaves of *Chloroxylon swietenia*, which has significant anti-inflammatory activity in a rat paw oedema test (Kumar *et al.*, 2006). Another example is *Zanthoxylum chalybeum* extract, which has also shown anti-inflammatory activity (Agyare *et al.*, 2013). Both *Zanthoxylum usambarense* and *Ruta spp.* have antimalarial activity (Agyare *et al.*, 2013). In addition, many members of the Rutaceae family have reports of antimicrobial activity.

Based on reports by Vats *et al.* (2011), the extract of *Murraya koenigii* can inhibit *Staphylococcus aureus*, *Micrococcus luteus*, *Bacillus subtilis*, *Escherichia coli*, *Pseudomonas aeruginosa*, *Candida albicans* and *Aspergillus niger*. During the screening of 172 plant extracts for antimicrobial activity in Puerto Rico, it was found that two species, *Citrus aurantifolia* and *C. aurantodium* were effective (Meléndez and Capriles, 2006). In

another study in Texas, the USA, fruit extract of *C. aurantifolia* acted as an anticancer agent, the result showed that 100 µg/mL of *C. aurantifolia* after 48 h exposure can inhibit growth of colon cancer cells SW-480 up to 78%, by damaging the DNA and expanding the level of caspase-3 (Narang and Jiraungkoorskul, 2016). The Rutaceae members are also known for their antioxidant properties. In a study carried out in India on *Clausena excavate* plant extract the result showed that the plant has high antioxidant activity (Elumalai and Id, 2016).

1.7.1 The genus *Ruta*

The genus *Ruta* L. grows in rocky and arid grasslands (Bennaoum and Benhassaini, 2017), contains 14 species of herbs and shrubs that emit green leaves with strong odour and has a bitter taste (Pollio *et al.*, 2008). *Ruta angustifolia*, *Ruta chalepensis*, *Ruta graveolens* and *Ruta montana* are some of the better known species of this genus.

1.7.1.1 *Ruta chalepensis* L.

Ruta chalepensis L., commonly known as “rue or ruda” belongs to the Rutaceae family, and is a perennial herb and shrub having wild as well as cultivated forms. This species is characterised by glabrous, alternating bi-pennatisect leaves with narrow oblong-lanceolate or obovate segments and a cymose inflorescence (Figure 1.8). Rue has a strong odour due to the oil glands located in the leaves. The individual yellow flower consists of 4-5 sepals, 4-5 petals, 8-10 stamens and a superior ovary.



Figure 1.8: *Ruta chalepensis* (A: fruits, B: flowers and leaves)

Origin and distribution:

Ruta chalepensis is indigenous to the Mediterranean region and Canary Islands. In the tropics, it is cultivated as a potherb or medicinal plant. It has been introduced in several countries, which are listed in the Table 1.1.

Traditional uses:

Documented reports suggest various traditional uses of Rue as an anti-inflammatory, analgesic, antipyretic and for uses in rheumatism, nerves disease, neuralgia, dropsy, convulsions and mental disorder. Moreover, it is also used to treat hysteria, colic, headache, eye problems, ear infections, intestinal worms, stomachache, poleaxe, apprehension and intestinal worms (Al-Said *et al.*, 1990; Béjar *et al.*, 2001; Iauk *et al.*, 2004; Alanis *et al.*, 2005; Günaydin and Savci, 2005; Gonzalez-Trujano *et al.*, 2006; Mejri *et al.*, 2010; Ali *et al.*, 2013; Khoury *et al.*, 2014). According to the study of Ali *et al.* (2004), *Ruta* is one of nineteen Yemeni traditional herbal medicines with known traditional use for Malaria treatment. The results of Günaydin and Savci (2005) however emphasised that *Ruta* causes embryo toxicity in mice and many other harmful effects. In addition, published literature suggest that *Ruta* is an abortifacient (San Miguel; 2003; Mejri *et al.*, 2010; Martínez-Pérez *et al.*, 2017). The plant is also used as a laxative. In addition to all the above features, *R. chalepensis* is also famous as an insect and lice repellent (Ali *et al.*, 2013; Khoury *et al.*, 2014).

Veterinary uses:

Apart from the medicinal usage of *R. chalepensis*, it has an immense claimed effect in farm and poultry animal health. It helps in delivery of sheep, pigs, goat, ewe and cows. Also, it solves digestive problems, joints pain, wounds and acts against parasites. In addition, it is used to kill lice on chickens, and to kill parasites in sheep and pigs. Moreover, it is used to cure earache, eye-ache and to heal the vocal cords in birds (San Miguel, 2003). According to de Sa *et al.* (2000) certain dosages of *R. chalepensis* causes many changes in the placenta and causes disorders in the blood-brain barrier, resulting in embryonic toxicity.

Table 1.1: Worldwide distribution of *Ruta chalepensis*

Country	Reference
Algeria	Dod and Dahmane, 2008; Ferhat <i>et al.</i> , 2014; Khourya <i>et al.</i> , 2014
America	Günaydin and Savci, 2005; Khoury <i>et al.</i> , 2014
Argentina	Zeichen <i>et al.</i> , 2000; Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003
Central America	San Miguel, 2003
China	San Miguel, 2003; Ferhat <i>et al.</i> , 2014; Khoury <i>et al.</i> , 2014
Egypt	Ali <i>et al.</i> , 2013
Greece	Tzakou and Couladis, 2001; Dod and Dahmane, 2008; Ferhat <i>et al.</i> , 2014; Khourya <i>et al.</i> , 2014
India	Bagchi <i>et al.</i> , 2003; San Miguel, 2003; Dod and Dahmane, 2008; Khoury <i>et al.</i> , 2014; Ferhat <i>et al.</i> , 2014
Iran	Rustaiyan <i>et al.</i> , 2002; Dod and Dahmane, 2008; Rustaiyan <i>et al.</i> , 2011; Ferhat <i>et al.</i> , 2014; Khourya <i>et al.</i> , 2014; Tedone <i>et al.</i> , 2014
Iraq	Al-Majmaie <i>et al.</i> , 2005
Israel	Ali-Shtayeh and Abu Ghdeib, 1999; Landaua <i>et al.</i> , 2004
Italy	Ali <i>et al.</i> , 2013; Iauk <i>et al.</i> , 2014; Tedone <i>et al.</i> , 2014
Jordan	Alomary <i>et al.</i> , 2013
Latin America	San Miguel, 2003
Lebanon	Khourya <i>et al.</i> , 2014
Mediterranean area	Ulubelen <i>et al.</i> , 1986; Iauk <i>et al.</i> , 2014 ; Tedone <i>et al.</i> , 2014
Mexico	Ali <i>et al.</i> , 2013
Middle east	San Miguel, 2003; Khoury <i>et al.</i> , 2014; Bouabidi <i>et al.</i> , 2015
Morocco	Uphof, 1968; Günaydin and Savci, 2005
North of India	Ferhat <i>et al.</i> , 2014
New York	San Miguel, 2003
North California	San Miguel, 2003
North America	Rustaiyan <i>et al.</i> , 2002; San Miguel, 2003
North Africa	Khourya <i>et al.</i> , 2014
Palestine	Ali-Shtayeh and Abu Ghdeib, 1999; Landaua <i>et al.</i> , 2004
Portugal	Ferhat <i>et al.</i> , 2014
Saudi Arabia	Shah <i>et al.</i> , 1991; Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Günaydin and Savci, 2005
Spain	San Miguel, 2003; Khoury <i>et al.</i> , 2014; Ferhat <i>et al.</i> , 2014
Southern Europe	Rustaiyan <i>et al.</i> , 2002
Texas	San Miguel, 2003
Tunisia	Mejri <i>et al.</i> , 2010; Ferhat <i>et al.</i> , 2014; Khourya <i>et al.</i> , 2014; Tedone <i>et al.</i> , 2014
Turkey	Ulubelen <i>et al.</i> , 1986; Ulubelen and Terem, 1988; Baser <i>et al.</i> , 1996; Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Günaydin and Savci, 2005; Dod and Dahmane, 2008; Ali <i>et al.</i> , 2013
Yemen	Alzorekya and Nakahara, 2003; Awadh <i>et al.</i> , 2004
Western Atlantic	San Miguel, 2003

Phytochemicals of Ruta chalepensis:

Previous phytochemical studies on *R. chalepensis* have shown the presence of essential oils (Al-Said *et al.*, 1990; Gonzalez-Trujano *et al.*, 2006; Ali *et al.*, 2013), alkaloids (Al-Said *et al.*, 1990; San Miguel, 2003; Iauk *et al.*, 2004; Khoury *et al.*, 2014) including furoquinoline alkaloids, 5-methoxydictamine (**40**) (Günaydin and Savci, 2005), skimmianine (**41**), quinolone alkaloid, γ -fagarine (**42**), saponins (Al-Said *et al.*, 1990; Ali *et al.*, 2013), sterols, tannins, triterpenes (Gonzalez-Trujano *et al.*, 2006; Ali *et al.*, 2013), coumarins (Khoury *et al.*, 2014) including furanocoumarins, bergapten (**43**) isopimpinellin (**44**) (Ulubelen *et al.*, 1986; Günaydin and Savci, 2005) chalepin (**45**), chalepinsin (**46**) rutamarin (**47**), and the flavonoid rutin (**48**) (Figure 1.9). Additionally, amino acids (de Sa *et al.*, 2000; Iauk *et al.*, 2004; Ali *et al.*, 2013), anthraquinones and various simple phenols (Ali *et al.*, 2013; Khoury *et al.*, 2014) were purified from this species. The essential oils of *Ruta chalepensis* contain many compounds some of which are listed in Table 1.2.

Bioactivity:

Ruta chalepensis as a repository of numerous compounds motivated the researchers from time to time to investigate the bioactivity of these compounds. It has been reported that the plant possesses antifungal, antibacterial, antimalarial and antiparasitic activities (Mejri *et al.*, 2010; Khoury *et al.*, 2014; Bouabidi *et al.*, 2015). Antitumor (Krayni *et al.*, 2015) and anticancer activity, especially for colon cancer (Tedone *et al.*, 2014), have also been reported. In a study in Italy, fifty-six patients suffering from different stages of colon cancer were treated with an ethanolic extract of *R. chalepensis*. As a result, there was a decrease in disease progression (Acquaviva *et al.*, 2011; Tedone *et al.*, 2014). In Palestine, as a folk medicine, aqueous extracts of 20 plants were evaluated for their antifungal activity, and out of these *R. chalepensis* was one of the most potent plants to inhibit *Microsporum canis*, *Trichophyton mentagrophytes* and *Trichophyton violaceum* (Ali-Shtayeh and Abu Ghdeib, 1999). Furthermore, according to Alzoreky and Nakahara (2003), in an experiment with the

extracts of 26 plants collected from different countries, *R. chalepensis* was effective in eliminating *E. coli* and *Salmonella infantis*. *Ruta chalepensis* is also effective against *Bacillus subtilis*, but it is less active against *Staphylococcus aureus* and has no activity against *E. coli* (Al-Bakri and Afifi, 2007). During an antibacterial activity study using the disc diffusion method (Perez and Anesini, 1994), it was found that there was no activity against *Salmonella typhi*.

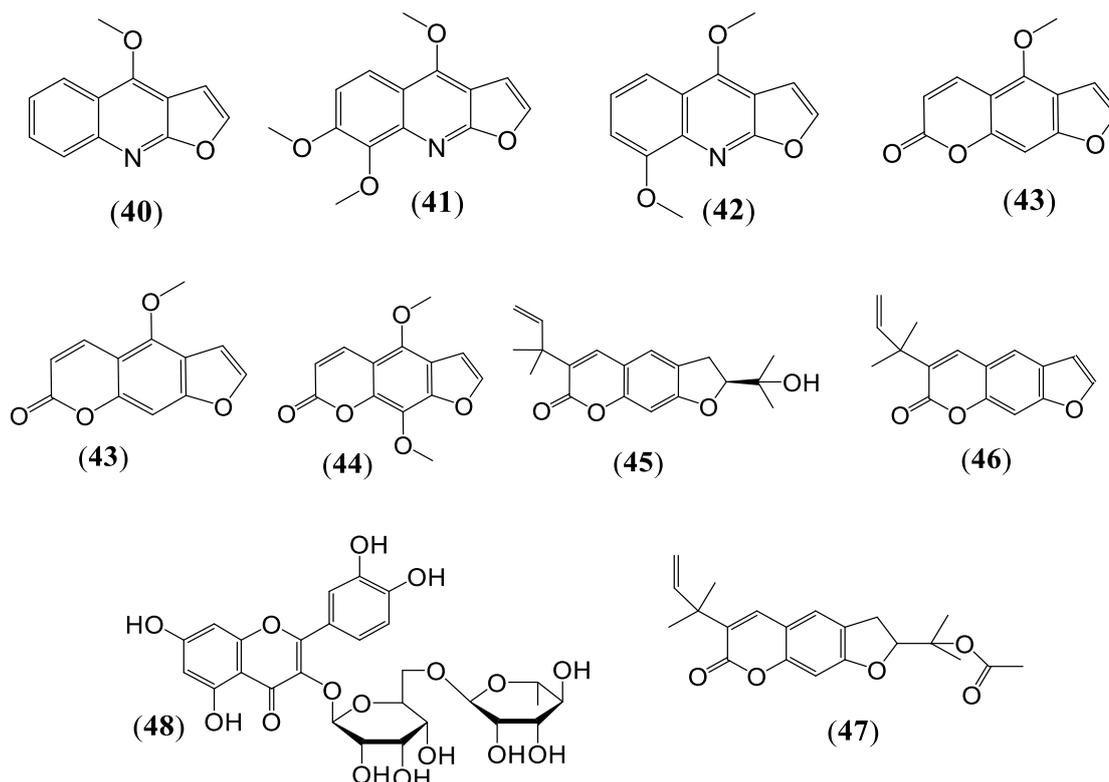


Figure 1.9: Structures of compounds 40-48

5-methoxydictamine (40), skimmianine (41), γ -fagarine (42), bergapten (43), isopimpinellin (44), chalepin (45), chalepinsin (46), rutamarin (47) and rutin (48)

Table 1.2: Compounds identified in essential oils of *R. chalepensis*

Chemical compound	Reference
2-Acetoxytridecane	Ferhat <i>et al.</i> , 2014
2-Acetoxytetradecane	Ferhat <i>et al.</i> , 2014
Benzaldehyde	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
(Z)-Bisabolene	Tzakou and Couladis, 2001
γ -Bisabolene	Khoury <i>et al.</i> , 2014
δ -Cadinene	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014

Camphor	Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Mejri <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Carnphene	Tzakou and Couladis, 2001
β -Caryophyllene	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Carvacol	Bagchi <i>et al.</i> , 2003
Chamazulene	Dob and Dahmane, 2008
1,8-Cineole	Bnina <i>et al.</i> , 2010
<i>p</i> -Cymene	Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008, Bnina <i>et al.</i> , 2010
Curcuphenol	Bnina <i>et al.</i> , 2010
γ -Decalactone	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
Decanal	Dob and Dahmane, 2008
Decanol	Bnina <i>et al.</i> , 2010
2-Decanol	Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008
2-Decanone	Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Mejri <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014; Ferhat <i>et al.</i> , 2014
2-Decyl acetate	Tzakou and Couladis, 2001; Rustaiyan <i>et al.</i> , 2002; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014
2-Decanyl acetate	Ferhat <i>et al.</i> , 2014
Dodecanol	Tzakou and Couladis, 2001; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
2-Dodecanone	Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Mejri <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Ferhat <i>et al.</i> , 2014; Khoury <i>et al.</i> , 2014
Dodecan-3-one	Bagchi <i>et al.</i> , 2003; Mejri <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Elemol	Tzakou and Couladis, 2001; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
2-Ethylheptyl acetate	Tzakou and Couladis, 2001
Ethyl linoleate	Dob and Dahmane, 2008
Ethyl octadecanoate	Bnina <i>et al.</i> , 2010
2-Ethylloctyl acetate	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
α -Eudesmol	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
β -Eudesmol	Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010
γ -Eudesmol	Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010
(<i>E,E</i>)- α -Farnesene	Tzakou and Couladis, 2001; Bnina <i>et al.</i> , 2010
(<i>Z,E</i>)- α -Farnesene	Tzakou and Couladis, 2001
Furfural	Dob and Dahmane, 2008
Geijerene	Tzakou and Couladis, 2001; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013
Geraniol	Tzakou and Couladis, 2001
Geranyl acetate	Tzakou and Couladis, 2001
Heptanal	Dob and Dahmane, 2008

Heptan-2-one	Bnina <i>et al.</i> , 2010
2-Heptanol	Dob and Dahmane, 2008
2-Heptyl acetate	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bouabidi <i>et al.</i> , 2015
Hexadecanoic acid	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
2-Hexanal	Dob and Dahmane, 2008
(Z)-3-Hexanal	Dob and Dahmane, 2008
Hex-3-en-1-ol	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Hex-2-enylacetate	Bnina <i>et al.</i> , 2010
Hex-3-enyl acetate	Bnina <i>et al.</i> , 2010
α -Humulene	Bnina <i>et al.</i> , 2010
β -Ionone	Dob and Dahmane, 2008
Isophytol	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Limonene	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014
Linalool	Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008
Mesitylene	Dob and Dahmane, 2008
<i>p</i> -Mentha-1,8-diene	Tzakou and Couladis, 2001
Methyl chavicol	Bnina <i>et al.</i> , 2010
2-Methyldecyl acetate	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Ferhat <i>et al.</i> , 2014
6-Methyl-5-hepten-2-one	Bagchi <i>et al.</i> , 2003
Methyl hexadecanolate	Dob and Dahmane, 2008
10-Methyl-2-undecanone	Ali <i>et al.</i> , 2013
2-Methylnonyl acetate	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
Methyl nonyl ketone	Ferhat <i>et al.</i> , 2014
Methyl octadecanoate	Bnina <i>et al.</i> , 2010
2-Methoxy-3-isopropyl-5-methylpyrazine	Dob and Dahmane, 2008
Myrcene	Tzakou and Couladis, 2001; Bnina <i>et al.</i> , 2010
Naphtalene	Bnina <i>et al.</i> , 2010
(<i>E</i>)-Nerolidol	Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010
Neryl acetate	Tzakou and Couladis, 2001
Nonan-2-yl	Khoury <i>et al.</i> , 2014
Nonyl acetate	Tzakou and Couladis, 2001; Rustaiyan <i>et al.</i> , 2002
Nonanal	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014
2-Nonanol	Tzakou and Couladis, 2001; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014
Nonan-3-ol	Bnina <i>et al.</i> , 2010
2-Nonanone	Tzakou and Couladis, 2001; Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Ferhat <i>et al.</i> , 2014; Khoury <i>et al.</i> , 2014
2-Nonyl acetate	Bagchi <i>et al.</i> , 2003; Ali <i>et al.</i> , 2013; Ferhat <i>et al.</i> , 2014

(<i>E</i>)- β -Ocimene	Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010
(<i>E</i>)- <i>p</i> -Ocimene	Tzakou and Couladis, 2001
β -Ocirnene	Tzakou and Couladis, 2001
Octanol	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
2-Octanol	Dob and Dahmane, 2008
2-Octanone	Rustaiyan <i>et al.</i> , 2002; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Octyl acetate	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
2-Octyl acetate	Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013
3-Octyl acetate	Rustaiyan <i>et al.</i> , 2002
3-Odecanone	Dob and Dahmane, 2008
β -Phellandrene	Tzakou and Couladis, 2001; Ferhat <i>et al.</i> , 2014
Phenylacetaldehyde	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
Phytol	Ali <i>et al.</i> , 2013; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Phytyl acetate	Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
α -Pinene	Tzakou and Couladis, 2001; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014; Bouabidi <i>et al.</i> , 2015
β -Pinene	Tzakou and Couladis, 2001
Pregeijerene	Tzakou and Couladis, 2001; Bnina <i>et al.</i> , 2010
Pregeijerene B	Ali <i>et al.</i> , 2013
Psoralen	Dob and Dahmane, 2008
Pulegone	Bnina <i>et al.</i> , 2010; Mejri <i>et al.</i> , 2010
<i>cis</i> -Sabinene hydrate	Tzakou and Couladis, 2001; Bnina <i>et al.</i> , 2010
α -Terpinene	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
β -Terpinene	Tzakou and Couladis, 2001; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008
Terpinolene	Tzakou and Couladis, 2001
α -Terpineol	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
Terpinen-4-ol	Tzakou and Couladis, 2001; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Khoury <i>et al.</i> , 2014
Tetradecanol	Bnina <i>et al.</i> , 2010
α -Thujene	Tzakou and Couladis, 2001; Dob and Dahmane, 2008
Thymol	Bagchi <i>et al.</i> , 2003
2-Tridecanone	Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Bnina <i>et al.</i> , 2010; Mejri <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Khoury <i>et al.</i> , 2014
1,2,4-Trimethylbenzene	Dob and Dahmane, 2008
2-Undecanol	Ali <i>et al.</i> , 2013; Dob and Dahmane, 2008
2-Undecanone	Tzakou and Couladis, 2001; Rustaiyan <i>et al.</i> , 2002; Bagchi <i>et al.</i> , 2003; Dob and Dahmane, 2008; Mejri <i>et al.</i> , 2010; Ali <i>et al.</i> , 2013; Bouabidi <i>et al.</i> , 2015
2-Undecanyl acetate	Rustaiyan <i>et al.</i> , 2002

Side effects of Ruta chalepensis:

Ruta chalepensis is considered to be one of the dangerous plants, especially in high dosage it may lead to death. Not recommended to use by pregnant or lactating women. It may cause abortion, hyperaemia in the uterus and high oxytocic action (San Miguel, 2003). According to El Sayed *et al.* (2000), *R. chalepensis* causes mutagenicity due to its quinolone alkaloids content.

1.7.2 The genus *Citrus* L.

In 1753, Carl Linneaus presented the genus “*Citrus*” to the world (Swingle, 1967). These plants grow in tropical and subtropical areas and in the borderline around the world (The *Citrus* and Date Crop Germplasm Committee, 2004; Sun *et al.*, 2015). The *Citrus* L. has 16 species (Sun *et al.*, 2015), and all of them are trees with a height around 12-30 feet and width ranging between 8-25 feet. The plants have rounded crowns of dark green leaves, which are simple, trifoliate, alternate, and source of essential oils. In addition, *Citrus* has fragrant blossoms (Watson, 1993). This genus is the source of *Citrus* fruits, which has high economical value (The *Citrus* and Date Crop Germplasm Committee, 2004; Sun *et al.*, 2015). Between 2009 and 2010, the world’s total production was up to 7.4 million metric tons (Sun *et al.*, 2015).

The plants from the *Citrus* genus have been used in traditional medicine for treating different diseases as an analgesic (Lima *et al.*, 2007; Correa *et al.*, 2016) for us in rheumatism and malaria (Martín *et al.*, 2011). *Citrus* species are reported to have different medicinal, physiological and pharmacological properties, such as antimicrobial (Espina *et al.*, 2011), antioxidant (Goulas and Manganaris, 2012), anticancer (Manthey and Guthrie, 2002; Cirmi *et al.*, 2017) and anti-inflammatory (Menichini *et al.*, 2011). The most important part of the *Citrus* plants is the edible fruit, used for foods, drinks, perfumes, cosmetic, soaps, and in many other aromatic products (Mohammed *et al.*, 2017). *Citrus* natural products are widely perceived as a critical part of the human eating regimen. It is cholesterol free and contains

sodium, vitamin C, folic acid, potassium, flavonoids, coumarins, pectins and dietary fibres (Dugo and Di Giacomo, 2002; Roy *et al.*, 2014).

Phytochemical studies on the genus *Citrus* have shown the presence of essential oils (Gancel *et al.*, 2005), and phenolic compounds (Fejzić and Čavar, 2014). This genus is considered as a main source of vitamin C (**49**), which provides various health benefits (Zhang *et al.*, 2011). It contains large amounts of polyphenolics, especially flavonoids, such as hesperidin (**50**), narirutin or naringin (**51**), neohesperidin (**52**), eriocitrin (**53**), neoeriocitrin (**54**), rutin (**48**), diosmin (**55**), neoponcirin (**56**), and nobiletin (**57**), which have significant biological activity (Albach *et al.*, 1969; Jourdan *et al.*, 1985; Wang *et al.*, 2007; Zhang *et al.*, 2011) (Figure 1.10). Many other compounds have also been documented from *Citrus* including phenolic acids, β -carotene (**58**), lutein (**59**), lycopene (**60**) cryptoxanthin (**61**) and citric acid (**62**) (Lee *et al.*, 2001; Xu *et al.*, 2006; Kurita *et al.*, 2008). Lycopene is of note for preventing vacuolization in human lens epithelial cells (Mohanty *et al.*, 2002).

In addition, this genus produces hydroxycinnamic acids (**63**), which can inhibit oxidation of low-density lipoprotein, such as caffeic acid (**64**), ferulic acid (**65**), chlorogenic acid (**66**), and *p*-coumaric acid (**67**). In general, *Citrus* fruits are rich in carotenoids, for example, pink fruit has a high concentration of β -carotene (**58**), while other fruits contain high level of lutein (**59**), cryptoxanthin (**68**), zeaxanthin (**69**), and β -cryptoxanthin (**70**) (Ma *et al.*, 2013) (Figure 1.10).

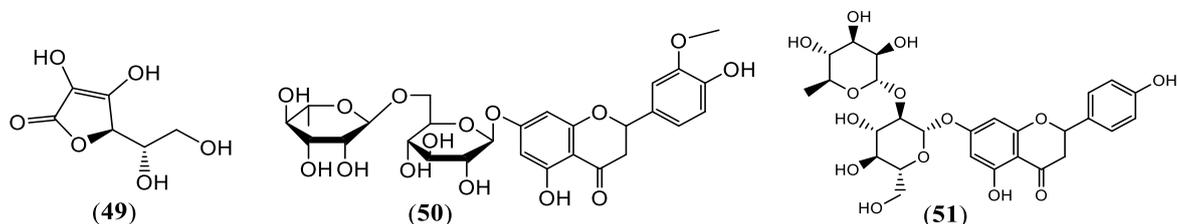


Figure 1.10: Structures of compounds 49-70

Vitamin C (**49**), hesperidin (**50**), narirutin or naringin (**51**), neohesperidin (**52**), eriocitrin (**53**), neoeriocitrin (**54**), diosmin (**55**), neoponcirin (**56**), nobiletin (**57**), β -carotene (**58**), lutein (**59**), lycopene (**60**), cryptoxanthin (**61**), citric acid (**62**), hydroxycinnamic acids (**63**), caffeic acid (**64**), ferulic acid (**65**), chlorogenic acid (**66**), *p*-coumaric acid (**67**), cryptoxanthin (**68**), zeaxanthin (**69**) and β -cryptoxanthin (**70**)

1.7.2.1 *Citrus sinensis* L.

Citrus sinensis L. is a common plant in tropical and subtropical parts of the world. It is an evergreen tree growing up to 9-10 meters and have spines on branches. Leaves are elliptical oval to oblong in shape with narrowly winged-petioles measuring 3-5 mm wide, 6.5-15 cm long with an alternate position. The leaves contain oil gland. Axillary flowers include 20-25 yellow stamens and five white petals. The fruit is ball-shaped to oval (6.5 to 9.5 cm wide) and when mature it becomes orange or yellow. Fruits are sweet orange with a solid centre (Flamini *et al.*, 2003; Sharon-Asa *et al.*, 2003; Connell, 2008; Goudeau *et al.*, 2008; Steduto *et al.*, 2012; Ventura *et al.*, 2012).

Origin and distribution:

Citrus sinensis is a tropical plant; Brazil and the USA are considered as the countries of its origin (Véronique Zech-Matterne (Éditeur), 2018). The plant's geographical distribution is shown in Table 1.3.

Table 1.3: Geographical distribution of *Citrus sinensis*

Country	Reference
Brazil	Kundsen <i>et al.</i> , 2011
China	Niu <i>et al.</i> , 2009
Czech	Hillebrand <i>et al.</i> , 2004; Milind <i>et al.</i> , 2012
Germany:	Mirhosseini <i>et al.</i> 2008; Naila <i>et al.</i> , 2014
India	Hillebrand <i>et al.</i> , 2004; Gattuso <i>et al.</i> , 2007
Italy	Hillebrand <i>et al.</i> , 2004; Gattuso <i>et al.</i> , 2007
Iraq	Al-Snafi, 2016
Japan	Matsubara <i>et al.</i> , 1991
Mexico	Favela-Hernández <i>et al.</i> , 2016
Pakistan	Stöggl <i>et al.</i> , 2006; Intekhab and Aslam, 2009;
Spain	Kanaze <i>et al.</i> , 2009; Barreca <i>et al.</i> , 2014
Sweden	Kolhed and Karlberg, 2005
Turkey	Gómez-Ariza <i>et al.</i> , 2004; Selli, <i>et al.</i> , 2008
USA	Moore <i>et al.</i> , 2001; Abbate <i>et al.</i> , 2001; Peterson <i>et al.</i> , 2006; Saleem <i>et al.</i> , 2010

Traditional uses of Citrus sinensis:

Citrus sinensis can be found widely throughout the world as a good source of vitamin C, which is the famous antioxidant compound that actively participates in building the immune system of the person (Etebu and Nwauzoma, 2014). It has been utilised generally to treat sicknesses like stoppage, cramps, colic, bronchitis, tuberculosis, hack, cold and flu, diarrhoea, obesity, menstrual disorder, angina, hypertension, nervousness, depression and stress (Milind and Dev, 2012).

Phytochemicals of Citrus sinensis

Citrus sinensis is a treasure trove of secondary metabolites and have different biological activities. Several nutritional elements, such as potassium, magnesium, calcium and sodium (Zhao, 2009) have been identified in this species. Major secondary metabolites present in this species are flavonoids, such as kaempferol (**71**) and its derivatives, naringin (**51**), diosmetin (**72**), and daidzein (**73**) (Figure 1.11) (Hillebrand *et al.*, 2004; Gattuso *et al.*, 2007; Favela-Hernández *et al.*, 2016); coumarins like scoparone (**75**), xanthotoxin (**76**), bergapten (**43**), isopimpinellin (**44**), rutamarin (**47**) and triterpene friedelin (**77**) (Figure 1.11) (Gil-Izquierdo *et al.*, 2001; Ribeiro *et al.*, 2008; Dugrand-Judek *et al.*, 2015); carbohydrates, *e.g.*, sucrose (**78**), fructose (**79**), glucose (**80**) and galactose (**81**) (Figure 1.11) (Kolhed *et al.*, 2005); others including steroids, β -sitosterol (**82**), fatty acids (Rani *et al.*, 2009), peptides (Matsubara *et al.*, 1991), carotenoids, such as lutein (**59**) (Aschoff *et al.*, 2015) and essential oils (Gómez-Ariza *et al.*, 2004; Selli *et al.*, 2007). The major components of the essential oils of *C. sinensis* are listed in Table 1.4.

Bioactivity of Citrus sinensis

Citrus sinensis is known to possess various bioactivities. Different extracts of *C. sinensis* showed a wide range of antibacterial activity against *Escherichia coli*, *Pseudomonas aeruginosa*, *Staphylococcus aureus* (Kaviya *et al.*, 2011), *Bacillus subtilis* and *Shigella* (Arooj *et al.*, 2014). According to Trovato *et al.* (1996), *C. sinensis* worked as an antifungal agent by inhibiting the growth of *Candida albicans*. The methanolic extract of this plant

successfully demonstrated its ability to inhibit the malaria parasite *Plasmodium falciparum* (Bhat and Surolia, 2001). *C. sinensis* is considered as one of the most potent plants with antioxidant activity and this activity has been proven in various studies (Atrooz, 2009; Barreca *et al.*, 2014; Garcia *et al.*, 2017). In addition, it has claimed anticancer (Xiao *et al.*, 2009), anti-obesity (Cardile *et al.*, 2015), hypocholesterolaemic (Trovato *et al.*, 1996), relaxant, sedative (Lehrner *et al.*, 2000) and anxiolytic activities (Goes *et al.*, 2012). According to Asgary and Keshvari (2013), *C. sinensis* can reduce the risk of cardiovascular problems. Morrow *et al.* (2009) found a relationship between orange and osteoporosis. When they fed oranges to male rats, they showed improved characteristics in bone structure. *C. sinensis* can be a good candidate to use in sun protection products and may protect from skin cancer, because studies proved that they gave protective activity against ultraviolet rays. Moreover, *C. sinensis* has insect repellent activity (Rossi and Palacios, 2013).

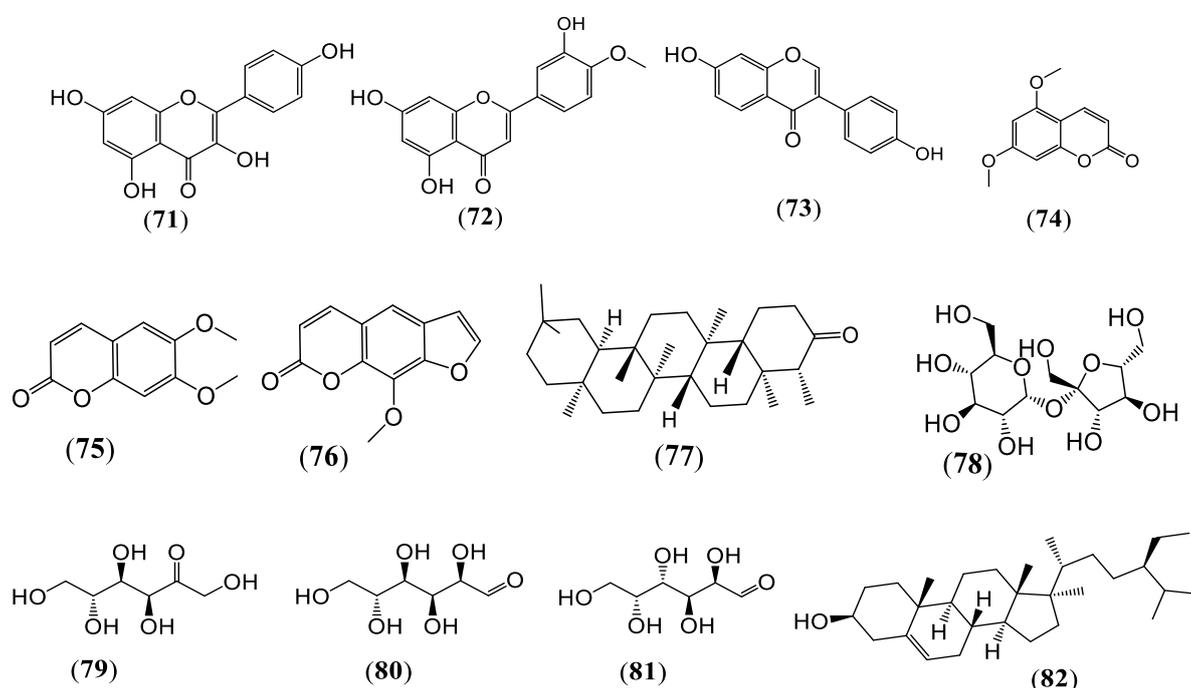


Figure 1.11: Structures of compounds 71-82

Kaempferol (71), diosmetin (72), daidzein (73), limettin (74), scoparone (75), xanthotoxin (76), triterpene friedelin (77), sucrose (78), fructose (79), glucose (80), galactose (81) and β -sitosterol (82)

Table 1.4: Compounds identified from the essential oils of *Citrus sinensis*

Chemical compounds	Reference	
4-Acetyl-1-methylcyclohexene	Selli and Kelebek, 2011	
(-)-Carvone		
<i>p</i> -Coumaric acid		
α -Caryophyllene		
3-Carene		
Dimethyl trisulfide		
Ethyl ethanoate	Gómez-Ariza <i>et al.</i> , 2004; Gómez-Ariza <i>et al.</i> , 2005; Favela-Hernández <i>et al.</i> , 2016	
β -Elemene	Selli and Kelebek, 2011	
Farnesol		
Geraniol		
Geranyl acetate		
Geranyl pyrophosphate		
Guaiacol		
Homofuraneol		
Limonene		
Linalool		Gómez-Ariza <i>et al.</i> , 2004
Linalyl acetate	Selli and Kelebek, 2011	
Malic acid		
3-Mercapto-2-butanone		
Myrecene		
<i>N</i> -Methyltyramine		
3-Methyl-1-pentanol		
Octopamine		
1-Octanol		
(<i>E</i>)- β -Ocimene		
2-Phenylethanol		
α -pinene		
Sabinene		
Sinensal		
Synephrine		
Terbutaline		
Terpinen-4-ol		
α -Terpinene	Gómez-Ariza <i>et al.</i> , 2004	
γ -Terpinene	Selli and Kelebek, 2011	
Tyramine		
Valencene		
Vanillin		

1.7.2.2 *Citrus grandis* Merr.

Citrus grandis is a perennial tree, with a height of around 5-15 m and has , big yellow round pear-shaped fruit of about 15 cm in diameter. The tree needs around 180-400 days from flowering to the mature fruit stage (Ecocrop, 2018).

Original and distribution:

Citrus grandis is considered as the first ancestor of all the *Citrus* fruits. It originated in Malaysia and the Malay Archipelago and has tropical distribution (Giovanni and Angelo, 2002). Table 1.5 presents the global distribution of the plant.

Table 1.5: Global distribution of *Citrus grandis*

Country	References
America	Sawamura and Kuriyama, 1988
Bangladesh	Orwa <i>et al.</i> , 2009, Singh <i>et al.</i> , 2015
Cambodia	Orwa <i>et al.</i> , 2009
Chile	Orwa <i>et al.</i> , 2009
China	Shih-Chen, 1973; Vinning and Moody, 1997; Zhang and Shaolin, 2000
Haiti	Liogier, 1974
India	Orwa <i>et al.</i> , 2009
Indonesia	Orwa <i>et al.</i> , 2009
Japan	Mokbel and Hashinaga, 2006
Keyna	Njoroge <i>et al.</i> , 2005
Laos	Orwa <i>et al.</i> , 2009
Malaysia	Hameed <i>et al.</i> , 2008
Philippines	Orwa <i>et al.</i> , 2009
South Korea	Lim <i>et al.</i> , 2006; Kim <i>et al.</i> , 2009
Taiwan	Wu <i>et al.</i> ,1983; Kuo <i>et al.</i> , 2017
Turkey	Steinmetz,1975
Venezuela	Gonzalez <i>et al.</i> , 2002
Vietnam	Orwa <i>et al.</i> , 2009

Traditional uses of Citrus grandis

Citrus grandis is one of the oldest traditional medicines, which has been used for treating different types of diseases such as oedema, abdominal pain and stomachache. In addition, it is taken orally to prevent or to treat throat pain, fever, anorexia and to cure body

weakness (Rahmatullah *et al.*, 2011). The fruit is used as a cardiac stimulant, antitoxic and to treat stomachache (Arias and Ramón-Laca, 2005). The pulp of *C. grandis* is used as an appetizer (Arias and Ramón-Laca, 2005). The flesh of the fruits is separated from the skin parts and eaten with or without sugar (Lim *et al.*, 2006; Kim *et al.*, 2010). This plant is a part of Chinese medicine for healing cold and to relieve fatigue (Taiping and Shaolin, 2000). In South Korea, leaves are used as a food flavouring and used in making tea (Kim *et al.*, 2010).

Phytochemicals of Citrus grandis

Important phytochemical constituents of *C. grandis* are flavonoids (Xi *et al.*, 2014), which are present in high amounts in fruit juice in the form of neohesperidin (**52**), hesperidin (**50**), naringenin (**83**), and naringin (**51**) (Kanes *et al.*, 1993; Kawaii *et al.*, 1999; Xi *et al.*, 2014). In addition, a high level of flavanones, flavones and flavonols are reported to be present in both the free and glycosidic forms (Kanes *et al.*, 1993; Bocco *et al.*, 1998; Kim *et al.*, 2009). On the other hand, most of the studies have mentioned that *C. grandis* contains coumarins and furocoumarins (Kanes *et al.*, 1993; Bocco *et al.*, 1998; Russo *et al.*, 2016). This plant also has sterols and acridone alkaloids, *e.g.*, grandisinine (**84**), prenylcitpressine (**85**), glycocitrine-I (**86**), citpressine-I (**87**), citrusinine-I (**88**), 5-hydroxynoracronycine (**89**), citracridone-I (**90**), *N*-methylalanine (**91**), preskimmianine (**92**), xanthyletin (**93**), xanthoxyletin (**94**), clausarin (**96**) and *p*-hydroquinone (**97**) (Figure 1.12) (Tian-Shung *et al.*, 1983; Mokbel and Sukanuma, 2006). The plant is reported to contain limonoid (**98**) (Boo *et al.*, 2007) and volatile oils (Njoroge *et al.*, 2005) (Table 1.6).

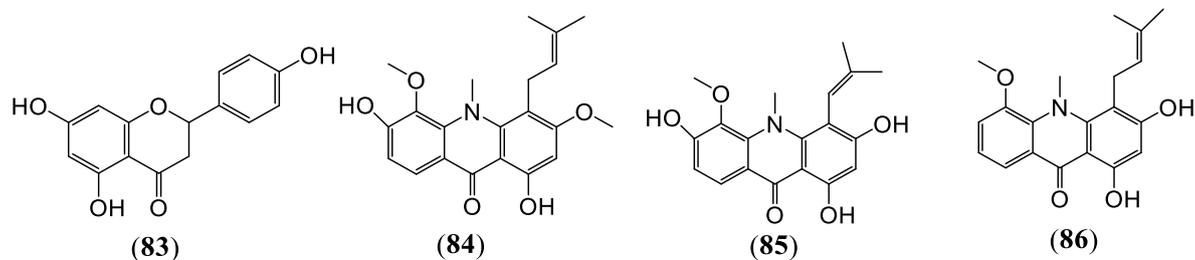


Figure 1.12: Structures of compounds 83-98

Naringenin (**83**), grandisinine (**84**), prenylcitpressine (**85**), glycocitrine-I (**86**), citpressine-I (**87**), citrusinine-I (**88**), 5-hydroxynoracronycine (**89**), citracridone-I (**90**), *N*-methylalanine (**91**), preskimmianine (**92**), xanthyletin (**93**), xanthoxyletin (**94**), clausarin (**96**), *p*-hydroquinone (**97**) and limonoid (**98**)

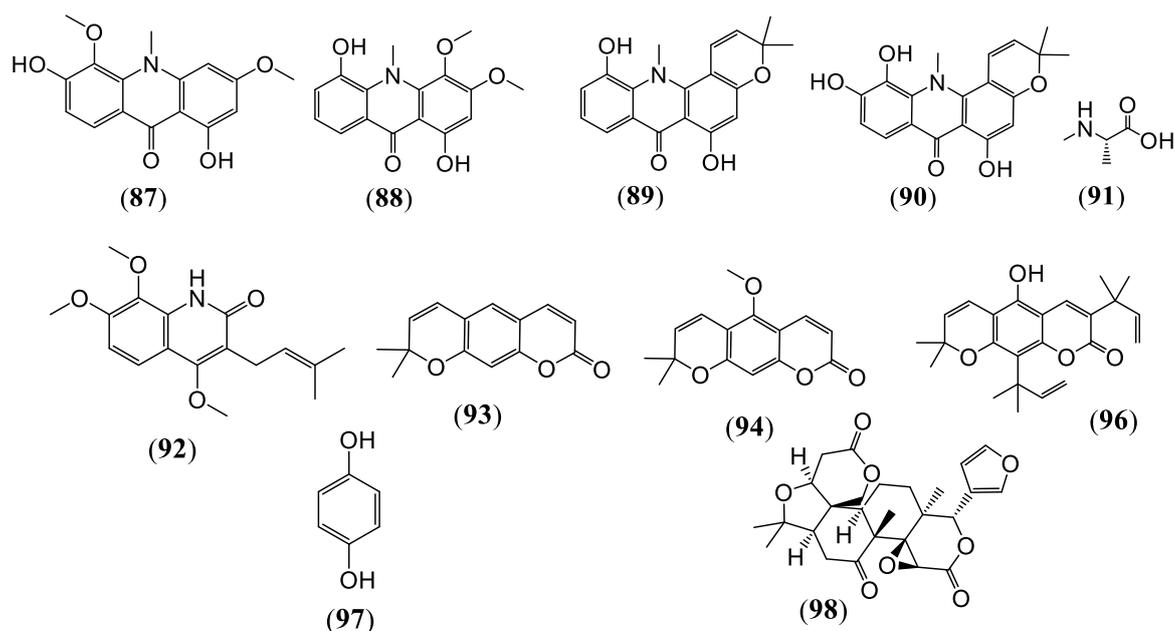


Figure 1.13 (Continued): Structures of compounds 83-98

Naringenin (**83**), grandisinine (**84**), prenylcitpressine (**85**), glycoctrine-I (**86**), citpressine-I (**87**), citrusinine-I (**88**), 5-hydroxynoracronycine (**89**), citracridone-I (**90**), *N*-methylalanine (**91**), preskimmianine (**92**), xanthyletin (**93**), xanthoxyletin (**94**), clausarin (**96**), *p*-hydroquinone (**97**) and limonoid (**98**)

Table 1.6: Compounds identified in essential oils of *Citrus grandis*

Compounds *	Compounds*	Compounds*
(<i>E</i>)-Carveol	Dodecanal	Neral
(<i>Z</i>)-Carveol	Elemol	Nerol
(<i>E</i>)-Carvone	β -Elemene	Nerolidyl acetate
(<i>Z</i>)-Carvone	(<i>E,E</i>)- α -Farnesene	Nonanal
Caryophyllene oxide	Geranial	Nonanoic acid
β -Caryophyllene	Heptyl acetate	α -Pinene
(<i>E</i>)-Cryophyllene epoxide	Limonene	β -Pinene
Cedrol	Limonen-10-ol	(<i>Z</i>)-Piperitol
α -Cedrene	Linalool	Sabinene
Citronellal	Linalyl acetate	Sabina ketone
α -Copaene	Isothujol	β -Sinensal
α -Cubebene	Methyl- <i>N</i> -methylantranilate	α -Terpinene
Cumin aldehyde	<i>p</i> -Mentha-1-en-9-ol	γ -Terpinene
Decanal	(<i>E</i>)- <i>p</i> -2,8-Menthadien-1-ol	α -Terpineol
Dihydrocarveol	(<i>E</i>)- <i>p</i> -Mentha-2,8-dienol	Tetradecenal

*(Njoroge *et al.*, 2005)

Bioactivity of Citrus grandis

Citrus grandis shows potentially useful effects on human health. The antiglycation effect of *C. grandis* extract (0.25-2.00 mg/mL) was approved against fructose-mediated protein oxidation and glycation (Caengprasath *et al.*, 2013) by controlling the blood glucose level of diabetic patients (Kim *et al.*, 2009). Various studies have proven the antioxidant (Lim *et al.*, 2006; Mokbel and Hashinaga, 2006), anti-inflammatory, antimicrobial activity (Kuo *et al.*, 2017) and anticancer activity (Kim *et al.*, 2010). It is also used in cosmetics.

A few studies have focused on the biological effects and chemical composition of *C. grandis* leaves (Kim *et al.*, 2009; Lim *et al.*, 2009). However, the anticancer effect of *C. grandis* fruit giving inhibition of human leukaemia cells (U937) was previously reported, and the result obtained indicated the hexane extract inhibited the cancer cells with IC₅₀ 60 µg/mL (Lim *et al.*, 2009). Another report showed the anticancer effect of *C. grandis* leaves against human gastric cancer cells (SNU-16), the inhibitory effect of the chloroform extract was IC₅₀ 92.15 µg/mL (Yong Moon *et al.*, 2009).

1.8 Aims and objectives

The aim of this work was to isolate and characterise the antimicrobial compounds present in three Iraqi medicinal plants from the family Rutaceae: *Citrus grandis*, *Citrus sinensis* and *Ruta chalepensis*. To achieve this aim the following objectives were set:

1. to prepare three types of crude extracts from ground plant material of every plant by successive Soxhlet extraction using the solvents, *n*-hexane, DCM and MeOH;
2. to assess the antimicrobial property of these extracts against various strains of Gram-positive, Gram-negative bacteria and fungi;
3. to follow bioassay-guided fractionation by vacuum liquid chromatography (VLC) and/or solid phase extraction (SPE) leading to the identification of active extracts for isolation of active compounds;

4. to isolate and identify the active compounds using chromatographic and spectroscopic techniques;
5. to determine the minimum inhibitory concentration (MIC) of the active compounds using the resazurin 96-well assay;
6. to carry out anti-MRSA activity testing against various clinical isolates of MRSA strains;
7. to try to understand possible mechanism of actions of active compounds.

Chapter 2
Materials and Methods

2.1 Plant materials

Three Iraqi medicinal plants from the family Rutaceae, *Citrus grandis*, *C. sinensis* and *Ruta chalepensis* (Table 2.1), were selected on the basis of their traditional uses to treat infections, and collected from Diyala, central Iraq at longitude 33.79684° N, and latitude 44.623337° E (Figure 2.1). Plant material were collected in September 2015 and identified in comparison with the voucher specimens held at the National Herbarium of Iraq, where appropriate voucher specimens of this collection were deposited (Table 2.1).

Table 2.1: Plant used in study

Plant name	Voucher number
<i>Citrus grandis</i>	15324
<i>Citrus sinensis</i>	6534
<i>Ruta chalepensis</i>	33396



Figure 2.1: Iraqi political map

2.2. Materials and reagents for phytochemical studies

To carry out the phytochemical analysis of the selected plants, a range of instruments, materials and various chemicals have been used. Table 2.2 summarises instrument, materials and reagents for the plant extract preparation and processing.

Table 2.2: Materials and reagents used in phytochemical work

	Name	Details
Materials	Rotary evaporator	Cole-Parmer, UK
	Strata C-18-E cartridge	Phenomenex, California-USA
	Freeze dryer	Telstar, UK
	Silica gel	60 H, Merck, Germany
	Solid phase extraction cartridge	Strata C18-E (55 μm , 70 \AA), 20 g/60 mL, Giga tubes, Phenomenex, USA
	TLC plates	Silica gel on Aluminium foils, Sigma-Aldrich, Germany
	Oven	Sciquip, Shrewsbury-UK
	UV lamps	Camag, Switzerland
	Analytical HPLC	Agilent Technologies, 1200 Infinity Series, Germany Dionex Ultimate 300 diode array detector, Thermo Scientific, Germany.
	Preparative HPLC	Agilent Technologies, 1200 Infinity Series, Germany.
Columns	Described in point 2.3.3.4	
Nuclear Magnetic Resonance Spectrometers	Bruker AMX 300 MHz and 600 AMX MHz, Germany	
Solvents	<i>n</i> -Hexane	Fisher Scientific, UK
	Ethyl acetate (EtOA)	
	Dichloromethane (DCM)	
	Chloroform (CHCl ₃)	
	Methanol (MeOH), HPLC grade	
	Water (H ₂ O), HPLC grade	
	Chloroform- <i>d</i> ₃ (CDCl ₃)	Cambridge Isotope Laboratories, Inc., USA
	Dimethyl sulfoxide- <i>d</i> ₆ (C ₂ D ₆ OS)	
Methanol- <i>d</i> ₄ (CD ₃ OD)		
Reagents	Anisaldehyde	Sigma-Aldrich, UK

2.3 Phytochemical work

2.3.1 Plant preparation

The plant parts were cleaned, sectioned and shade-dried at room temperature. The dried parts were ground in a grinder to prepare the powdered material for Soxhlet extraction.

2.3.2 Soxhlet extraction

Extraction of a plant material is the most important initial step in phytochemical and bioactivity processing. Selection of a suitable solvent and extraction technique is necessary to get an optimum extraction yield.

The Soxhlet extraction technique used for extraction of small to moderate volumes of plant materials, consumes less solvent and time. Furthermore, this extraction technique is a continuous extraction which can use solvents of different polarities or mixed solvents. Selection of a suitable solvent and use of reasonable continuous heat is helpful to obtain different types of extracts with high yields (Sarker and Nahar, 2012; Rostagno and Prado, 2013).

Powdered plant material was packed in a thimble made up of filter paper and loaded in the main chamber of a Soxhlet apparatus. The distillation flask filled with an organic solvent was placed on a heating mantle. The Soxhlet chamber along with the condenser was fitted on top of the flask. During each cycle, the chamber gradually filled with warm solvent and after reaching a certain point, the siphon side arm immediately discharges the solvent back down to the distillation flask. The compounds of interest dissolved in that warm solvent were concentrated in the distillation flask (Figure 2.2) (De Castro and Priego-Capote, 2010).

In the present study, the successive extractions were performed using three different solvents of increasing polarity, *viz* *n*-hexane, DCM and MeOH. Ten cycles were allowed for each extraction. The obtained extracts were concentrated to dryness using a rotary evaporator and stored at 4°C for further usage.

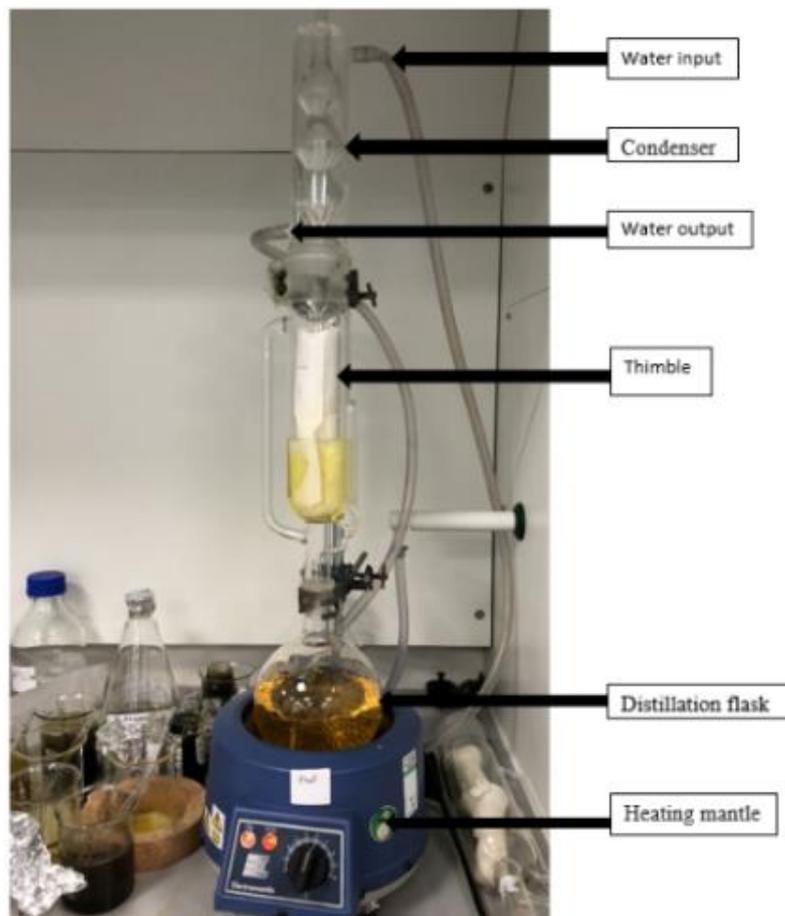


Figure 2.2: Soxhlet apparatus

2.3.3 Chromatographic techniques

Chromatography is an analytical technique, which can be used to separate mixture of compounds into individual components. It is used to fractionate a crude extract into different discrete fractions depending on the polarity. Application of a single fractionation technique is usually inadequate for isolation of a single compound from a crude extract. Therefore, different chromatographic techniques were used in this project depending on the polarity of the extracts.

2.3.3.1 Vacuum liquid chromatography (VLC)

Vacuum liquid chromatography (VLC) is a convenient and modified column chromatography method of fractionation on a Silica gel bed as described by Pelletier *et al.* (1986). The VLC method consisted of a sintered glass Buchner filter funnel connected via the flask side arm to a vacuum system (Figure 2.3). The stationary phase on VLC was the

Buchner filter funnel filled with two-thirds of dry silica gel 60H, rinsed with solvent and a vacuum applied to compress the surface and to allow the silica to pack. Selection of eluting solvent usually depends on the solvent that has been used during sample extraction (for example if the sample is a *n*-hexane extract, the rinsing solvent will be *n*-hexane) (Reid and Sarker, 2012). The preparation of sample for the VLC method required the sample to be dissolved in a small amount of the appropriate solvent and then mixed with normal silica gel (70-230 mesh). The mixture was dried and loaded as a uniform thin layer on the top of the funnel. The mobile phase was mixtures of various solvents of increasing polarity and was run under vacuum. The eluted fractions were collected in flasks for further work (Table 2.3). All fractions were evaporated to dryness using a rotary evaporator and analysed by TLC.

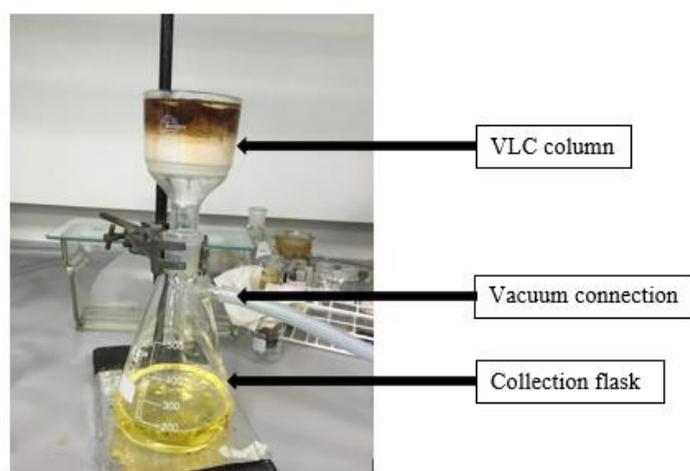


Figure 2.3: Vacuum liquid chromatography (VLC)

2.3.3.2 Solid-phase extraction (SPE)

Solid-phase extraction (SPE) is an economical and sometimes automated chromatographic technique used to fractionate liquid mixtures. The SPE technique was very similar to VLC with the only difference being the stationary phase, which was reversed-phase silica (C₁₈). Also, in SPE the mobile phase was a mixture of solvents of decreasing polarity (Sarker *et al.*, 2006). Both SPE and VLC operate under vacuum.

A portion of active methanolic extract was dissolved or suspended in 10 mL of 10% MeOH in H₂O and applied to the cartridge, which was previously washed with 50 mL of MeOH and then equilibrated with 100 mL of H₂O sequentially (Figure 2.4). A step gradient protocol was applied with 200 mL of different concentrations of MeOH in water (Table 2.3). Four fractions were collected, dried using a rotary evaporator along with freeze drying and kept for further work (Sarker *et al.*, 2006).

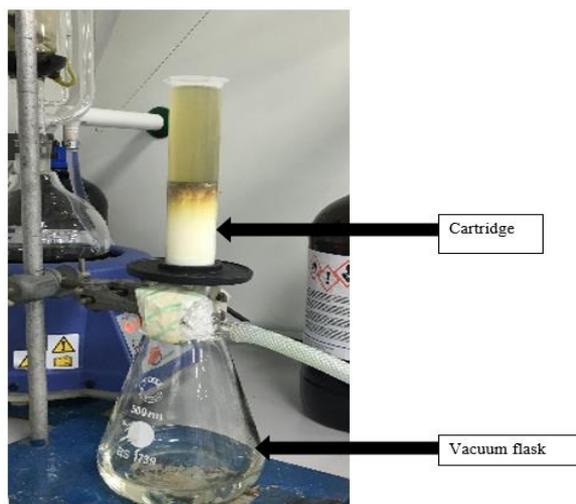


Figure 2.4: Solid-phase extraction (SPE)

2.3.3.3 Thin layer chromatography (TLC)

Thin layer chromatography (TLC) is one of the most simple, easy and economical techniques used to determine and monitor compounds during the separation process depending on polarity, absorption and binding to silica. In the current study, this method was used to analyse the secondary metabolites in plant extracts and check the purity of the isolated compounds.

Furthermore, preparative-TLC (PTLC) was applied as an isolation technique for the purification of nonpolar compounds. The TLC plates used in this study were characterised as precoated silica gel 60 PF₂₅₄ aluminium plates with thickness range of 0.5-2.0 mm.

Table 2.3: Methods used in VLC and SPE techniques

Method	Type and method of extracts	Fractions										
		F1	F2	F3	F4	F5	F6	F7	F8	F9	F10	F11
A	<i>n</i> -hexane /VLC	100% hex	10% Et/hex	20% Et/hex	40% Et/hex	60% Et/hex	80% Et/hex	100% Et	_____	_____	_____	_____
B	<i>n</i> -hexane /VLC	100% hex	5% Et/hex	10% Et/hex	15% Et/hex	20% Et/hex	50% Et/hex	80% Et/hex	100% Et	50% Et/Ch	100% Ch	2% Me/Ch
C	DCM/VLC	100% Ch	2% Ch/Me	4% Ch/Me	6% Ch/Me	8% Ch/Me	10% Ch/Me	20% Ch/Me	30% Ch/Me	_____	_____	_____
D	MeOH/SPE	20% Me/H	50% Me/H	80% Me/H	100% Me	_____	_____	_____	_____	_____	_____	_____

hex: *n*-hexane; Me: MeOH, Et: EtOAc; Ch: CHCl₃; H: H₂O

The processing of a TLC plate involves spotting of the samples on the line around 1.5 cm from the bottom. Then, the plate is transferred to a glass chromatographic chamber containing developing solvent. As the solvent goes up through the TLC plate by the capillary action, the component's spots start developing. In this work, the procedure described above was followed, the TLC plates were dried and visualised under short wave length (254 nm) and long wave length (366 nm) using UV lamps to check the best separation for the target compounds. Anisaldehyde solution (1%) in aqueous H₂SO₄ was sprayed over the plate which was then exposed to oven heating at 105 °C for 5 min to detect the spots, which were not detected by UV lights (Gibbons, 2012).

PTLC is a common and most basic method available around the world for compound isolation. This plate can be made from silica, alumina, C₁₈ and cellulose, and have thickness range of 0.5-2.0 mm. It should be homogenous to get a good separation. 10-100 mg samples can separated with PTLC (Hostettmann *et al.*, 1986). Table 2.4 below shows the different methods used to analyse and isolate compounds using TLC and PTLC.

Table 2.4: Thin layer chromatographic methods

Methods	Extracts type	Mobile phase
E	<i>n</i> -hexane extracts	10% EtOAc / <i>n</i> -hexane
F	DCM extracts	5% MeOH/CHCl ₃
G	DCM extract	100% MeOH
H	Plant extracts	100% DCM
I	RFH4	15% EtOAc/ <i>n</i> -hexane
J	RFF2-11	20 % EtOAc/ <i>n</i> -hexane
K	F11 from method B	25% EtOAc/ <i>n</i> -hexane

2.3.3.4 High performance liquid chromatography (HPLC)

High performance liquid chromatography (HPLC) is one of the analytical and separation techniques that has been used for determination, separation and purification of compound mixtures.

Analytical high performance liquid chromatography

Analytical HPLC was used in this study for detection and preliminary analysis of secondary metabolites in plant extracts and to develop methods for preparative-HPLC. This work was carried out using a Dionex 3000 or an Agilent 1260, both instruments had a binary pump, an autosampler, a column chamber, a degasser and a UV/DAD detector. Before starting, the column was washed with 100% MeOH, 50%-50%, 30%-70% MeOH/water respectively, to wash out all unwanted contaminants. The analytical column used in the current study was a Phenomenex Gemini-NX 5 U C₁₈ column (150 × 4.6 mm, 5 μm, Phenomenex, USA) or a Hypersil 5 U C₁₈ column (150 × 4.6 mm, 5 μm, Phenomenex, USA) and the column temperature was set at 25 °C. The samples were dissolved in MeOH (1 mg/mL) and 10 μL was injected with a flow rate of 1 mL/min and the chromatograms were monitored by variable UV/vis wavelengths between 205 and 366 nm.

Preparative high performance liquid chromatography

Reversed-phase preparative HPLC was used for separation and purification of compounds following the developed methods from analytical HPLC. This work was accomplished on an Agilent 1260 infinity series include a binary pump, a degasser, a UV/DAD detector and a column chamber set at on 25°C. In the case of isolation of compounds from plant extracts, a Phenomenex column LC-18 stainless steel column (150 x 21.2 mm, 5 μm particle size, Phenomenex, USA) was used with a flow rate of 10 mL/min. For the purification of impure compounds or compound mixtures, a Phenomenex semi-preparative column, LC-18 stainless steel (150 x 10 mm, 5 μm particle size, Phenomenex, USA) was used with a flow rate of 2 mL/min. The plant samples were dissolved in MeOH (10 mg/mL) and the HPLC solvents H₂O, MeOH, ACN containing 0.1% TFA. Table 2.5 summarizes the developed methods used in HPLC.

Table 2.5: HPLC methods used in the current study

Methods	Description (All mobile phases containing 0.1% TFA)
L	Linear gradient 30-100% MeOH in water over 30 min, isocratic 100% MeOH for the next 10 min, linear gradient 100-30% MeOH in water for 5 min, and isocratic 30% MeOH in water for 5 min
M	Linear gradient 50-80% MeOH in water over 30 min, isocratic 80% MeOH for the next 5 min, linear gradient 80-30% MeOH in water for 5 min, and isocratic 30% MeOH in water for 5 min
N	Linear gradient 30-100% ACN in water over 30 min, isocratic 100% ACN for the next 5 min, linear gradient 100-30% ACN in water for 5 min, and isocratic 30% MeOH in water for 5 min
O	Linear gradient 20-80% MeOH in water over 30 min, isocratic 80% MeOH for the next 5 min, linear gradient 80-20% MeOH in water for 5 min, and isocratic 20% MeOH in water for 5 min
P	Linear gradient 30-100% MeOH in water over 20 min, isocratic 100% MeOH for the next 5 min, linear gradient 100-30% MeOH in water for 5 min, and isocratic 30% MeOH in water for 5 min
Q	Linear gradient 10-90% ACN in water over 30 min, isocratic 100% ACN for the next 5 min, linear gradient 100-30% ACN in water for 5 min, and isocratic 30% ACN in water for 5 min
R	Linear gradient 20-80% MeOH in water over 30 min, isocratic 80% MeOH for the next 2 min, linear gradient 80-20% MeOH in water for 2 min, and isocratic 100% MeOH in water for 2min, and isocratic 20% MeOH in water for 5 min
S	Linear gradient 20-50% ACN in water over 30 min, isocratic 50% ACN for the next 3 min, linear gradient 50-20% ACN in water for 5 min, and isocratic 20% ACN in water for 5 min
T	Linear gradient 20-80% ACN in water over 30 min, isocratic 80% ACN for the next 5 min, linear gradient 80-20% ACN in water for 5 min, and isocratic 20% ACN in water for 5 min
U	Linear gradient 30-100% MeOH in water over 18 min, isocratic 100% MeOH for the next 3 min, linear gradient 100-30% MeOH in water for 3 min, and isocratic 30% MeOH in water for 3 min

2.3.3 Isolation of compounds

2.3.3.1 Isolation of compounds from *R. chalepensis* fruits

The dried and ground *R. chalepensis* fruits (206.1 g) were subjected to successive Soxhlet extraction to obtain three extracts, *n*-hexane (9.0 g), DCM (9.37 g) and MeOH (5.36 g). The dried *n*-hexane and DCM extracts were applied to VLC, resulting in seven and eight fractions, respectively, while, the MeOH extract was fractionated by SPE. Active fractions 4 and 11 of the *n*-hexane extract, fraction 7 of the DCM extract, and fractions 2 and 3 of the MeOH extracts were analysed by different chromatographic methods leading to isolation of compounds. Table 2.6 outlines the isolated compounds from the different extracts and fractions from the fruits of *R. chalepensis*.

2.3.3.2 Isolation of compounds from *R. chalepensis* leaves

The leaves of *R. chalepensis* were dried and ground. A portion (293.31 g) of the plant material was subjected to Soxhlet extraction to obtain different extracts, *n*-hexane (9.06 g), DCM (8.18 g) and MeOH (5.33 g). The DCM and MeOH extracts were fractionated using VLC and SPE, respectively. Both fractions 2 and 3 from the MeOH extract along with fraction 7 of the DCM extract were subjected to reversed-phase preparative HPLC to isolate different compounds. Table 2.7 summarizes the isolated compounds from *R. chalepensis* leaves following different methods.

2.3.3.3 Isolation of compounds from *R. chalepensis* stem

To isolate compounds from *R. chalepensis* stem, powdered material (243.3 g) was extracted to obtain three extracts, *n*-hexane (5.4 g), DCM (3.01 g) and MeOH (5.40 g). Only fractions 2 and 3 from the MeOH extract were submitted to reversed-phase preparative-HPLC to isolate the compounds as listed in Table 2.8 with their retention time.

Table 2.6: Isolated compounds from *R. chalepensis* fruits

Extract	Fraction	Methods	R _f value	Retention time (<i>t</i> _R) in min	Compounds
<i>n</i> -Hexane	F4	I	0.6	-	46
			0.5	-	47
	F11	K	0.20	-	42
			0.19	-	43
DCM	F7	M	0.47	-	99
			-	7.19	107
			-	10.41	105
			-	12.77	99
			-	15.15	42
			-	18.65	41
Methanol	F2	L	-	8.30	106
			-	16.02	48
			-	16.99	101
			-	17.78	103
			-	18.64	105
			-	21.43	44
			-	22.73	42
	RFF2-11	J	-	0.19	43
			-	0.47	99
	F3	L	-	18.50	48
			-	24.67	41
			-	25.15	104
			-	26.37	46
			-	26.92	45
			-	27.94	47

Table 2.7: Isolated compounds from *R. chalepensis* leaves

Extract	Fraction	Methods	Retention time (<i>t_R</i>)	Compounds
DCM	F7	O	13.03	105
			14.70	99
			20.53	107
			24.85	45
Methanol	F2	L	13.34	43
			15.73	48
			20.41	42
			21.28	99
	RLF2-5	U	10.85	101
			11.84	102
	RLF2-7	P	10.52	48
			11.01	100
			14.56	105
	F3	L	12.53	48
			16.05	105
			17.69	43
			18.09	99
			19.03	42
20.05			107	
23.39			45	
24.57	46			

Table 2.8: Isolated compounds from *R. chalepensis* stem

Extract	Fraction	Retention times (<i>t_R</i>)	Compounds
Methanol	F2	11.56	108
		12.5	48
		14.7	102
	F3	20.64	99
		21.11	42
		22.38	43
		23.01	107
		27.03	45
		28.5	46

2.3.3.4 Isolation of compounds from *C. grandis* leaves

The air-dried and powdered leaves (351.0 g) of *C. grandis* were Soxhlet extracted sequentially with solvents of increasing polarity to obtain three extracts, *n*-hexane (5.9 g), DCM (13.5 g) and MeOH (15.9 g). The VLC fractionation of the DCM extract afforded seven fractions. Using the method **T** in reversed-phase preparative-HPLC, six pure compounds were separated from fraction 7 of the DCM extract. Table 2.9 summarizes the separated compounds along with their retention times.

Table 2.9: Isolated compounds from *C. grandis*

Extract	Fraction	Retention time (t_R)	Compounds
DCM	F7	14.58	112
		19.00	111
		20.27	115
		21.19	57
		23.93	114
		28.45	50

2.3.3.5 Isolation of compounds from *C. sinensis* leaves

The air-dried and powdered leaves of *C. sinensis* (194.58 g) were Soxhlet extracted sequentially with solvents of increasing polarity to obtain three extracts, *n*-hexane (6.15 g), DCM (10.2 g) and MeOH (14.3 g). The DCM extract was fractionated using VLC to afford seven fractions. Fraction 7 was subjected to reversed-phase preparative-HPLC using the method **N** to isolate five compounds (Table 2.10).

Table 2.10: Isolated compounds from *C. sinensis* leaves

Extract	Fraction	Retention time (t_R)	Compounds
DCM	F7	20.39	57
		21.36	114
		22.32	112
		31.61	115
		33.45	50

2.3.3.6 Isolation of compounds from *C. sinensis* peels

The Soxhlet extraction of the air-dried and ground peels of *C. sinensis* (393.39) produced three extracts, *n*-hexane (7.34 g), DCM (12.7 g) and MeOH (25.17 g). The DCM and the MeOH extracts were fractionated by VLC and SPE, respectively. Fractions 2 and 3 of the MeOH extract and fractions 6 and 7 of the DCM extract were subjected to reversed-phase preparative-HPLC using different methods to isolate various compounds (Table 2.11).

Table 2.11: Isolated compounds from *C. sinensis* peels

Extract	Fractions	Methods	Retention time (t_R)	Compounds
DCM	F6	S	28.28	111
			32.00	57
			33.21	117
			34.35	110
			37.03	116
			39.84	113
	F7	T	14.01	48
			17.70	51
			19.74	111
			21.75	57
			22.42	117
			23.03	110
			24.33	116
	MeOH	F2	R	13.78
15.03				119
15.50				101
18.28				51
19.18				50
F3		R	17.56	48
			26.13	116
			28.33	110
			28.82	57

2.4 Identification and characterisation of isolated compounds

To determine the molecular structures and chemical formula of the isolated compounds in this study, different spectroscopic methods were used, *e.g.*, ultraviolet (UV), Nuclear Magnetic Resonance (NMR) and Mass spectroscopy (MS).

2.4.1 Nuclear Magnetic Resonance (NMR) spectroscopy

NMR is a powerful spectroscopic tool that provides information to elucidate chemical structures of compounds. The method involves exposing atomic nuclei of a sample to a magnetic field linked with electromagnetic radiation with specific frequency. By detecting the absorption signals, one can acquire an NMR spectrum (Günther, 2013). Based on these signals, a compound structure can be elucidated. Structure can be confirmed by applying various types of NMR experiments 1 dimension (1D) experiments include proton (^1H) and carbon (^{13}C) NMR while 2 dimension (2D) NMR experiments include ^1H - ^1H COSY (Correlation Spectroscopy), DEPT 135 (Distortionless Enhancement by Polarization Transfer), HMBC (^1H - ^{13}C Heteronuclear Multiple Bond Correlation Spectroscopy) and HSQC (^1H - ^{13}C Heteronuclear Single Quantum correlation) (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993). The samples were dissolved in appropriate amounts from deuterated solvents and these solvents were calibrated by specific chemical shifts summarized on Table 2.12. The NMR experiments were performed on a Bruker AMX 300 MHz or AMX 600 MHz.

2.4.1.1 Proton (^1H) NMR spectroscopy

The ^1H (Figure 2.5) is one of the basic, essential and simple experiments providing important information about H number of protons (signals), chemical shift (measured in ppm), proton neighbour, coupling constant and integration. The different signals of protons indicates protons with different magnetic environments. The signals in chemical shift indicates proton position. The neighbouring ions causes shielding (up field) or deshielding (down field) of the protons, which effects its position in the chemical shift. Spin-spin splitting appears of equivalent hydrogens in a compound; the proton can be a single peak (s)

when there is no equivalent hydrogens otherwise it will be doublet (*dd*), triplet (*t*) or multiplet (*m*). The coupling constant (*J* value measured in Hz) is the distance value between the doublet, triplet or multiplied protons. Integration indicates number of hydrogens and is presented under the baseline of spectra (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993).

Table 2.12: Chemical shifts for deuterated solvents used in this study

Solvents	Chemical shifts δ in ppm	
	^1H	^{13}C
Chloroform- <i>d</i> (CDCl_3)	7.26	77.36
Dimethyl sulfoxide- <i>d</i> ₆ ($\text{DMSO-}d_6$)	2.5; 3.3	39.9
Methanol- <i>d</i> ₄ (CD_3OD)	3.33	49.15

2.4.1.2 Carbon (^{13}C) NMR spectroscopy

The ^{13}C NMR (Figure 2.6) is an important experiment to identify all carbon atoms in an organic molecule. This experiment provides the number and types of carbons present in the molecule. The position of the carbon signal by chemical shift helps to identify to which functional groups are connected to carbons. All carbon signals must be in one direction (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993).

2.4.1.3 Distortionless Enhancement by Polarisation Transfer (DEPT-135)

DEPT carbon (Figure 2.7) distinguishes carbons numbers and full analysis of carbon atoms. The carbon signal of methine ($-\text{CH}$), methyl ($-\text{CH}_3$) will be on one side and the methylene ($-\text{CH}_2$) be on the other side (Figure 2.7) (Silverstein and Bassler, 1962).

2.4.1.4 Homonuclear Correlation Spectroscopy (COSY)

This two-dimensional experiment (2D) (Figure 2.8) includes two proton axes, and the cross plot between them determines their correlation presenting the coupling protons on molecule. Thus, the *HH* COSY indicates the connectivities between *HH* to be *geminal*, *vicinal* or *w*-relationships of the H atoms of a molecular and the associated structural units (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993).

2.4.1.5 Heteronuclear Single Quantum Coherence Spectroscopy (HSQC)

HSQC (Figure 2.9) is a two-dimensional experiment (2D) revealing the direct correlation from proton to attached carbon. The HSQC has the ^1H NMR spectrum on top axis and ^{13}C NMR spectrum on side axis. The HSQC identifies protons which are coupled to carbons (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993).

2.4.1.6 Heteronuclear Multiple Bond Correlation (HMBC)

HMBC is powerful experiment that shows correlations between carbons and hydrogens that are separated by two or more chemical bonds. HMBC is used to assign the quaternary and carbonyl carbons (Figure 2.10) (Silverstein and Bassler, 1962; Breitmaier and Sinnema, 1993).

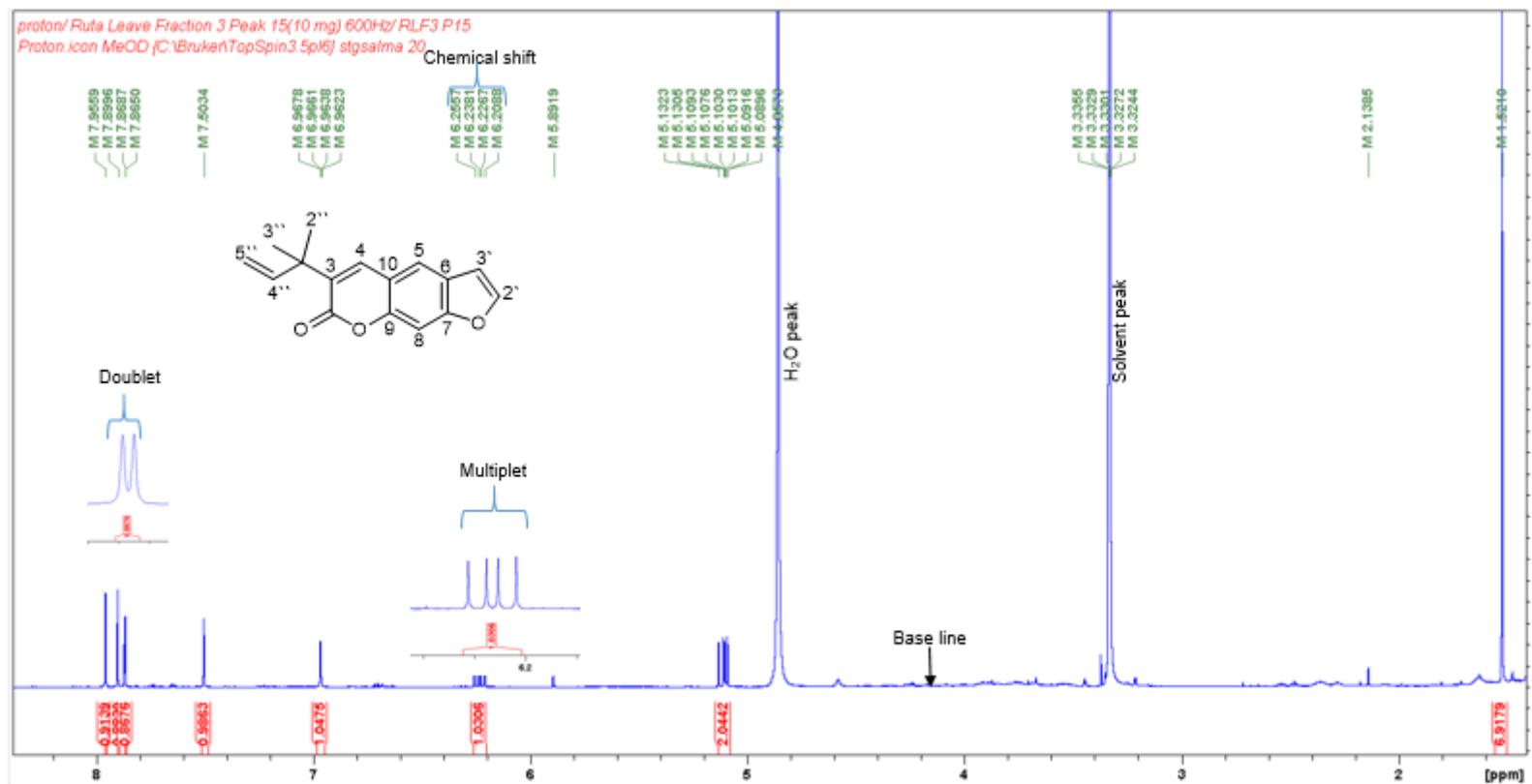


Figure 2.5: Proton (^1H) NMR experiment

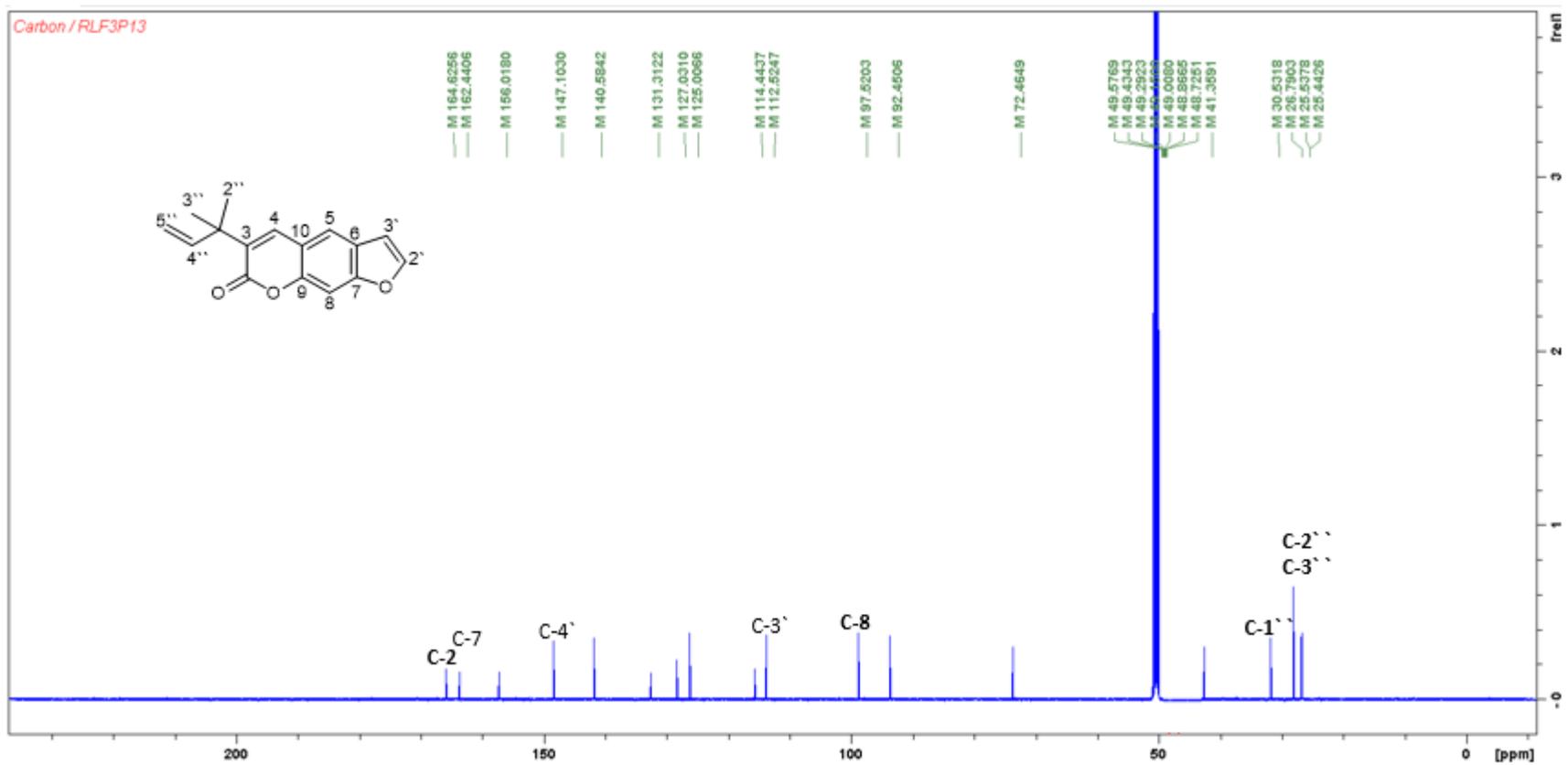


Figure 2.6: Carbon (^{13}C) NMR experiment

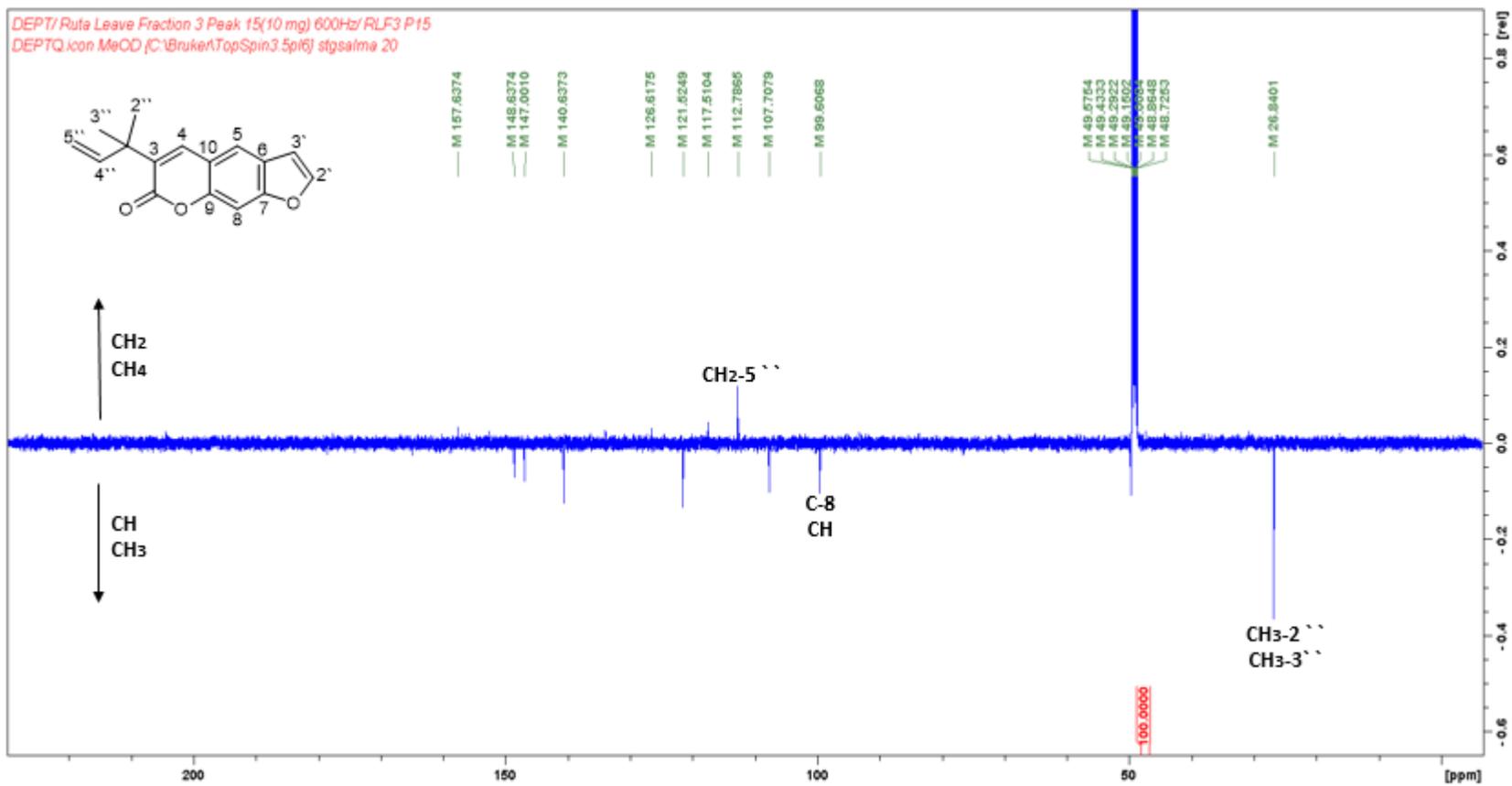


Figure 2.7: DEPTQ NMR experiment

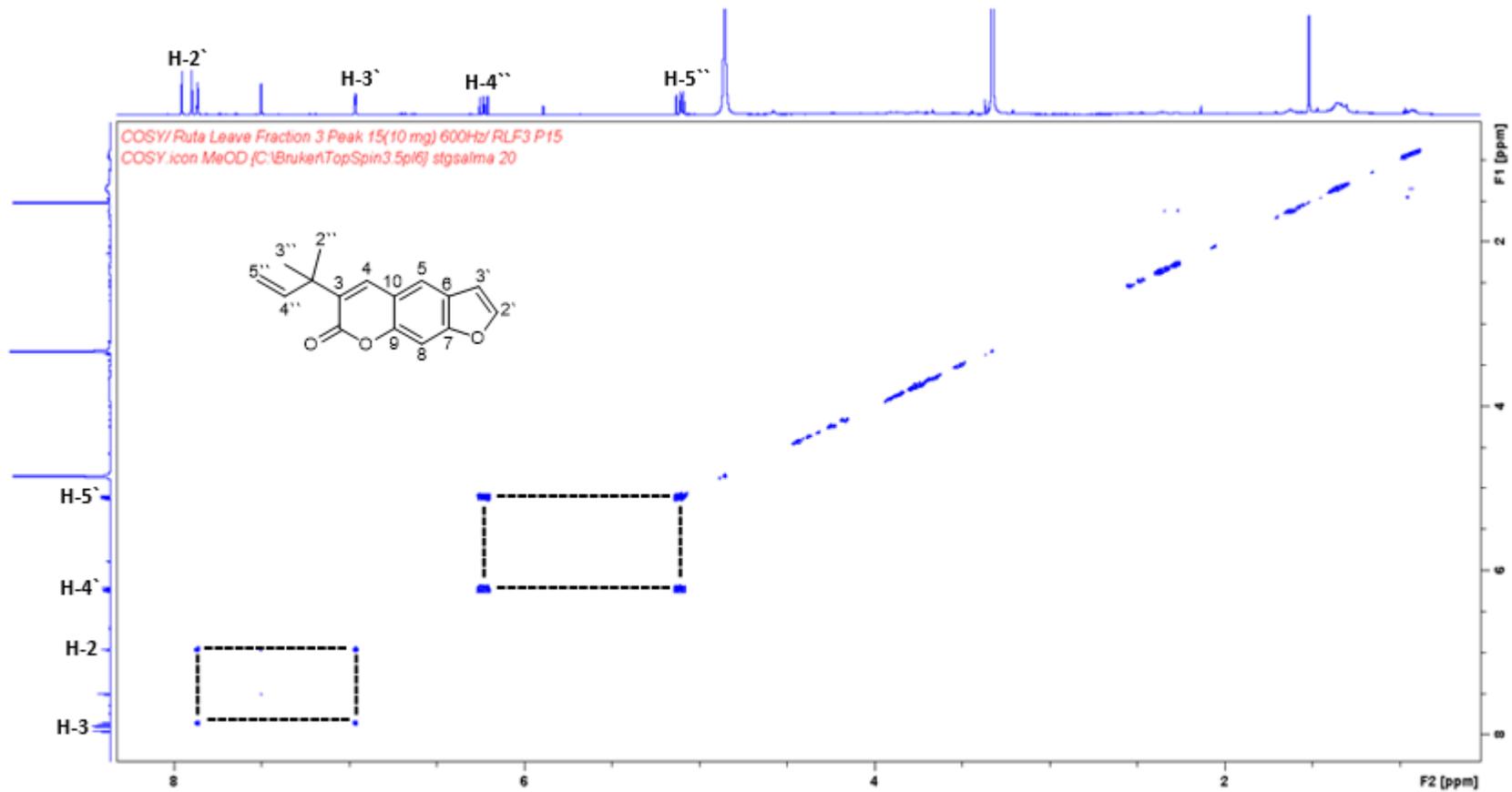


Figure 2.8: COSY NMR experiment

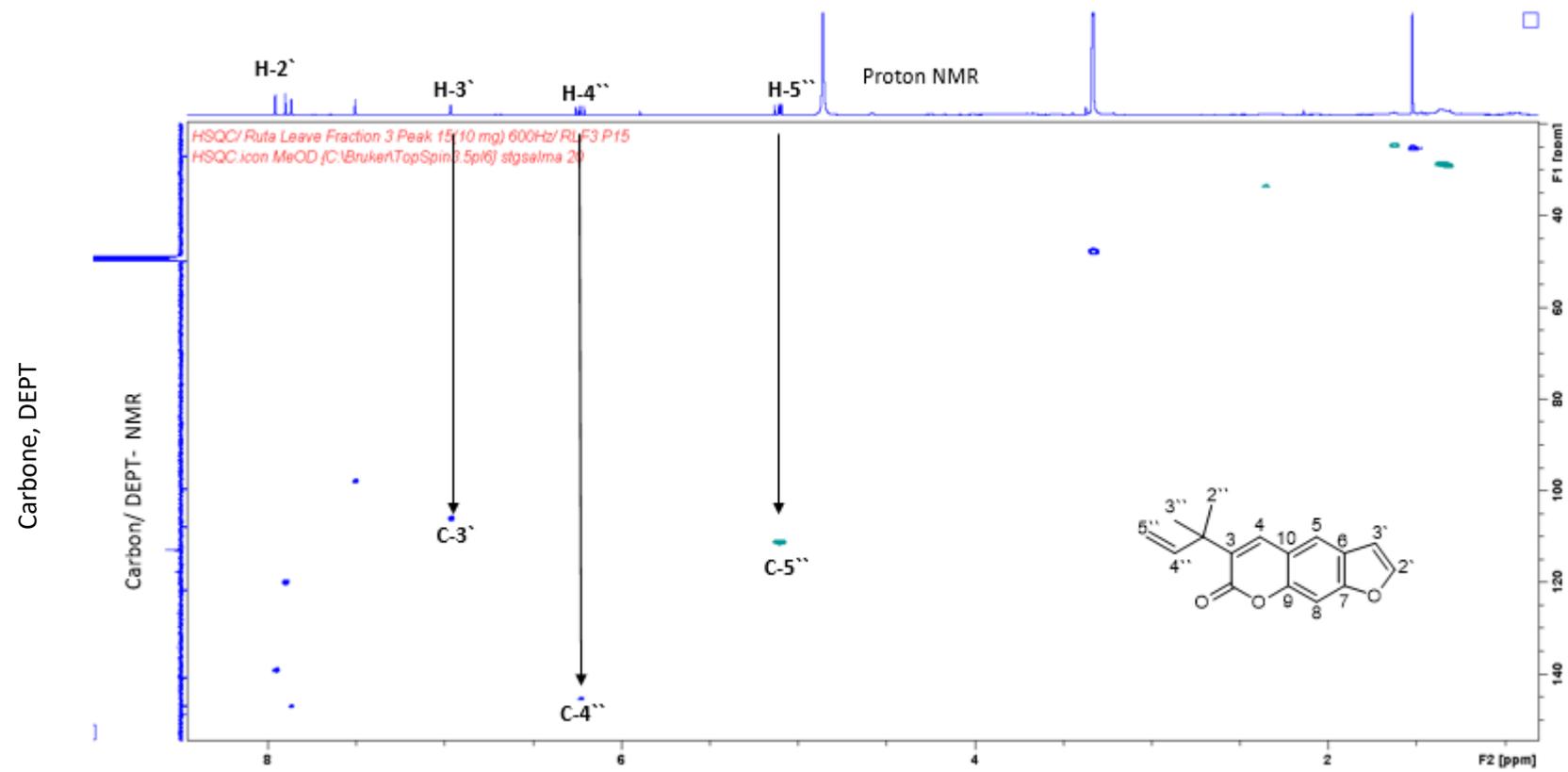


Figure 2.9: HSQC NMR experiment

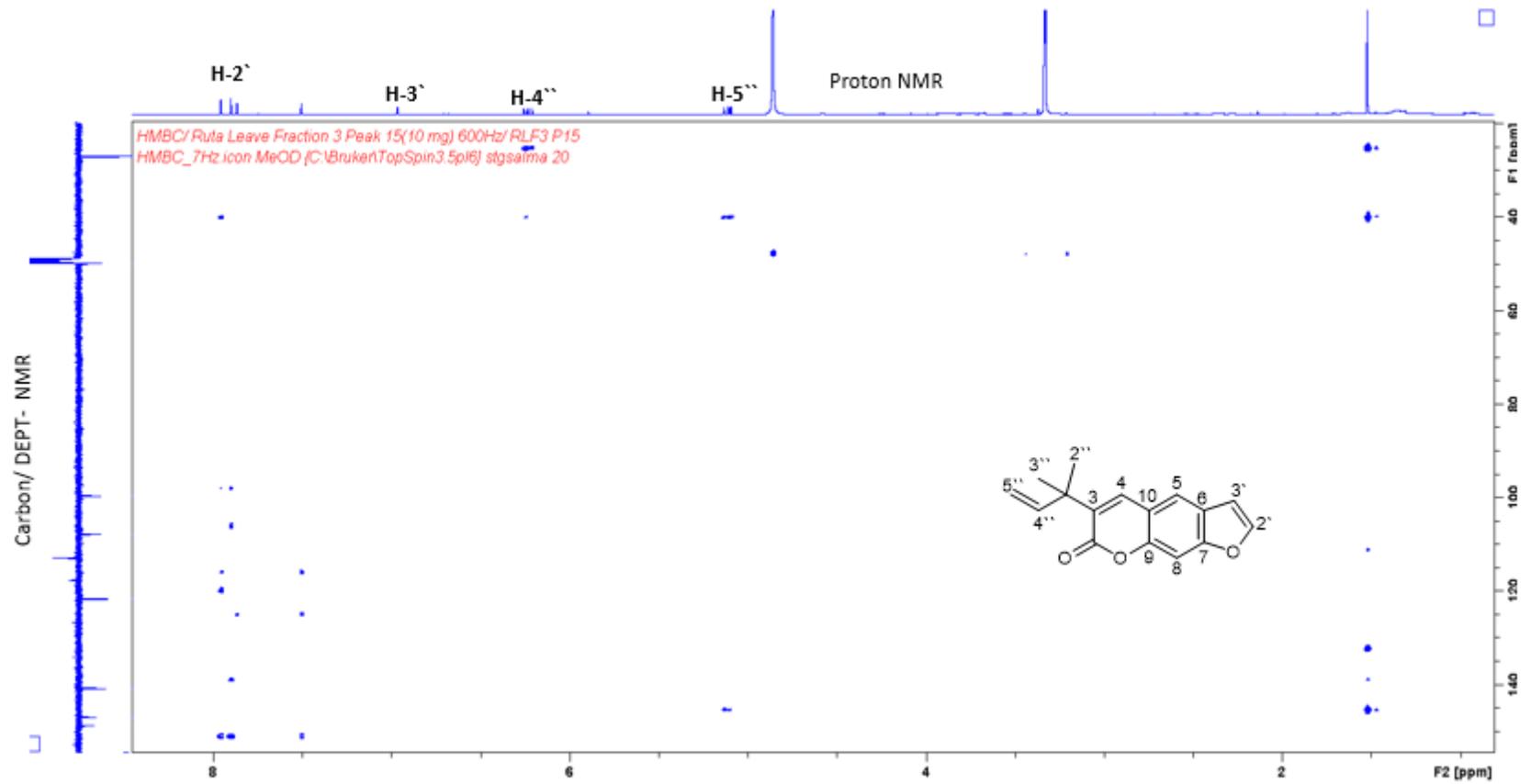


Figure 2.10: HMBC NMR experiment

2.4.2 Mass spectroscopy (MS)

Mass spectroscopy is one of the most powerful useful authoritative analytical techniques to determine the molecular weight of compounds and the molecular formula. The MS instrument contains an ion source, mass analyser and a detector operated under high vacuum conditions. The main principle of MS is generates of ions either from organic or from an organic compounds. Each molecular ion undergoes a fragmentation process and, each of these ions are then separated by mass spectrometry depending on the mass ratio to their charge. Then the quantity of each type of ion is measured (Gross, 2017). In this study MS was carried out using High resolution mass spectra (HR MS) by the National Mass Spectrometry Facility (NMSF) (Swansea, UK) on Xevo G2-S ASAP or LTQ Orbitrap XL1 spectrometers. Low and High resolution MS analyses were also obtained at LJMU; HR-MS using Agilent 6200 Series Accurate-Mass Time-of-Flight (TOF) LC/MS using electro spray ionisation (ESI) in positive ion mode connected to an Agilent auto-sampler injection system. The samples were dissolved in MeOH and the mass spectrum was recorded relating to their mass to charge range at 1700 m/z . The column temperature was 30°C.

2.5 Antimicrobial study

Various *in vitro* laboratory methods are in use to screen the antimicrobial activity of extracts, fractions and pure compound. In the current study, two of the major phenotypic and sensitive methods were used: disk diffusion method and resazurin assay, for determining the susceptibility of a bacterial isolate. Assessment of the nature of the inhibitory effect (bactericidal or bacteriostatic) and molecular method like methicillin-resistant *Staphylococcus aureus* (MRSA) were applied to establish the direct effect of the antimicrobial agent. All details of the strains with their National Collection types of Culture (NCTC) and American Type Cell Culture Collection (ATCC) number, cultured media, materials and other reagents used in this process are described in Table 2.13. The microbial strains were cultured in nutrient media and all the extracts, fractions and pure compounds

were screened for their potential antimicrobial activity against the selected microbes. A group of antibiotics were used as positive controls.

2.5.1 Disk diffusion method

The primary screening of all extracts were performed using the disk diffusion method, which is one of the common clinical microbiology tests. In this procedure, one single colony from the stock was inoculated in to 100 mL of nutrient broth and incubated overnight at 35 ± 2 °C. After the incubation, microbial strains were adjusted to a McFarland Standard of 0.5 using a Spectrophotometer. Using a micropipette, 200 μ L of calibrated nutrient broth was added to a pre-prepared Petri dish with 20 mL agar broth and evenly spread using a swap stick. Disks prepared by filter paper (Whatman, no.1) were cut in circles of 6 mm diameter and sterilized by autoclave. 20 μ L from the extract (10 mg/mL) was added to the disk and left to dry for 30 mint in a flow hood or 3h at room temperature. The inoculated disc was placed on the agar surface of the Petri dish and incubated under suitable conditions. In general, the antimicrobial agent is spread in the agar and inhibits the growth of the microorganisms, with the inhibition zone diameters measured by a ruler.

2.5.2 Resazurin assay

The modified resazurin test described by Sarker *et al.* (2007) was used to determine the MIC utilising a microtitre-plate. The key feature of this assay is the use of a standard concentration of bacterial suspension. This assay was performed under aseptic conditions.

2.5.2.1 Preparation of standard microbial colonies

The classical method to measure turbidity of microbial strains is to adjust the sample to Macfarland standards at 0.5. In fact, this process is unable to give a standardised number of colony-forming unit (CFU) for all strains due to the difference in optical densities of the different microbe species. In the current study, microbial strains were cultured in 20 mL nutrient agar on Petri dishes and incubated for 12-48 h at 35 °C. Single colonies from incubated plates were transferred to sterilized tubes containing 100 mL nutrient broth and incubate tubes at 35 °C for 24 -48 h. After the incubation, the tubes were centrifuged at 4000 rpm for 5 min. The supernatant was discarded and 20 mL sterile normal saline was added to

the tubes and again centrifuged at the same conditions. Centrifugation was repeated until the supernatant become clear. A spectrophotometer was used at 500 nm to determine the optical density of the bacterial suspension with dilution factor and calculations to obtain a concentration of 5×10^6 cfu/mL.

Table 2.13: Strains, antibiotics, reagents and materials used for antimicrobial activity

Microbial strains	Bacterial strain		NCTC/ ATCC number	
	Gram-positive	<i>Micrococcus luteus</i>	NCTC 7508	
		<i>Staphylococcus aureus</i>	NCTC 12981	
	Gram-negative	<i>Escherichia coli</i>	NCTC12241	
		<i>Pseudomonas aeruginosa</i>	NCTC 12903	
	Fungal strain			
	<i>Candida albicans</i>		ATCC 90028	
Antibiotics	Name		Product details	
	Ciprofloxacin		Sigma-Aldrich, Israel	
	Nalidixic acid		Sigma, Germany	
	Chloramphenicol		Acros, USA	
	Gentamicin		Sigma-Aldrich, USA	
	Nystatin		Sigma, Germany	
Reagents	Nutrient broth		Sigma-Aldrich, Spain	
	Agar broth			
	Resazurin sodium salt		Aldrich, USA	
	Normal saline prepared from sodium chloride		Sigma, UK	
Materials	Spectrophotometer		BMG Lab Teach, UK	
	96 well plate		Falcon, USA	

2.5.2.2 Preparation of resazurin solution

Resazurin is a blue dye sodium salt. The resazurin solution was prepared by dissolving 1 mg of resazurin in 5 mL of sterile distilled water. A vortex mixer was used to ensure that it was a well-dissolved and homogenous solution. Resazurin was used in this assay as an indicator of cell growth.

2.5.2.3 Preparation of tested materials

The stock concentration was prepared by dissolving the tested materials in 10% (v/v) DMSO or sterilized water. The stock concentration was 10 mg/mL for crude extracts, while 1 mg/mL for the fractions and pure compounds.

2.5.2.4 Preparation of 96 well plates

All wells on 96 plates were filled with 50 μ L sterilized normal saline. The test material (100 μ L) was added to the first row of the plate and serial dilutions were made using multichannel pipettes by transferring 50 μ L. Resazurin (10 μ L) was added to all wells and finally 10 μ L of bacterial suspension (5×10^6 cfu/mL) was added to each well. To prevent bacterial dehydration, each plate was wrapped loosely with cling film. Every plate contained an antibiotic as a positive control for the bacterial stain, and they were ciprofloxacin, nalidixic acid, gentamycin and chloramphenicol, while for *C. albicans*, nystatin was used.

2.5.2.5 Result interpretation

The normal resazurin colour is blue. During incubation, if the test materials inhibited the microorganisms they acquire the resazurin's blue colour, or become purple or colourless, which is considered as a positive result. Whereas, the development of pink colour represents no effect of the test materials on the microbes. The lowest concentration at which the colour change occurs is considered as an MIC (minimum inhibitory concentration) value. The mean of three values were calculated.

2.6 Determination of bacteriostatic or bactericidal property

To determine whether the compounds are bacteriostatic or bactericidal, 20 mL nutrient agar were poured in empty Petri dish and incubated at 37°C for 24 h. Under aseptic condition, 100 µL of culture from the rows of MIC in the microtitre plates was transferred to a Petri dish and incubated at 37°C for 24 h. Any microbial regrowth indicated bacteriostatic, and no growth indicated bactericidal activity of the test sample.

Chapter 3
Results and Discussion

3.1 *Ruta chalepensis* L.

3.1.1 Extraction

The Soxhlet extraction, sequentially with *n*-hexane, dichloromethane (DCM) and methanol (MeOH), of four different parts of *Ruta chalepensis* afforded three different extracts for each of the plant parts, giving a total of 12 different extracts of varying extraction yields (Table 3.1). The highest percentage yield of extraction of *R. chalepensis* was for the *n*-hexane fruit extract (4.44%), and the lowest percentage was 0.45% for the DCM stem extract.

Table 3.1: Percentage yield of *R. chalepensis*

<i>Ruta chalepensis</i> parts	Powder weight (g)	Extract type	% Yield
Fruit	103.05	<i>n</i> -Hexane (RFH)	4.44
		DCM (RFD)	4.05
		MeOH(RFM)	5.81
Stem	81.1	<i>n</i> -Hexane (RSH)	0.64
		DCM (RSD)	0.45
		MeOH(RSM)	10.0
Leaves	97.77	<i>n</i> -Hexane (RLH)	3.93
		DCM(RLD)	2.18
		MeOH (RLM)	3.87
Root	109.89	<i>n</i> -Hexane (RRH)	2.42
		DCM (RRD)	1.50
		MeOH (RRM)	2.13

3.1.2 Preliminary analytical TLC screening

All the *n*-hexane and DCM extracts of *R. chalepensis* parts were subjected to analytical TLC analysis (Suliman, 2018), following the methods **E** and **G** (Figure 3.1). The developed TLC plates were viewed under short (254 nm) and long (366 nm) UV light followed by spraying with anisaldehyde reagent and then heating at 100°C for 5 min to reveal various coloured spots (Sarker and Nahar, 2012)

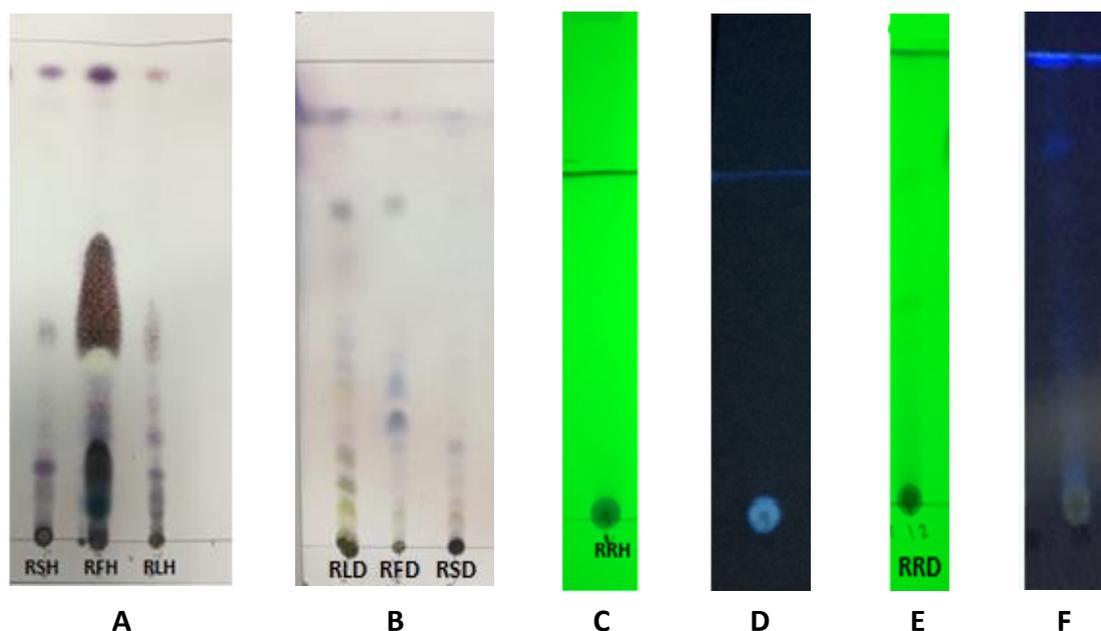


Figure 3.1: The TLC plates for *R. chalepensis* extracts

A: TLC plate for n-hexane extract after spraying; *B:* TLC plate for DCM extract after spraying, *C:* TLC plate for root n-hexane extract under short UV; *D:* TLC plate for root n-hexane extract long UV; *E:* TLC plate for root DCM extract under short UV; *F:* TLC plate for root n-hexane extract under long UV.

3.1.3 Analytical HPLC screening of the MeOH extracts of *R. chalepensis*

The MeOH extracts of the fruits, leaves and stems of *R. chalepensis* (10 mg/mL) were subjected to Dionex Ultimate 3000 analytical HPLC coupled with a photodiode array detector, while the root extract of *R. chalepensis* (10 mg/ml) was analysed using the analytical Agilent HPLC. Method **L** was used to analyse all the MeOH extracts for 45 min with a volume of injection of 20 μ L and a flow rate of 1 mL/min. (Figure 3.2).

3.1.4 The antimicrobial screening of extracts of various parts of *R. chalepensis*

The modified microtitre plate assay as described by Sarker *et al.* (2007) was used on this study using four types of bacterial strains, *Escherichia coli* (NCTC 12241), *Pseudomonas aeruginosa* (NCTC 12903), *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981) and one fungal strain *Candida albicans* (ATCC 90028). A group of antibiotics were used in this assay as positive controls and their MIC values were summarized in Table 3.2, while the negative control was a mixture of nutrient

broth and normal saline. The *n*-hexane, DCM and MeOH extracts of the leaves, stems, fruits and roots of *R. chalepensis* were assessed for antimicrobial proprieties against all these microbial strains. This result revealed that all extracts of the leaves, stems and fruits could inhibit all the microbial strains used in this study with different MIC values (Table 3.3). The *R. chalepensis* root extracts did not show any inhibitory activity, while the MeOH extract of the leaves (RLM) displayed considerable antimicrobial activity against both *C. albicans* and *M. luteus* with an MIC value of 1.95×10^{-2} mg/mL. Both the DCM extract of the leaves (RLD) and the *n*-hexane extract (RLH) revealed inhibitory activity against *M. luteus* with an MIC value of 1.56×10^{-1} mg/mL.

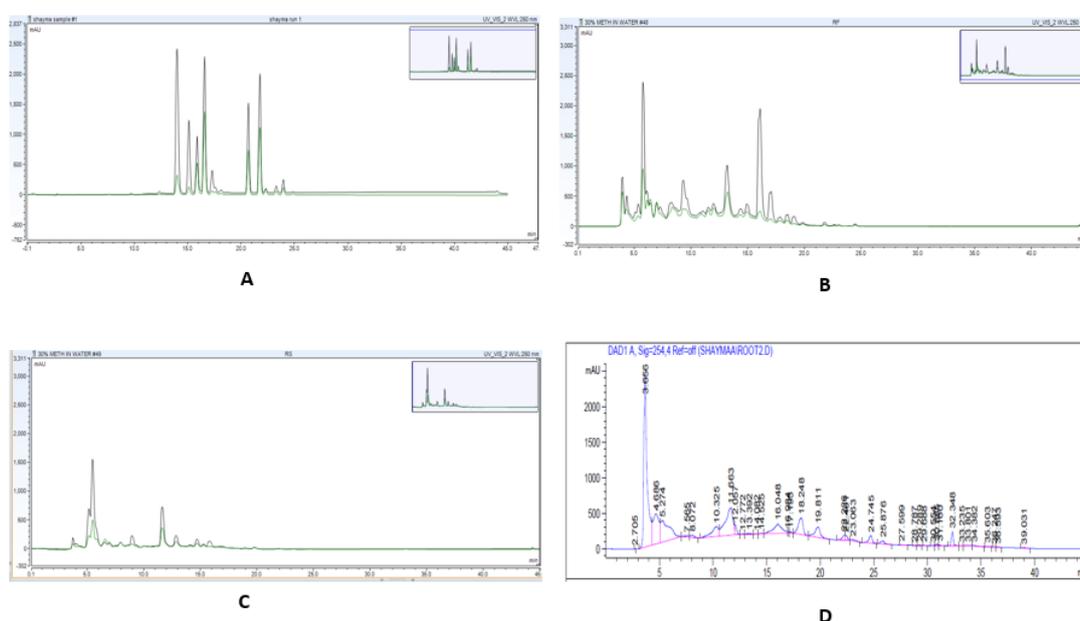


Figure 3.2: Analytical HPLC results for *R. chalepensis*

A: RLM; B: RFM; C: RSM; D: RRM

Table 3.2: The MIC values for antibiotics (positive controls)

Microbes	Antibiotics	MIC mg/mL	Ciprofloxacin mg/mL
<i>Escherichia coli</i>	Nalidixic acid	1.55×10^{-2}	1.55×10^{-2}
<i>Pseudomonas aeruginosa</i>	Gentamycin	1.22×10^{-4}	1.95×10^{-3}
<i>Micrococcus luteus</i>	Chloramphenicol	9.76×10^{-4}	3.90×10^{-3}
<i>Staphylococcus aureus</i>	Chloramphenicol	9.76×10^{-4}	3.90×10^{-3}
<i>Candida albicans</i>	Nystatin	9.76×10^{-4}	N/A

The most potent antimicrobial activity was observed with the MeOH extract of the stem (RSM) with an MIC value of 7.81×10^{-2} mg/mL against *M. luteus* and *C. albicans*, while the DCM extract of the stem (RSD) exhibited a moderate level of inhibition against all the tested microbes. The MIC value of the *n*-hexane stem extract (RSH) against all microbes as 6.25×10^{-1} mg/mL except against *C. albicans* with an MIC value of 1.95×10^{-1} mg/mL. Both the *n*-hexane (RRH) and DCM (RRD) extracts of the roots (RRH) were unable to show any inhibitory efficacy at test concentrations, while the MIC value of the MeOH extract (RRM) was 1.25 mg/mL against *M. luteus*. This result indicated that most of the extracts had certain levels of antimicrobial activity. Many previous studies on *R. chalepensis* documented its antimicrobial activity using different plant parts and methods (Alzoreky and Nakahara, 2003; Priya *et al.*, 2009; Babu-Kasimala *et al.*, 2014; Suliman, 2018). However, there is no report on the antimicrobial activity studies of *R. chalepensis* using the modified microtitre resazurine assay. This current piece of work generated the first phytochemical report on the analysis of the fruits of Iraqi *R. chalepensis* species along with their antimicrobial activity using modified resazurine microtitre assay.

After the antimicrobial screening of the crude extracts of *R. chalepensis*, the MIC 6.25×10^{-1} mg/mL was chosen as the minimum threshold of activity for any extract for further analysis leading to the isolation of compounds responsible for their antimicrobial activity.

3.1.5 Chromatographic fractionation of the extracts

3.1.5.1 Vacuum liquid chromatography fractionation (VLC)

The active *n*-hexane and DCM extracts were fractionated by VLC over silica gel using the method **A** and the method **C**, respectively. Table 3.4 summarizes the yield of various fractions. The VLC fractions of the *n*-hexane extracts were analysed by TLC using the method **E** (Figure 3.3), while the VLC fractions of the DCM extracts were analysed by the TLC method **F** (Figure 3.4).

Table 3.3: The minimum inhibitory concentration (MIC) values for crude extracts of *R. chalepensis*

Bacteria and fungi		Extract	Plant parts (mg/mL)			
			Leaves	Stems	Fruits	Roots
Gram-negative	<i>E. coli</i>	<i>n</i> -Hexane	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	N/A
		DCM	3.12 x 10 ⁻¹	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	N/A
		Methanol	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	6.25 x 10 ⁻¹	5
	<i>P. aeruginosa</i>	<i>n</i> -Hexane	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	N/A
		DCM	3.12 x 10 ⁻¹	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	N/A
		Methanol	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	1.56 x 10 ⁻¹	2.5
Gram-positive	<i>M. luteus</i>	<i>n</i> -Hexane	1.56 x 10 ⁻¹	6.25 x 10 ⁻¹	3.12 x 10 ⁻¹	N/A
		DCM	1.56 x 10 ⁻¹	6.25 x 10 ⁻¹	7.81 x 10 ⁻²	N/A
		Methanol	1.95 x 10 ⁻²	7.81 x 10 ⁻²	3.90 x 10 ⁻²	1.25
	<i>S. aureus</i>	<i>n</i> -Hexane	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	N/A
		DCM	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	6.25 x 10 ⁻¹	N/A
		Methanol	3.12 x 10 ⁻¹	3.12 x 10 ⁻¹	3.12 x 10 ⁻¹	5
Pathogenic yeast	<i>C. albicans</i>	<i>n</i> -Hexane	6.25 x 10 ⁻¹	1.95 x 10 ⁻¹	1.56 x 10 ⁻²	N/A
		DCM	3.12 x 10 ⁻¹	3.12 x 10 ⁻¹	3.90 x 10 ⁻²	N/A
		Methanol	1.95 x 10 ⁻²	7.81 x 10 ⁻²	3.90 x 10 ⁻²	2.5

*N/A: No Activity

Various chromatographic fractionation techniques were performed for the active crude extracts using vacuum liquid chromatography fractionation (VLC) and solid phase extraction (SPE) as described in the experimental section.

3.1.5.2 Solid-phase extractions (SPE)

The active MeOH extracts were fractionated by SPE using the method **D** to collect four fractions from every extract. Table 3.4 summarizes the yield of fractions. All fractions were then subjected to analytical HPLC.

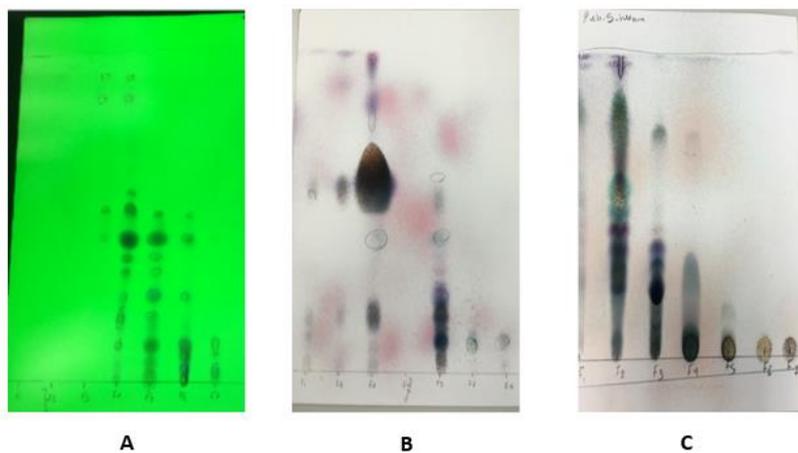


Figure 3.3: TLC analysis of n-hexane extracts of *R. chalepensis*

A: Fruit; B: Leaves; C: Stems

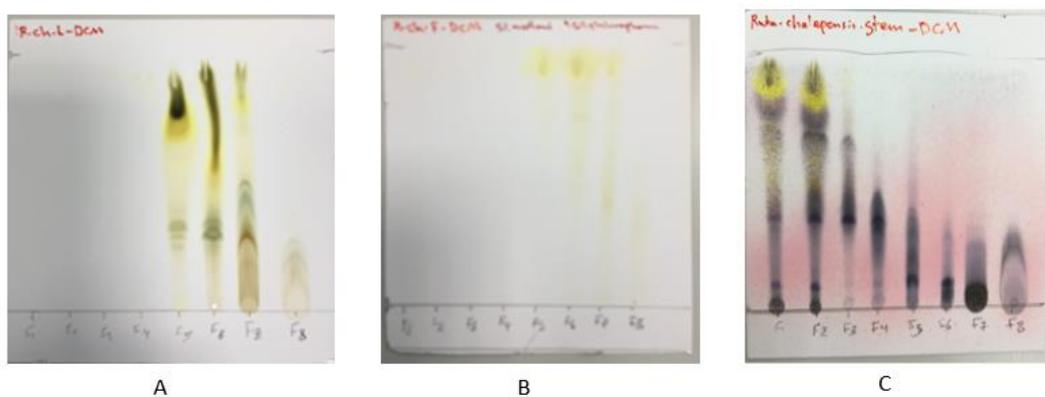


Figure 3.4: TLC analysis of the DCM extracts of *R. chalepensis*

A: Leaves; B: Fruits; C: Stem

Table 3.4: The yields of fractions of the active extracts of various parts of *R. chalepensis*

Plant parts	Type of extract	Weight of extract (g)	Fractions yield %							
			F1	F2	F3	F4	F5	F6	F7	F8
Fruits	<i>n</i> -Hexane	3.080	0.46	0.60	46.70	23.93	14.44	16.33	3.61	-
	DCM	3.1244	0.40	0.30	3.00	4.26	30.88	39.77	6.55	8.80
	Methanol	1.7894	42.64	20.68	4.87	2.85	-	-	-	-
Leaves	<i>n</i> -Hexane	3.002	0.66	2.25	5.38	25.00	13.08	10.94	6.66	-
	DCM	2.7291	0.40	0.50	0.52	1.01	5.43	28.79	3.89	8.50
	Methanol	1.776	28.80	18.45	2.26	4.15	-	-	-	-
Stems	<i>n</i> -Hexane	2.9634	3.86	6.26	3.96	8.97	6.01	2.16	0.62	-
	DCM	2.1195	12.87	2.44	1.41	1.43	1.52	1.01	2.08	2.25
	Methanol	1.8021	31.93	9.12	3.98	1.83	-	-	-	-

3.1.6 Screening of *R. chalepensis* fractions for antimicrobial activity

All VLC and SPE fractions were screened for their antimicrobial activity against *Escherichia coli* (NCTC 12241), *Pseudomonas aeruginosa* (NCTC 12903), *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981) and *Candida albicans* (ATCC 90028).

The VLC fraction 4 from the *n*-hexane extract of the fruits showed significant inhibitory activity against all the microbial strains used in this study, specifically against *M. luteus* with an MIC value of 7.81×10^{-3} mg/mL. Moreover, the VLC fraction 7 from the DCM extract of the fruits presented strong inhibitory activity to inhibit all the microbes used in this test with different MIC values. Regarding the MeOH extract of the fruits, the SPE fractions 2 and 3 displayed remarkable reduction of growth of the tested microbial strains with different MIC values (Table 3.5). Table 3.5 outlines the MIC values of the fractions of various extracts of the fruits of *R. chalepensis*.

Most of the fractions of the *n*-hexane extract of the leaves did not show significant antimicrobial activity, while VLC fraction 7 (F7) of the DCM extract of the leaves displayed good inhibition of growth of all five microbial strains, particularly against *M. luteus* with an MIC value of 3.12×10^{-2} mg/mL. Furthermore, both SPE fractions 2 and 3 from MeOH extract of the leaves exhibited significant antimicrobial activity (Table 3.6). Table 3.6 summarizes the MIC values of the fractions of the extracts of the leaves of *R. chalepensis* leaves. As for the stems, the VLC fractions originating from both *n*-hexane and DCM extracts had weak antimicrobial activity, but the SPE fractions 2 and 3 of the MeOH extract displayed notable inhibitory activity against all microbial strains used in this study, particularly, the SPE fraction 2 inhibited *C. albicans* with an MIC value of 3.12×10^{-2} mg/mL. Table 3.7 outlines the MIC values of various fractions of extracts of stems of *R. chalepensis*.

Table 3.5: MIC values (in mg/mL) of *Ruta chalepensis* fruit fractions

Microbes	Type of extract	F1	F2	F3	F4	F5	F6	F7	F8
<i>Escherichia coli</i>	<i>n</i> -Hexane	N/A	N/A	N/A	6.25×10^{-2}	N/A	N/A	N/A	-
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	3.12×10^{-2}	N/A
	MeOH	N/A	7.81×10^{-3}	1.56×10^{-2}	N/A	-	-	-	-
<i>Pseudomonas aeruginosa</i>	<i>n</i> -Hexane	N/A	N/A	N/A	1.25×10^{-1}	N/A	N/A	N/A	
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	6.25×10^{-2}	N/A
	MeOH	N/A	6.25×10^{-2}	2.5×10^{-1}	N/A	-	-	-	-
<i>Micrococcus luteus</i>	<i>n</i> -Hexane	N/A	1	1	6.25×10^{-2}	5×10^{-1}	5×10^{-1}	1	-
	DCM	N/A	N/A	1	1	1	1	3.12×10^{-2}	1
	MeOH	N/A	6.25×10^{-2}	3.12×10^{-2}	N/A	-	-	-	-
<i>Staphylococcus aureus</i>	<i>n</i> -Hexane	N/A	N/A	N/A	7.81×10^{-3}	N/A	N/A	N/A	
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	N/A
	MeOH	N/A	5×10^{-1}	1.25×10^{-1}	N/A	-	-	-	-
<i>Candida albicans</i>	<i>n</i> -Hexane	N/A	N/A	N/A	3.12×10^{-2}	5×10^{-1}	5×10^{-1}	0.1	
	DCM	0.5	0.25	N/A	N/A	6.25×10^{-2}	3.12×10^{-2}	0.5	N/A
	MeOH	N/A	3.12×10^{-2}	1.56×10^{-2}	N/A	-	-	-	-

*N/A: No Activity

Table 3.6: MIC values (in mg/mL) of *Ruta chalepensis* leaves fractions

Microbes	Type of extract	F1	F2	F3	F4	F5	F6	F7	F8
<i>Escherichia coli</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	-
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	MeOH	N/A	5×10^{-1}	2.5×10^{-1}	N/A	-	-	-	-
<i>Pseudomonas aeruginosa</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	
	DCM	N/A	N/A	N/A	N/A	N/A	N/A		N/A
	MeOH	N/A	1.25×10^{-1}	2.5×10^{-1}	N/A	-	-	-	-
<i>Micrococcus luteus</i>	<i>n</i> -Hexane	N/A	1	N/A	N/A	1	5×10^{-1}	5×10^{-1}	
	DCM	5×10^{-1}	5×10^{-1}	1	N/A	N/A	N/A	N/A	N/A
	MeOH	N/A	5×10^{-1}	5×10^{-1}	N/A	-	-	-	-
<i>Staphylococcus aureus</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	
	DCM	5×10^{-1}	5×10^{-1}	N/A	N/A	N/A	N/A		N/A
	MeOH	N/A	5×10^{-1}	2.5×10^{-1}	N/A	-	-	-	-
<i>Candida albicans</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	1	5×10^{-1}	
	DCM	5×10^{-1}	5×10^{-1}	N/A	N/A	N/A	N/A	N/A	5×10^{-1}
	MeOH	N/A	3.12×10^{-2}	1.25×10^{-1}	N/A	-	-	-	-

*N/A: No Activity

Table 3.7: MIC values (in mg/mL) of *Ruta chalepensis* stems fractions

Microbes	Type of extract	F1	F2	F3	F4	F5	F6	F7	F8
<i>Escherichia coli</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	-
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	N/A
	MeOH	N/A	5×10^{-1}	2.5×10^{-1}	N/A	-	-	-	-
<i>Pseudomonas aeruginosa</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	-
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	N/A
	MeOH	N/A	1.25×10^{-1}	5×10^{-1}	N/A	-	-	-	-
<i>Micrococcus luteus</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	1	5×10^{-1}	
	DCM	N/A	N/A	1	1	5×10^{-1}	5×10^{-1}	3.12×10^{-2}	5×10^{-1}
	MeOH	N/A	1.56×10^{-2}	6.25×10^{-2}	N/A	-	-	-	-
<i>Staphylococcus aureus</i>	<i>n</i> -Hexane	N/A	N/A	N/A	N/A	N/A	N/A	N/A	
	DCM	N/A	N/A	N/A	N/A	N/A	N/A	2.5×10^{-1}	5×10^{-1}
	MeOH	N/A	1.25×10^{-1}	2.5×10^{-1}	N/A	-	-	-	-
<i>Candida albicans</i>	<i>n</i> -Hexane	N/A	N/A	N/A	5×10^{-1}	N/A	1	N/A	-
	DCM	N/A	N/A	1	1	1	1.56×10^{-2}	1.25×10^{-1}	N/A
	MeOH	N/A	1.25×10^{-1}	3.12×10^{-2}	1	-	-	-	-

*N/A: No Activity

3.1.7 Phytochemistry of *Ruta chalepensis*

Chromatographic separation of the active fractions of the *n*-hexane, DCM and MeOH extracts of different parts of *R. chalepensis* afforded nineteen compounds (Figure 3.5): bergapten (43), kokusaginine (99), isokokusaginine (100), skimmianine (41), rutin (48), rutin 3'-methyl ether (101), rutin 7,4'-dimethyl ether (102), 6-hydroxy-rutin-3'-7-dimethyl ether (103), chalepin (45), chalepentin (46), rutamarin (47), isopimpinellin (44), γ -fagarine (42), imperatorin (104), graveoline (105), ribalinium (106), arborinine (107), 3',6-disinapoylsucrose (108) and hexadecane (109) (Figure 3.5). All the isolated compounds were identified by spectroscopic means (NMR and HRESIMS) and the spectral data of known compounds were compared with the respective published data. Of the isolated compounds, except 6-hydroxy-rutin-3'-7-dimethyl ether (103), all isolated compounds are known natural products. To the best of our knowledge, this is the first report on the phytochemical studies on the fruits of the Iraqi species of *R. chalepensis*. Compounds 100-102 and 108 are reported here for the first time from *R. chalepensis*.

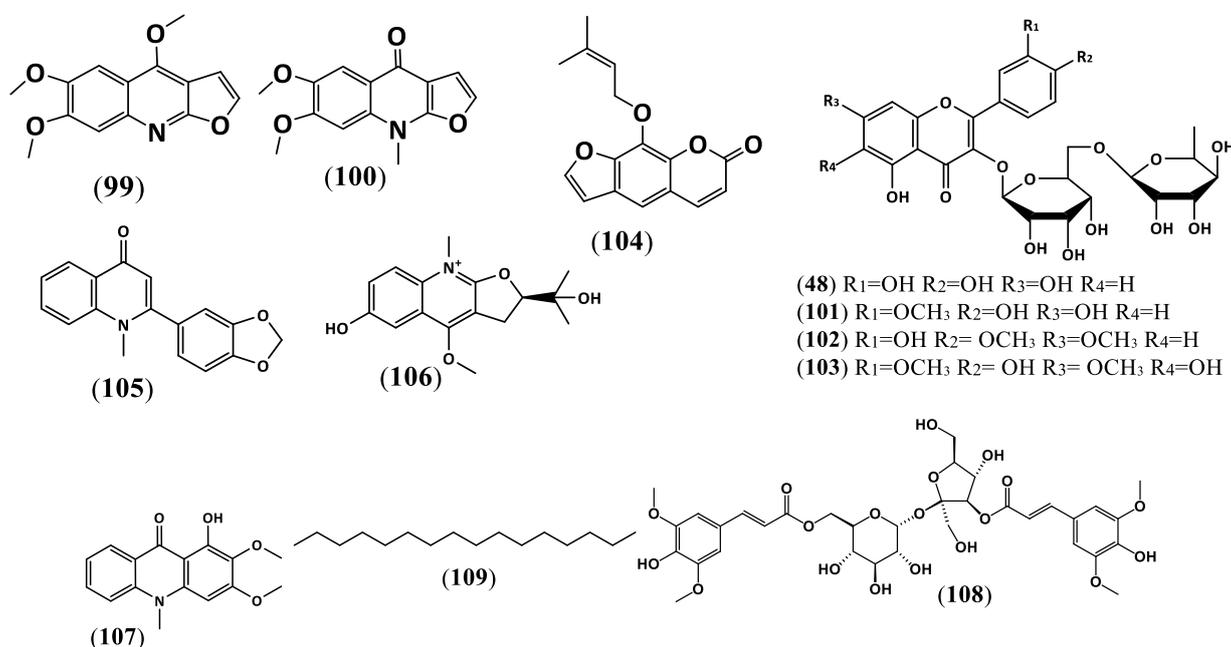


Figure 3.5: Isolated compounds from *R. chalepensis*

Kokusaginine (99), isokokusaginine (100), rutin 3'-methyl ether (101), rutin 7,4'-dimethyl ether (102), 6-hydroxy-rutin-3'-7-dimethyl ether (103), imperatorin (104), graveoline (105), ribalinium (106), arborinine (107), 3',6-disinapoylsucrose (108) and hexadecane (109)

3.1.8 Isolation compounds from active fractions of *Ruta chalepensis*

All the active fractions were subjected to different chromatographic methods, *e.g.*, PTLC and/or prep-HPLC to isolate the active compounds.

3.1.8.1 *R. chalepensis* fruit *n*-hexane fraction 4

The VLC fraction 4 obtained from the *n*-hexane extract of fruits was subjected to analytical TLC using the method **E** (Figure 3.6) because of its significant antimicrobial activity against all the five strains used in this study.

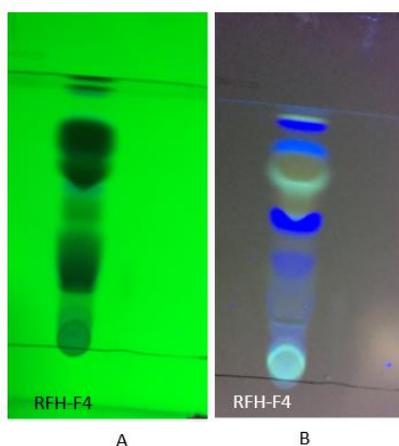


Figure 3.6: Analytical TLC (using method **E**) of the VLC fraction 4 of the *n*-hexane extract of *R. chalepensis* fruits

The method **B** was selected for further fractionation based on the analysis of the TLC chromatography that had been developed using method **E**. The application of the method **B** gave 11 fractions from fraction 4 (Figure 3.7) and then these fractions were subjected to tests for their antimicrobial activities. Out of these 11 fractions, only the fractions RFH4-3, RFH4-4, RFH4-5 and RFH4-11 was showed a significant antimicrobial activity (Table 3.8). To determine the responsible active compound, method **I** was used to develop PTLC, in RFH4-3, RFH4-4 and RFH4-5, while RFH4-11 subjected to method **K**. (Figure 3.8). The final compounds (**42**), (**43**), (**46**), (**47**) and (**99**) were obtained from these chromatographic techniques suggesting the possible antimicrobial activity of these fractions.

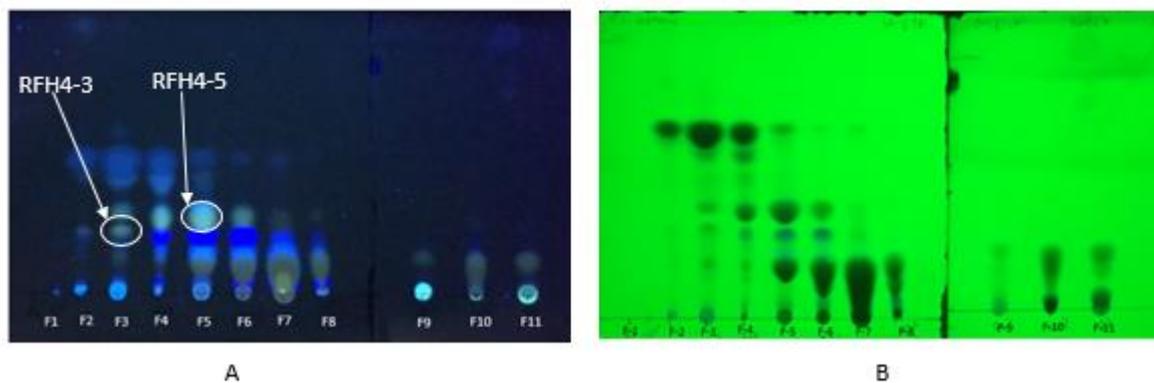


Figure 3.7: Analytical TLC (using method I) of the VLC fractions of *R. chalepensis* fruit fraction 4

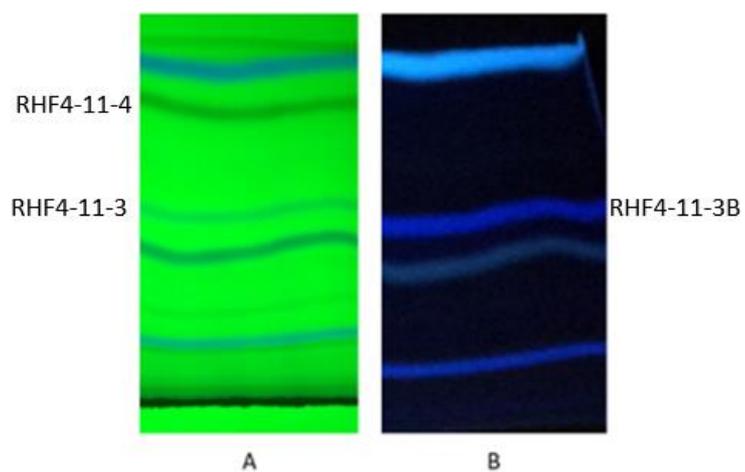


Figure 3.8: Preparative TLC (using method K) of *R. chalepensis* fruit fraction 11 derived from fraction 4

Table 3.8: The MIC values of the VLC fractions of *R. chalepensis* fruit fraction 4, and compounds isolated from the active fractions

Fractions	Microbes (MIC in mg/mL)					Methods		Extracts name	Isolated compounds	
	<i>E. coli</i>	<i>P. aeruginosa</i>	<i>M. luteus</i>	<i>S. aureus</i>	<i>C.albicans</i>	Type	Name		Name	Number
F1	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F2	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F3	N/A	5×10^{-1}	1.25×10^{-1}	5×10^{-1}	2.5×10^{-1}	PTLC	I	RFH4-3	Chalepensin	46
F4	N/A	N/A	5×10^{-1}	N/A	2.5×10^{-1}	-	I	RFH4-4	Chalepensin	46
F5	N/A	N/A	5×10^{-1}	N/A	5×10^{-1}	-	I	RFH4-5	Rutamarin	47
F6	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F7	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F8	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F9	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F10	N/A	N/A	N/A	N/A	N/A	-	-	-	-	-
F11	5×10^{-1}	2.5×10^{-1}	1.56×10^{-2}	1.25×10^{-1}	3.12×10^{-2}	PTLC	K	RFH4-11-4	Bergapten	43
								RFH4-11-3	γ -Fagarine	42
								RFH4-11-3B	Kokusaginine	99

*N/A: No Activity

3.1.8.2 *R. chalepensis* fruit DCM fraction 7

The antimicrobial screening of *R. chalepensis* fruit DCM fractions revealed that the most active fraction was fraction 7, which was subjected to the prep-HPLC (Agilent) using method **M** for 35 min. The injection volume was 300 μ L and the flowrate was 10 mL/min (Figure 3.9). This separation method was gave six compounds, arborinine (**107**), (0.3 mg) (Knölker, 2017; Kumar, 2018), graveoline (**105**, 0.2 mg) (Ulubelen *et al.*, 1986), kokusaginine (**99**, 0.3 mg) (Adamska-Szewczyk *et al.*, 2016), γ -fagarine (**42**, 0.5 mg) (Ulubelen *et al.*, 1986), skimmianine (**41**, 1.2 mg) (Ulubelen *et al.*, 1986) and chalepentin (**46**, 0.4 mg) (Adamska-Szewczyk *et al.*, 2016).

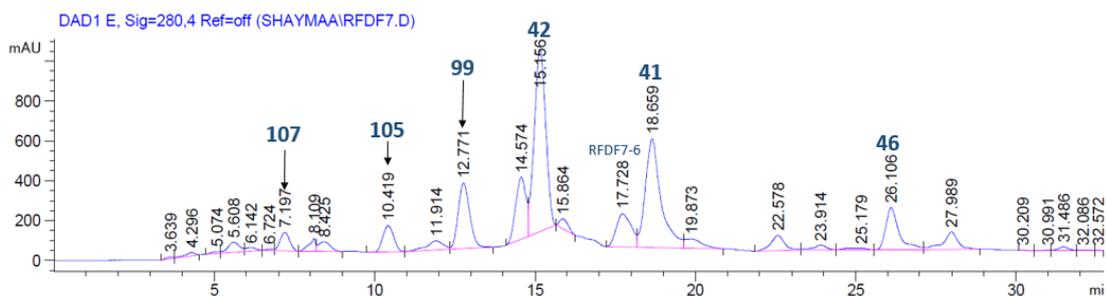


Figure 3.9: Preparative- HPLC chromatogram of isolated compounds from *R. chalepensis* fruit DCM extract fraction 7, using method **M**

3.1.8.3 *R. chalepensis* fruit MeOH fraction 2

The method **L** in prep-HPLC was used to isolate the active compounds from fraction 2 of the fruit MeOH extract (Figure 3.10). The injection volume was 200 μ L and the flow rate was 10 mL/min. The separation process obtained gave mixed peak (RFF2-11) and seven compounds, ribalinium (**106**, 0.5 mg), rutin (**48**, 1.2 mg) (Hamad, 2012), rutin 3'-methyl ether (**101**, 1.0 mg), 6-hydroxy-rutin-3'-7-dimethyl ether (**103**, 0.3 mg), graveoline (**105**, 1.2 mg) (Ulubelen *et al.*, 1986), isopimpinellin (**44**, 1.0 mg) (Richardson *et al.*, 2016) and γ -fagarine (**42**, 0.4 mg) (Adamska-Szewczyk *et al.*, 2016).

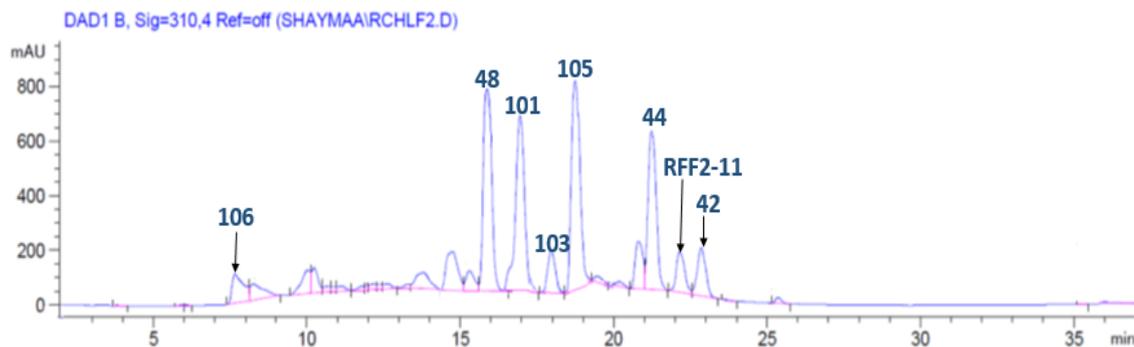


Figure 3.10: Preparative-HPLC chromatogram of isolated compounds from *R. chalepensis* fruit MeOH extract fractions 2 using method **L**

Purification of RFF2-11

The peak RFF2-11 was isolated as a mixture of compounds. This mixture was subjected to PTLC using the method **J** to isolate two pure compounds **43** (3 mg; $R_f = 0.19$) and **99** (2 mg; $R_f = 0.47$), which were analysed by MS and NMR spectroscopic techniques.

3.1.8.4 *R. chalepensis* fruit MeOH fraction 3

This active fraction was subjected to prep-HPLC using method **L** for 35 min. The injection volume was 300 μ L and the flowrate was 10 mL/min (Figure 3.11). The separation process was obtained six compounds rutin (**48**, 0.6 mg) (Hamad, 2012), skimmianine (**41**, 0.5 mg) (Ulubelen *et al.*, 1986), imperatorin (**104**, 1.8 mg), chalepentin (**46**, 0.2 mg), chalepin (**45**, 1.2 mg) (Richardson *et al.*, 2016) and rutamarin (**47**, 1.5 mg) (Adamska-Szewczyk *et al.*, 2016).

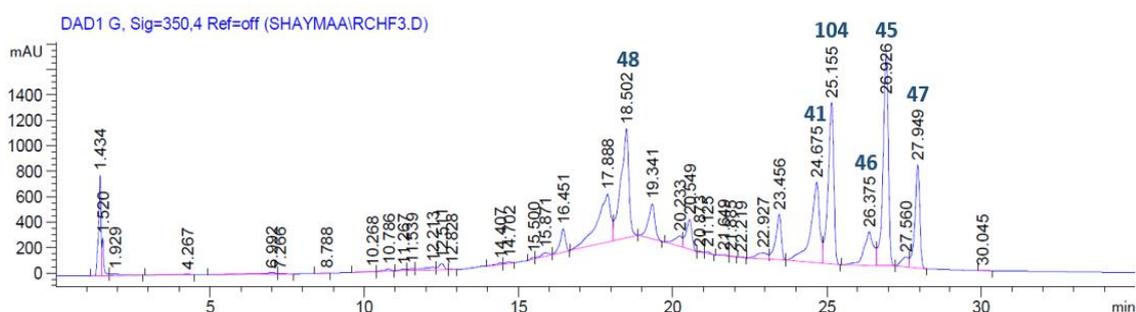


Figure 3.11: Preparative- HPLC chromatogram of isolated compounds from *R. chalepensis* fruit MeOH extract fraction 3, using method **L**

3.1.8.5 *R. chalepensis* leaves DCM fraction 7

The method **O** in prep-HPLC was used to separate the target compounds from fraction 7 of leaves DCM extract (Figure 3.12). The injection value was 350 μ L and the flowrate was 10 mL/min. The isolation techniques were obtained four pure compounds, graveoline (**105**, 1.0 mg) (Ulubelen *et al.*, 1986), kokusaginine (**99**, 0.6 mg) (Adamska-Szewczyk *et al.*, 2016), arborinine (**107**, 0.4 mg) (Knölker, 2017; Kumar, 2018) and chalepin (**45**, 0.9 mg) (Richardson *et al.*, 2016). The injection value was 350 μ L and the flowrate was 10 mL/min.

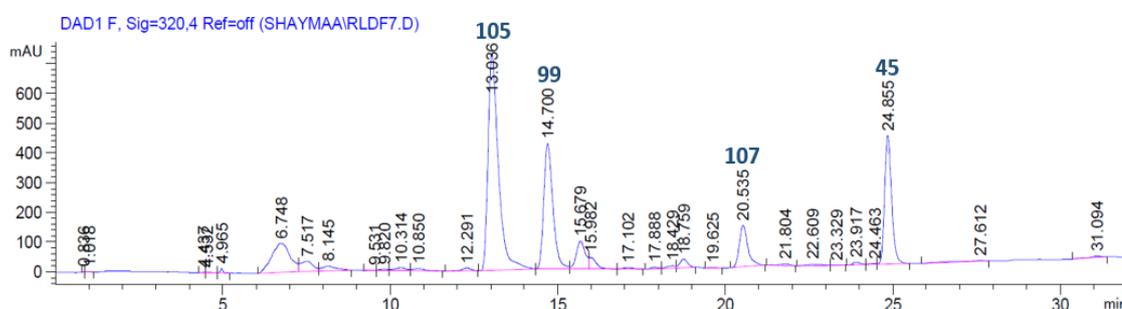


Figure 3.12: Preparative- HPLC chromatogram of isolated compounds from *R. chalepensis* leaves DCM extract fraction 7, using method **O**

3.1.8.6 *R. chalepensis* leaves MeOH fraction 2

This *R. chalepensis* leaves fraction 2 of MeOH extract was analysed by prep-HPLC using method **L**. The injection value was 250 μ L and the flowrate was 10 mL/min for 33min. The separation process were obtained two fractions RLF2-5, RLF2-7 and four compounds bergapten (**43**, 1.3 mg) (Gonzalez *et al.*, 1977), rutin (**48**, 0.3 mg) (Hamad, 2012), γ -fagarine (**42**, 0.3 mg) (Adamska-Szewczyk *et al.*, 2016) and kokusaginine (**99**, 1.0 mg) (Adamska-Szewczyk *et al.*, 2016).

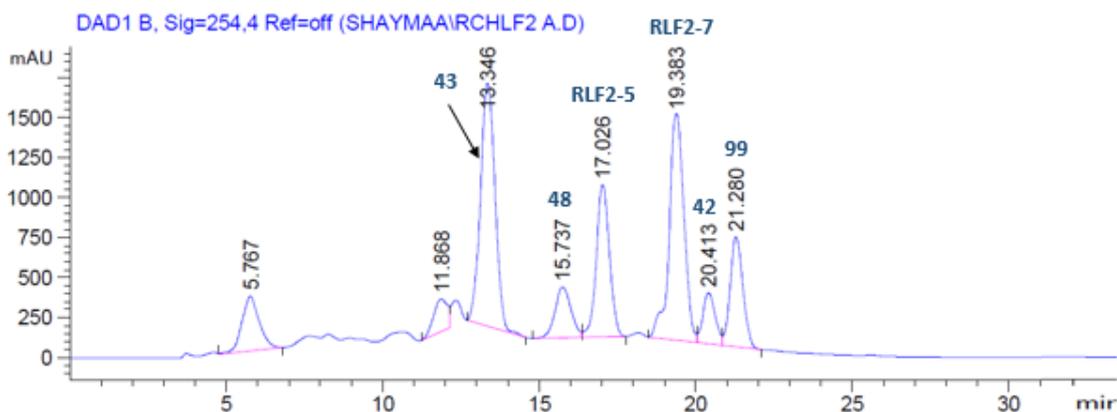


Figure 3.13: Preparative- HPLC chromatogram of isolated compounds from *R.chalepensis* leaves MeOH extract fraction 2, using method **L**

Purification of RLF2-5

The peak RLF2-5 was isolated as a mixture of compounds. This mixture was subjected to semi-prep-HPLC using method **U** to isolate two pure compounds rutin 3'-methyl ether (**101**, 0.5 mg) and rutin 7,4'- dimethyl ether (**102**, 0.3 mg). The injection volume was 100 μ L and the flow rate was 2 mL/min for 20 min (Figure 3.14).

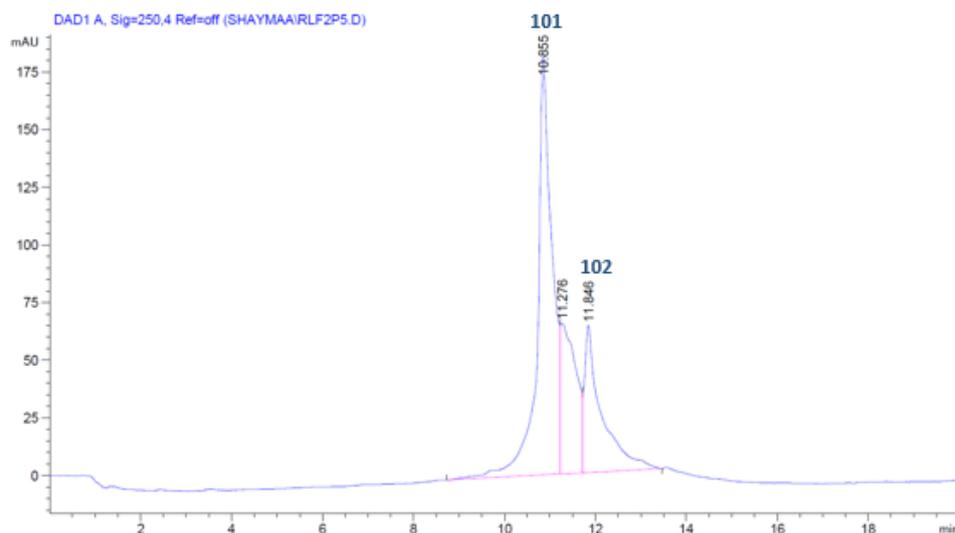


Figure 3.14: Semi-preparative-HPLC chromatogram of purification compounds from RLF2-5, using method **U**

Purification of RLF2-7

This peak RLF2-7 was separated as a mixture of compounds. This mixture was purified by semi-prep-HPLC using method **P** for 21 min. The injection value was 100 μ L and the flowrate was 2 mL/min (Figure 3.15). The separation process gave three compounds rutin (**48**, 0.2 mg) (Hamad, 2012), isokokusaginine (**100**, 1.5 mg) (Openshaw, 1967) and graveoline (**105**, 1.2 mg) (Ulubelen *et al.*, 1986).

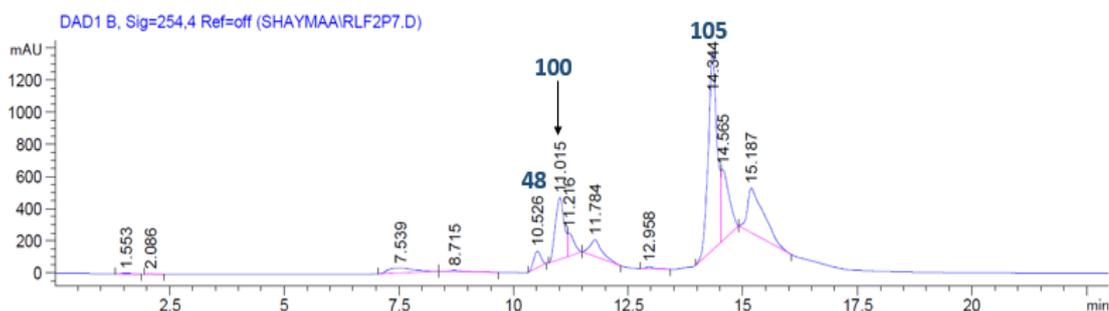


Figure 3.15: Semi-preparative- HPLC chromatogram of purification compounds from RLF2-7, using method **P**

3.1.8.7 *R. chalepensis* leaves MeOH fraction 3

The *R. chalepensis* leaves fraction 3 of the MeOH extract was subjected to prep-HPLC using method **L** for 37min. (Figure 3.16). The injection volume was 250 μ L, while the flow rate was 10 mL/min. The separation process produced eight compounds, rutin (**48**, 0.2 mg), graveoline (**105**, 1.3 mg) (Ulubelen *et al.*, 1986), bergapten (**43**, 0.2 mg) (Gonzalez *et al.*, 1977), kokusaginine (**99**, 1.5 mg) (Adamska-Szewczyk *et al.*, 2016), γ -fagarine (**42**, 0.5 mg) (Adamska-Szewczyk *et al.*, 2016), arborinine (**107**, 0.3 mg) (Knölker, 2017; Kumar, 2018), chalepin (**45**, 1.2 mg) (Richardson *et al.*, 2016) and chalepentin (**46**, 0.2 mg) (Adamska-Szewczyk *et al.*, 2016).

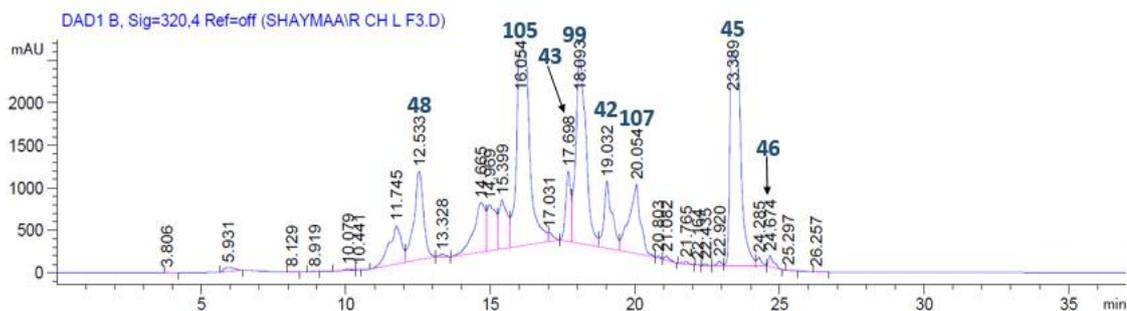


Figure 3.16: Preparative-HPLC chromatogram of isolated compounds of *R. chalepensis* leaves MeOH extract fraction 3, using method **L**

3.1.8.8 *R. chalepensis* stem MeOH fraction 2

The method **L** in semi-prep-HPLC was used to isolate the active compounds from fraction 2 of stem MeOH extract (Figure 3.17). The injection value was 100 μ L and the flow rate was 2 mL/min. The separation process gave three compounds 3',6-disinapoylsucrose (**108**, 1.0 mg), rutin (**48**, 0.8 mg) (Hamad, 2012) and rutin 7,4'- dimethyl ether (**102**, 0.2 mg).

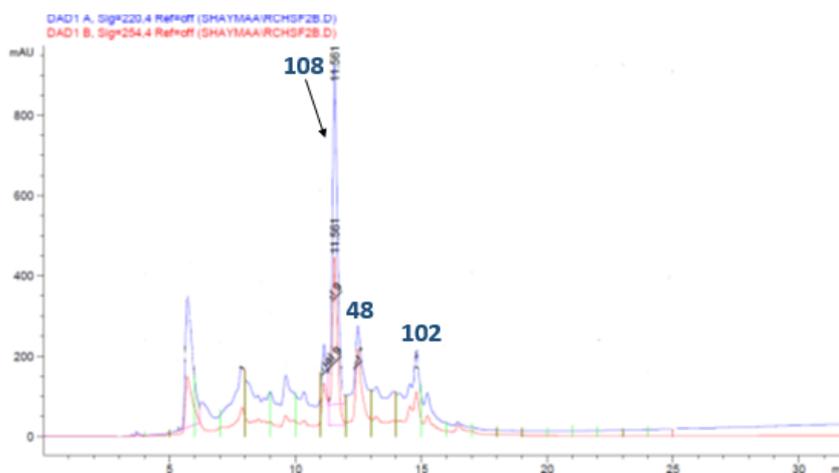


Figure 3. 17: Semi -preparative- HPLC Chromatogram of isolated compounds of *R. chalepensis* stem MeOH extract fraction 2, using method **L**

3.1.8.9 *R. chalepensis* stem MeOH fraction 3

The *R. chalepensis* stem fraction 3 of the MeOH extract was subjected to semi-prep-HPLC using method **L** for 40 min (Figure 3.18). The injection value was 200 μ L and flow rate was 2 mL/min. The separation process afforded six compounds, kokusaginine (**99**, 0.9 mg) (Adamska-Szewczyk *et al.*, 2016), γ -fagarine (**42**, 0.4 mg) (Adamska-Szewczyk *et al.*,

2016), bergapten (**43**, 1.6 mg) (Gonzalez *et al.*, 1977), arborinine (**107**, 0.7 mg) (Knölker, 2017; Kumar, 2018), chalepin (**45**, 0.2 mg) and chalepentin (**46**, 0.4 mg) (Adamska-Szewczyk *et al.*, 2016).

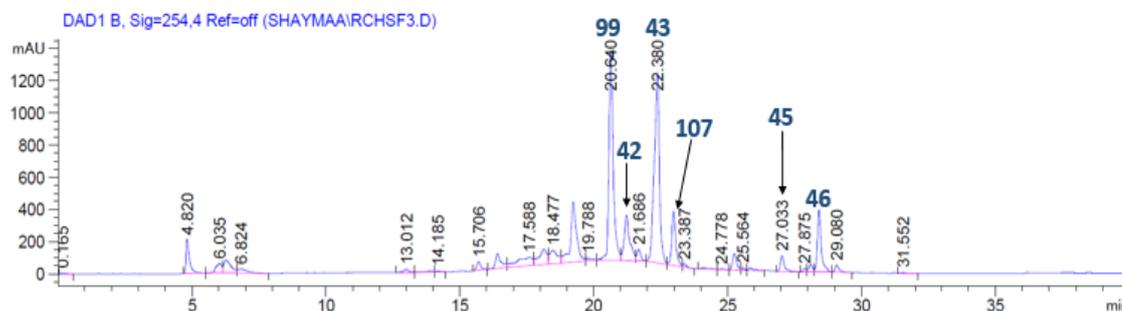


Figure 3.18: Preparative-HPLC chromatogram of isolated compounds from *R. chalepensis* stem MeOH extract fraction 3, using method L

3.1.9 Characterisation and structure elucidation of isolated compounds

3.1.9.1 Structure elucidation of kokusaginine (99)

The compound **99** was isolated as a colourless powder. The HRESIMS (Figure 3.19) suggested the empirical formula as $C_{14}H_{14}NO_4$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 260.0917 (calculated 260.0922). The 1H NMR spectrum (Figure 3.20, Table 3.9) exhibited signals for three methoxy groups at δ_H 4.44 (3H, s, 4-OMe), 4.03 (3H, s, 7-OMe), 4.02 (3H, s, 6-OMe) and two olefinic protons at δ_H 7.57 (1H, d, $J = 2.80$ Hz, H-2) and 7.04 (1H, d, $J = 2.80$ Hz, H-3'). Moreover, two aromatic protons resonated at δ_H 7.48 (1H, s, H-5) and 7.34 (1H, s, H-8). The ^{13}C -NMR (Figure 3.21, Table 3.9) presented seven quaternary carbons at δ_C 163.4 (C-2), 102.5 (C-3), 155.9 (C-4), 148.1 (C-6), 152.9 (C-7), 142.7 (C-9), 113.2 (C-10); three methyl groups at δ_C 59.1 (4-OMe), 56.3 (7-OMe), 56.3 (6-OMe) and two olefinic carbons at δ_C 142.74 (C-2') and 104.9 (C-3'). In addition, there were two aromatic methine carbon signals at δ_C 100.5 (C-5) and 107.0 (C-8). The 1H - 1H COSY spectrum showed the connection between H-2'/H-3'. The HMBC (Figure 3.22) displayed the correlation from H-3' to C-2, 3, 2' and H-2' showed 2J correlation to C-2 and C-3. Moreover, the long-rang correlation was observed from H-8 to C-6, C-7, C-9, C-10 and from H-5 to C-

4, C-6, C-7, C-9, C-10. The HMBC confirmed the positions of methoxy groups by revealing the correlation of OMe/C-4, OMe/C-6 and OMe/C-7. Thus, compound **99** was identified as 4,6,7-trimethoxyfuro[2,3-b]quinolone or kokusaginine. The spectroscopic data of compound **99** were in a good agreement with respective published data of kokusaginine (Wu *et al.*, 2003; Elaine Monteiro *et al.*, 2010). Kokusaginine (**99**) was previously isolated from other species of the Rutaceae including *Ruta* species. Kokusaginine (**99**) was reported to possess antioxidant, anti-leishmanic and anticholinesterase activities (Ulubelen *et al.*, 1986; El Sayed *et al.*, 2000; Wu *et al.*, 2003; Adamska-Szewczyk *et al.*, 2016).

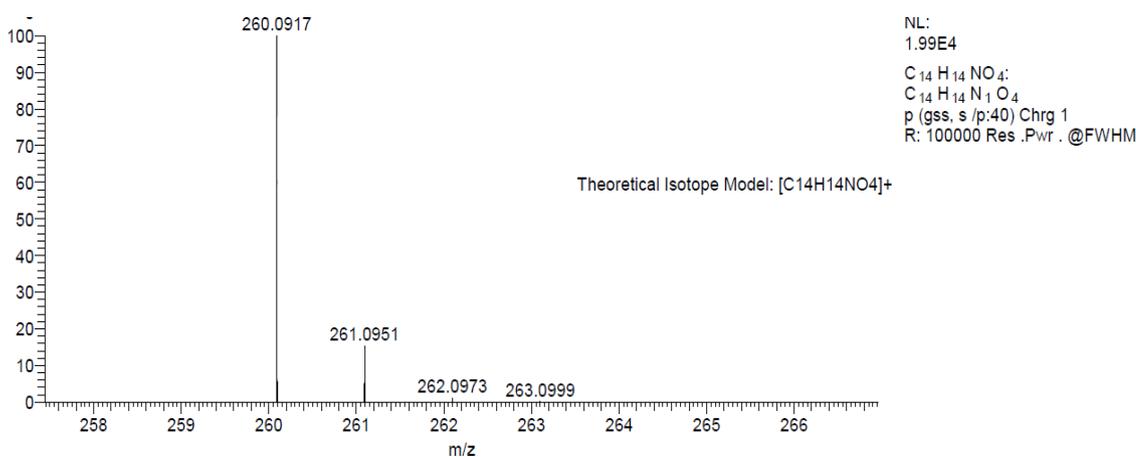


Figure 3.19: The HRESIMS spectrum of compound **99**

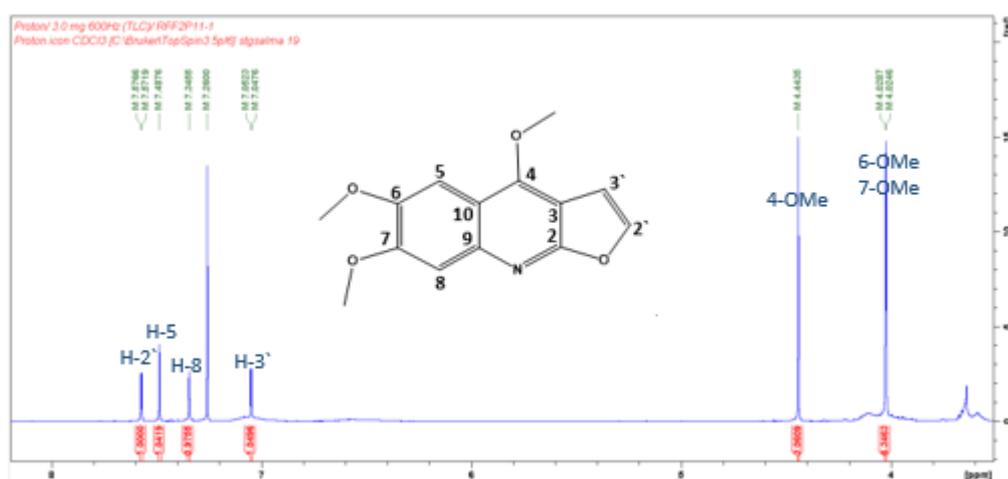


Figure 3.20: The ^1H NMR (600 MHz, CDCl_3) spectrum of compound **99**

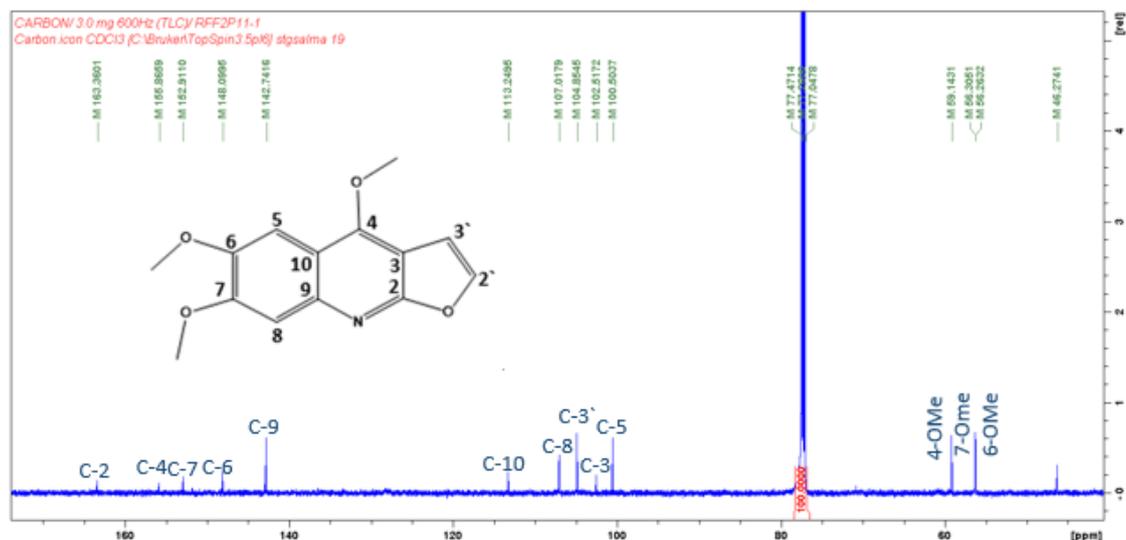


Figure 3.21: The ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **99**

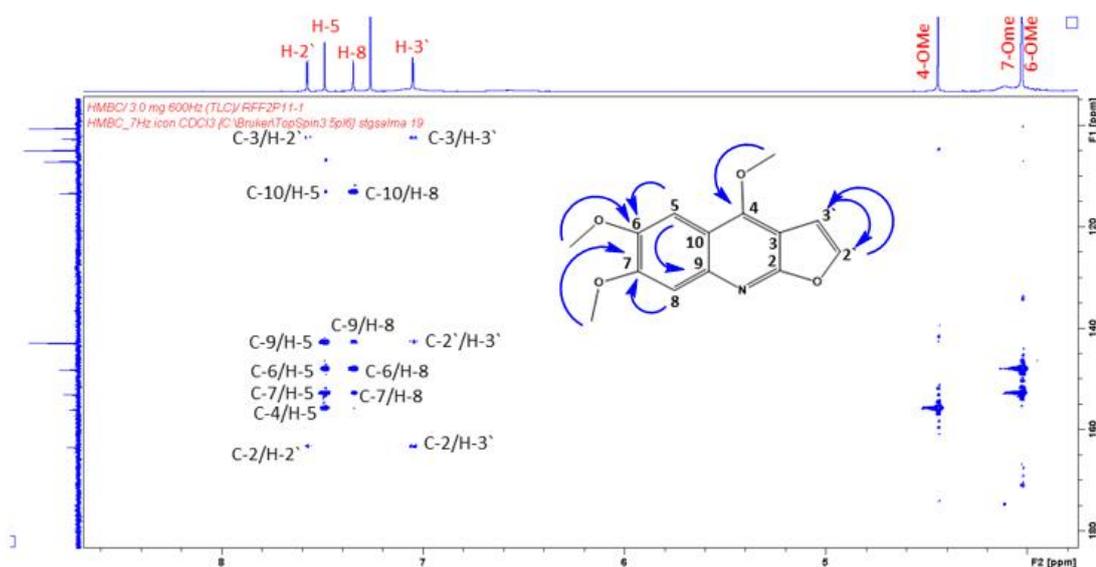


Figure 3.22: The HMBC correlations of compound **99**

3.1.9.2 Structure elucidation of isokokusagine (100)

The compound **100** was isolated as pale yellow powder. The HRESIMS (Figure 3.23) suggested the empirical formula as $\text{C}_{14}\text{H}_{13}\text{NO}_4$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 260.0919 (calculated 260.0922). The ^1H NMR spectrum (Figure 3.24, Table 3.9) displayed two sets of methyl signals at δ_{H} 3.98 (3H) and 4.07 (2X 3H). The deshielded methyl signals at δ_{H} 4.07 must be assigned as OMe groups whereas other methyl could be accounted for NMe. The ^1H NMR spectrum also two aromatic protons resonated at

δ_H 7.87, 7.21 and also olefinic protons at δ_H 7.60 and 7.09. The DEPTQ spectrum (Figure 3.25, Table 3.9) exhibited presence of a total of fourteen carbons which could be assigned based on the correlation observed in the HSQC and HMBC spectra. In the HSQC experiment, the three protons signals at δ_H 3.98 showed direct correlation to the carbon at δ_C 32.6 whilst other methyl signals at δ_H 4.07 revealed direct connection to the carbons at δ_C 56.7 and 57.0. The HMBC experiment (Figure 3.26) played a key role in the confirmation of the structure of the compound. In the HMBC experiment, the methoxyl group protons at δ_H 4.07 (2 x OMe) showed 3J connectivity to the oxygenated quaternary carbons at δ_C 135.6 (C-6) and 154.5 (C-7) while the quaternary carbons at *N*-methyl protons at δ_H 3.98 was connected to two quaternary carbons at δ_C 156.1 (C-2) and 146.2 (C-9) by 3J . The protons at δ_H 7.09 (H-3') and 7.87 (H-5) revealed a common 3J HMBC correlation to the carbonyl at δ_C 172.8 (C-4). Thus, compound **100** was identified as 6,7-dimethoxy-9-methylfuro[2,3-*b*]quinolin-4(9H)-one or isokokusaginine. The spectroscopic data of compound **100** were in a good agreement with respective published data of isokokusaginine. Isokokusaginine (**100**) was previously isolated from *Ruta graveolines* and other plants (Openshaw, 1967). Isokokusaginine (**100**) is here reported for the first time isolated from *R. chalepensis*.

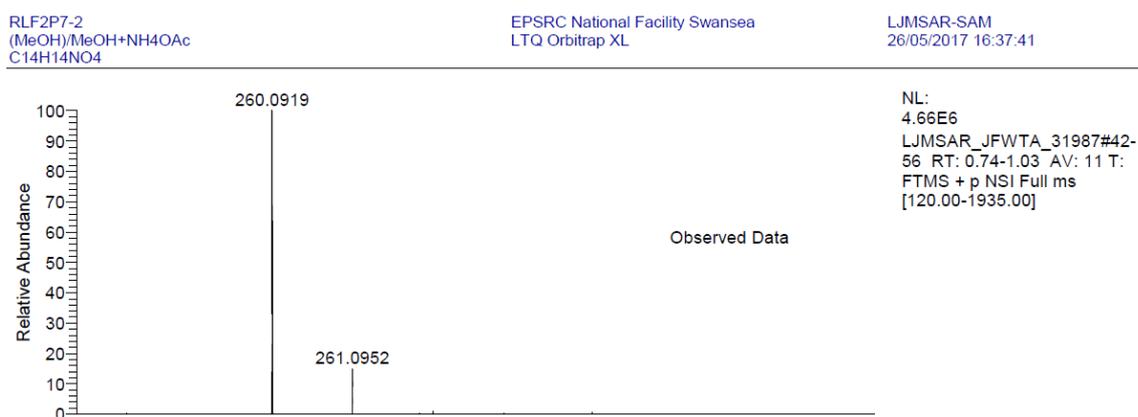


Figure 3.23: The HRESIMS spectrum of compound **100**

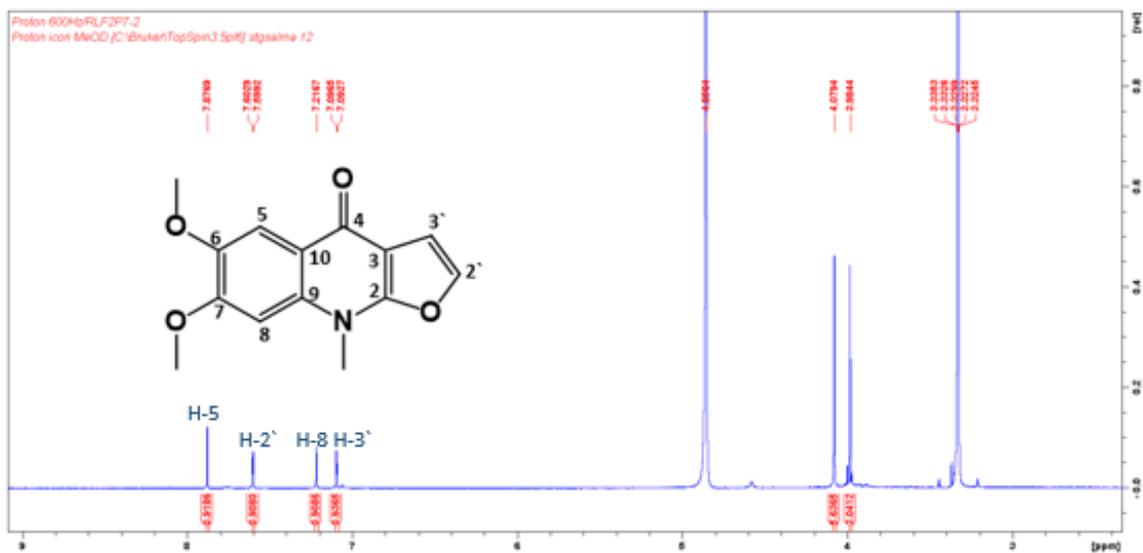


Figure 3.24: The ^1H NMR (600 MHz, CD_3OD) spectrum of compound **100**

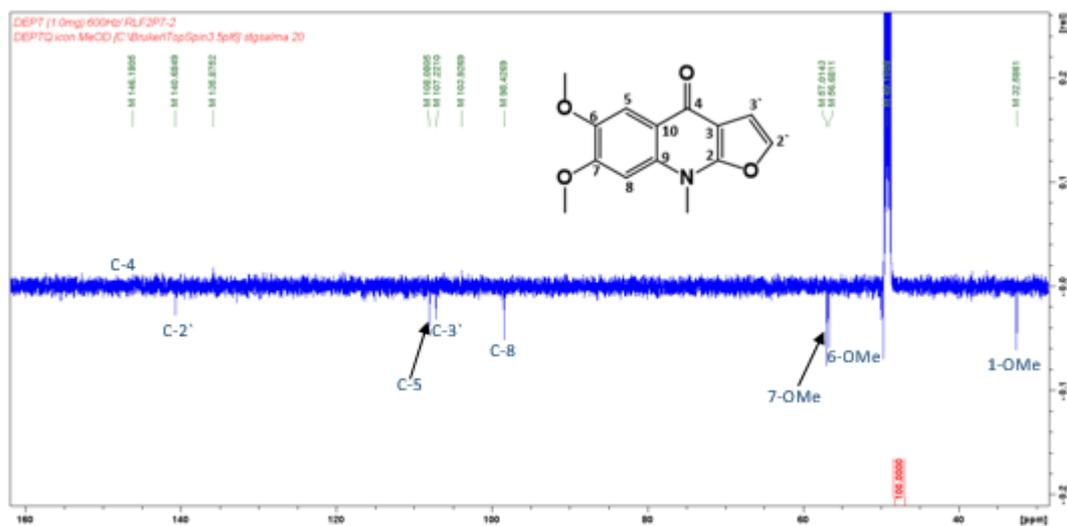


Figure 3.25: The DEPT NMR (150 MHz, CD_3OD) spectrum of compound **100**

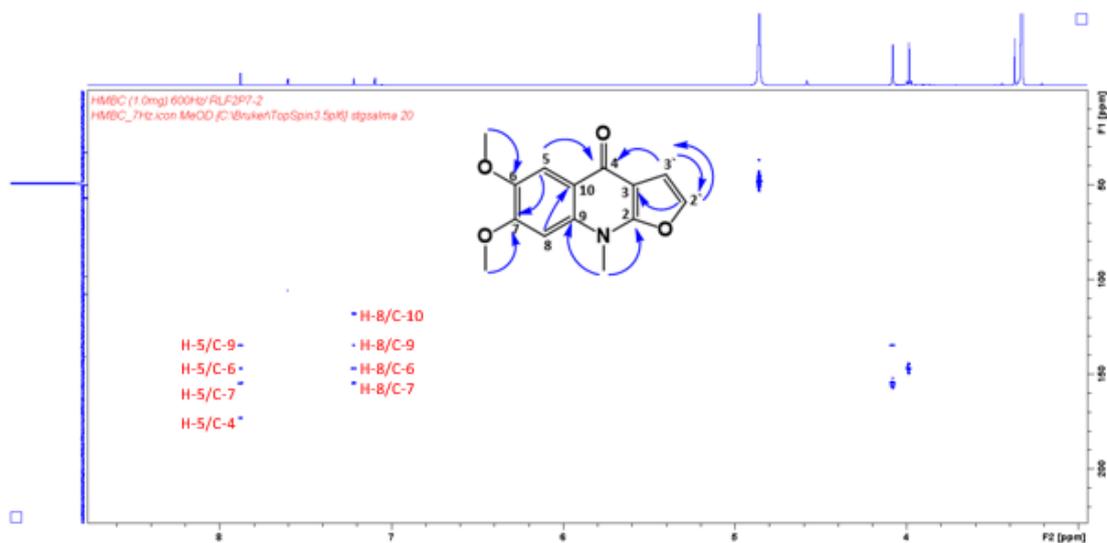


Figure 3.26: The HMBC correlations of compound **100**

3.1.9.3 Structure elucidation of γ -fagarine (**42**)

The compound **42** was isolated as a pale brown crystal from different part extracts of *R. chalepensis*. The HRESIMS (Figure 3.27) suggested the empirical formula as $C_{13}H_{11}NO_3$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 230.0814 (calculated 230.0817). The 1H NMR spectrum (Figure 3.28, Table 3.9) showed two sets of methoxy signals at δ_H 4.45 (3H) and 4.08 (3H), four aromatic protons resonated at δ_H 7.85, 7.36, 7.36 and 7.06 and two olefinic protons at δ_H 7.65 and 7.08 as a part of furan ring. The DEPTQ spectrum (Figure 3.29, Table 3.9) exhibited a total of thirteen carbons, two methoxyl, six quaternary carbons and five olefinic carbons. The protons at δ_H 4.08 and 4.45 revealed direct correlation to the corresponding carbons at δ_C 56.4 and 59.3 in the HSQC spectrum. In the 1H - 1H COSY spectrum (Figure 3.30), proton at δ_H 7.36 (H-6) showed interaction with protons at δ_H 7.85 (H-5) and 7.06 (H-7). 1H - 1H COSY interaction was also revealed between protons at δ_H 7.65 (H-2') and 7.08 (H-3'). In the HMBC experiment (Figure 3.31), H-5, H-3' and methoxyl protons at δ_H 4.45 (MeO-4) displayed 3J connectivity to the oxygenated quaternary carbon at δ_C 157.4 (C-4) whilst H-6 and methoxyl protons at δ_H 4.05 were collated to another oxygen bearing quaternary carbon at δ_C 157.4 (C-4). Thus, compound **42** was

identified as 8-methoxydictamnine or γ -fagarine. The spectroscopic data of compound **42** were in a good agreement with respective published data of γ -fagarine (Boyd *et al.*, 2013). γ -fagarine (**42**) was previously isolated from *Rutaceae* family *Ruta* genus and *Ruta chalepensis*. γ -fagarine (**42**) was reported as antimicrobial, antiplasmodial and anti-HCV effect on 20 $\mu\text{g/ml}$ concentration (Tavares *et al.*, 2014; Adamska-Szewczyk *et al.*, 2016).

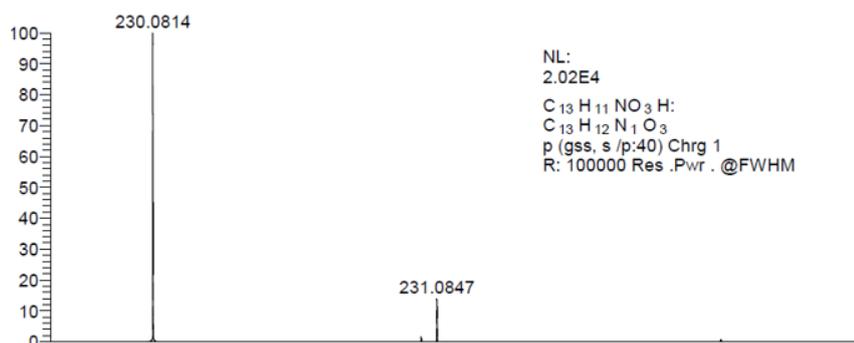


Figure 3.27: The HRESIMS spectrum of compound **42**

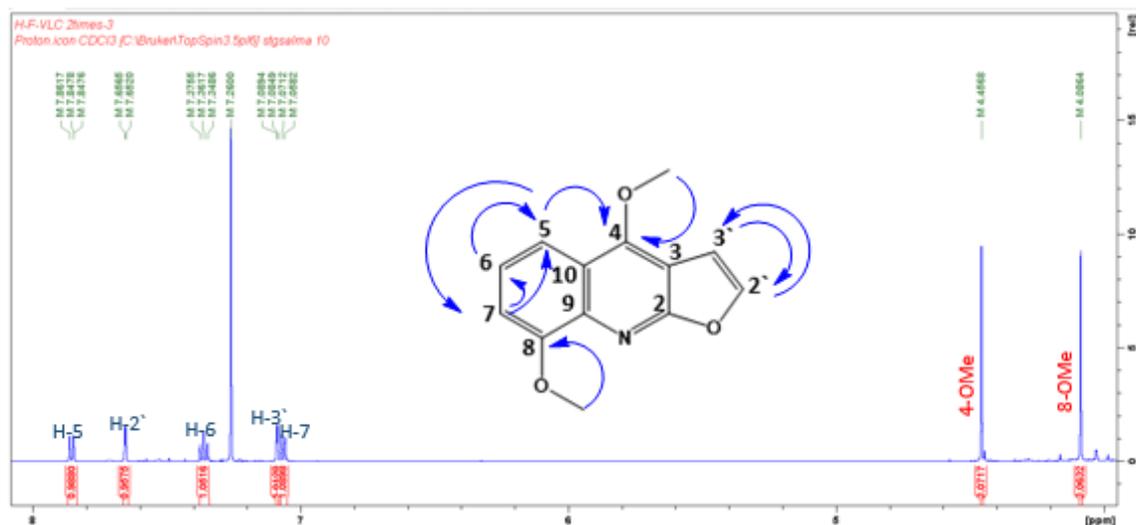


Figure 3.28: ^1H NMR (600 MHz, CDCl_3) spectrum of compound **42**

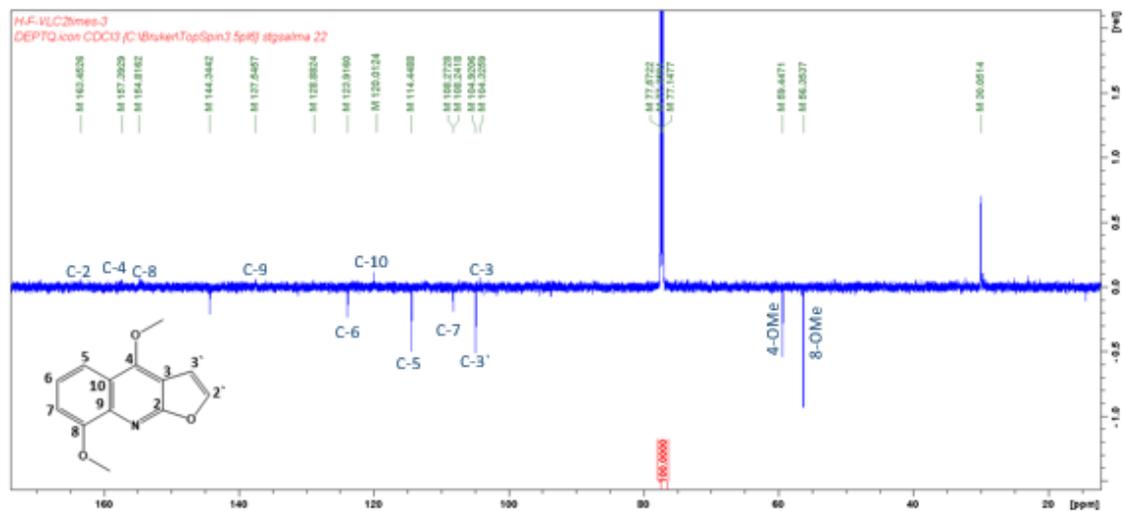


Figure 3.29: DEPT NMR (150 MHz, CDCl_3) spectrum of compound 42

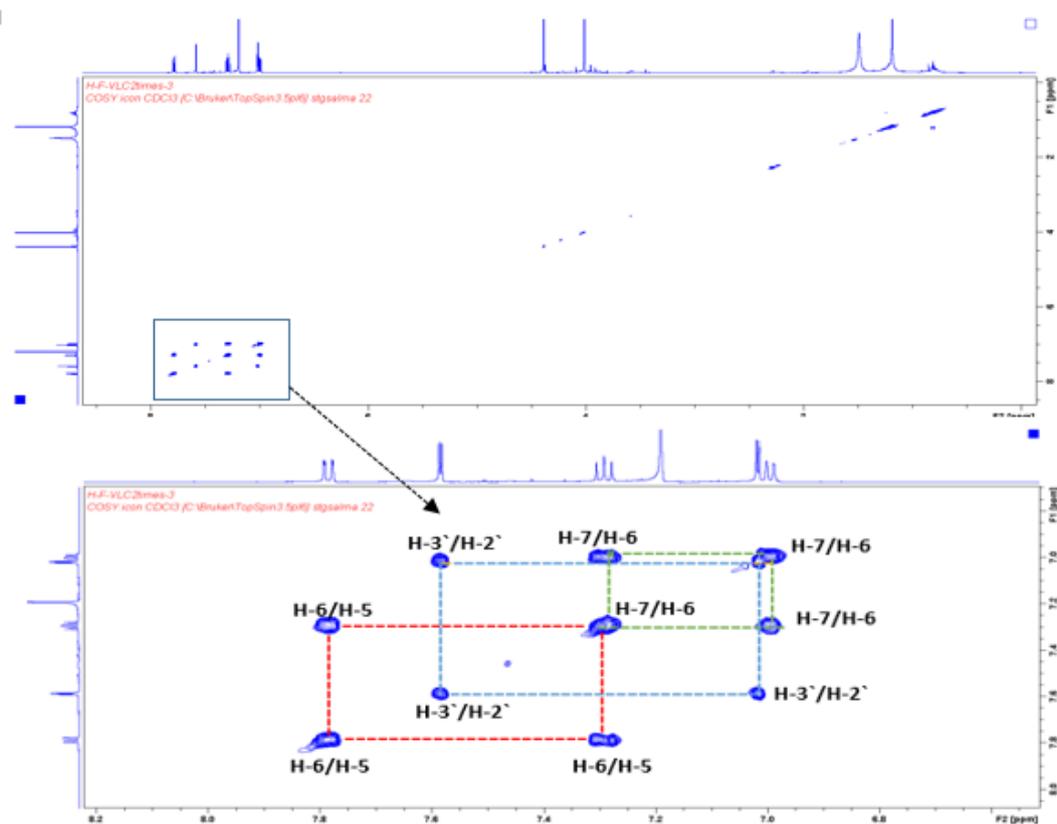


Figure 3.30: COSY NMR spectrum of compound 42

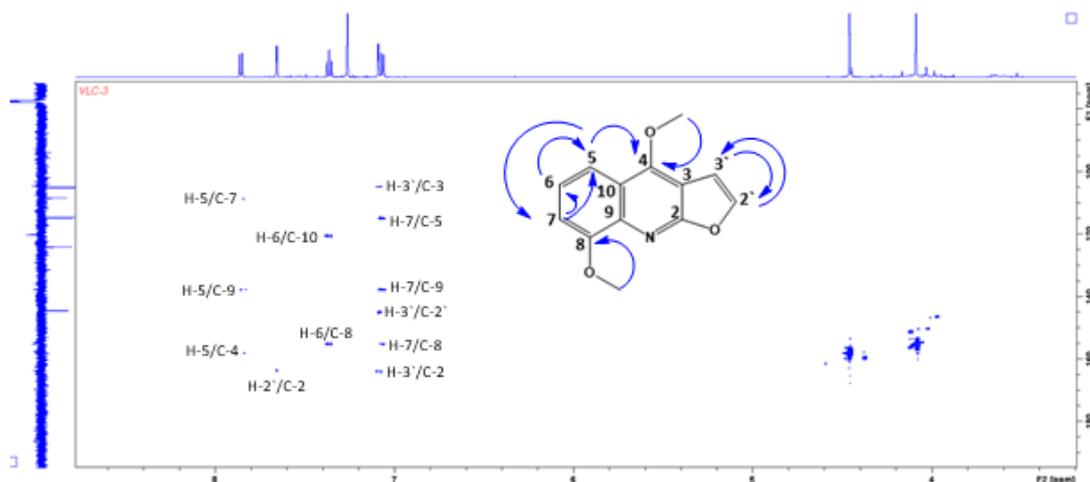


Figure 3.31: The HMBC correlations of compound **42**

3.1.9.4 Structure elucidation of skimmianine (**41**)

The compound **41** was isolated as a colourless powder. The HRESIMS (Figure 3.32) suggested the empirical formula as $C_{14}H_{13}NO_4$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 260.0919 (calculated 260.0923). The 1H NMR spectra (Figure 3.33, Table 3.9) showed three sets of methoxyl signals at δ_H 3.99 (3H), 4.04 (3H) and 4.51 (3H) and *ortho*-coupled ($J= 9.2$ Hz) protons at δ_H 7.40 (H-6) and 8.08 (H-5) also, two olefinic signals at δ_H 7.35 (H-3') and δ_H 7.78 (H-2') confirming the formation of furan ring. The ^{13}C -NMR (Figure 3.34, Table 3.9) spectra showed a total of fourteen carbons including three methoxyl (OMe) and seven quaternary carbons. In the HSQC, H-2', H-2', H-5 and H-6 exhibited direct connectivity to the corresponding methine carbons δ_C 106.5, 144.6, δ_C 119.8 and 113.6 whilst the methoxyl group protons at δ_H 3.99, 4.04 and 4.51 revealed direct connection to the carbons at δ_C 61.9, 57.4 and 60.0. In the HMBC experiment, H-5, H-3' and methoxyl protons at δ_H 4.51 revealed 3J correlation to the oxygenated quaternary carbon at δ_C 159.4 while the methoxyl protons at δ_H 4.04 and H-5 showed 3J correlation to another oxygen bearing quaternary carbon at δ_C 154.0. Based on these spectral data and their correlation in the HSQC and HMBC, the compound **41**, quinoline alkaloid, was identified as 5,4,7,8-trimethoxyfuro[2,3-b]quinolone, commonly known as skimmianine. The

spectroscopic data of compound **41** were in a good agreement with respective published data of skimmianine, which is considered one of the common compounds in the Rutaceae family (Ulubelen *et al.*, 1986). Skimmianine (**41**) was isolated from different species of the genus *Ruta*. Previous studies confirmed the anti-inflammatory effect and antimicrobial activity of skimmianine (**41**) (El Sayed *et al.*, 2000; Ratheesh *et al.*, 2013).

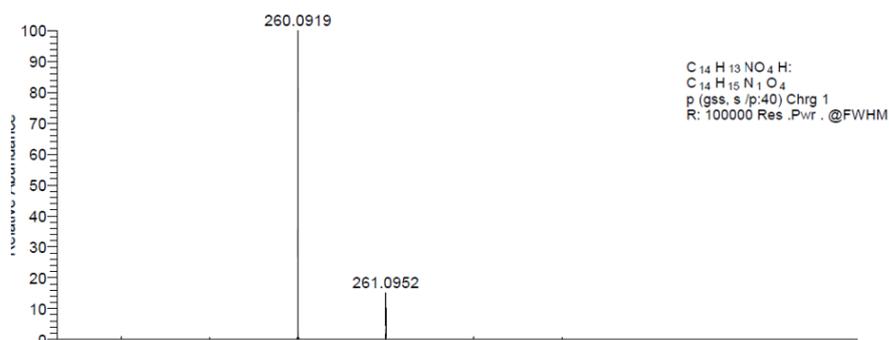


Figure 3.32: The HRESIMS spectrum of compound **41**

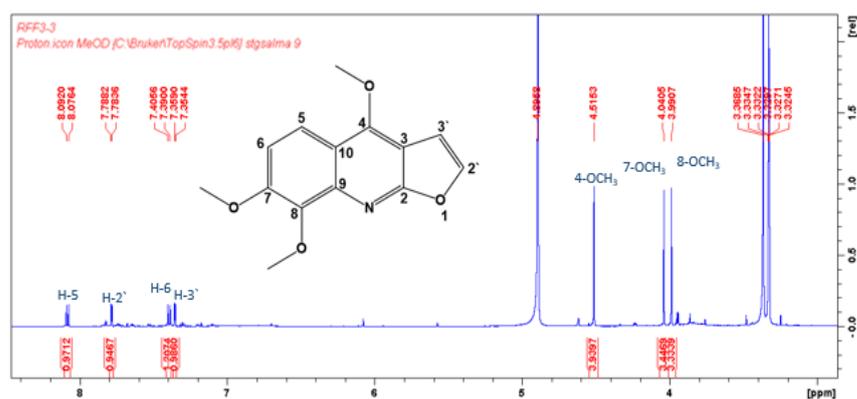


Figure 3.33: ¹H NMR (600 MHz, CD₃OD) spectrum of compound **41**

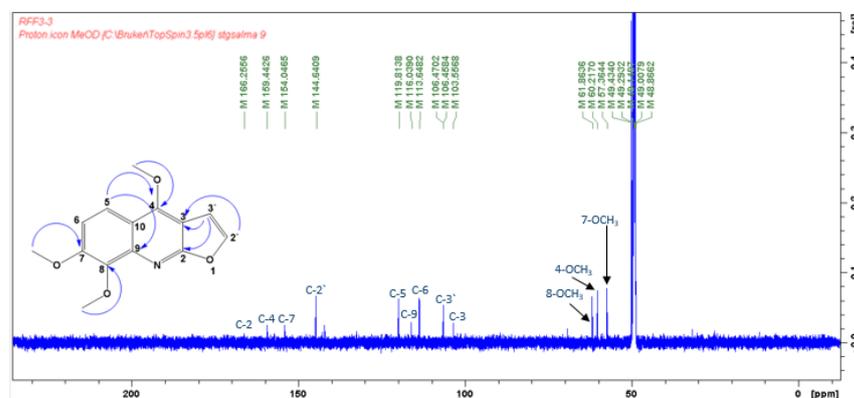


Figure 3.34: The ¹³C NMR (150 MHz, CD₃OD) spectrum of compound **41**

Table 3.9: ^1H NMR (600 MHz) and ^{13}C NMR data of compounds **41, 42, 99,100**

Position	δ_{H} m (<i>J</i> in Hz)				δ_{C}			
	99	100	42	41	99	100	42	41
2	-	-	-	-	163.3	156.1	163.5	166.2
3	-	-	-	-	102.5	106.2	104.3	103.5
4	-	-	-	-	155.8	172.8	157.4	159.4
5	7.48 s	7.87 s	7.85 <i>d</i> (8.34)	8.08 <i>d</i> (9.2)	100.5	107.2	114.3	119.8
6	-	-	7.36 <i>dd</i> (8.34,7.8)	7.40 <i>d</i> (9.2)	148.0	135.9	123.9	113.6
7	-	-	7.06 <i>d</i> (7.8)	-	152.9	154.5	108.3	154.0
8	7.34 s	7.21 s	-	-	107.0	98.4	154.8	144.7
9	-	-	-	-	142.7	146.2	137.5	116.0
10	-	-	-	-	113.2	117.6	120.0	106.4
2'	7.57 <i>d</i> (2.80)	7.60 <i>d</i> (2.22)	7.65 <i>d</i> (2.7)	7.78 <i>d</i> (2.2)	142.7	140.7	144.3	144.6
3'	7.04 <i>d</i> (2.80)	7.09 <i>d</i> (2.22)	7.08 <i>d</i> (2.7)	7.35 <i>d</i> (2.2)	104.8	108.1	105.0	106.5
N- CH ₃	-	4.07 s	-	-	-	32.6	-	-
4- OCH ₃	59.14 s	-	4.45 s	4.51 s	59.1	-	59.4	60.2
6- OCH ₃	4.02 s	3.98 s	-	-	56.2	56.7	-	-
7- OCH ₃	56.30 s	4.07 s	-	4.04 s	56.3	57.0	-	57.4
8- OCH ₃	-	-	4.08 s	3.99 s	-	-	56.3	61.9

3.1.9.5 Structure elucidation of bergapten (**43**)

The compound **43** was isolated as pale yellow crystals. The HRESIMS (Figure 3.36) suggested the empirical formula as C₁₂H₈O₄ and in the positive ion mode it showed [M+H]⁺ peak at *m/z* 217.0495 (calculated 217.0501). The UV spectrum showed absorption maxima (λ_{max}) at 210 nm, typical of a coumarin. The ¹H NMR spectrum (Figure 3.37, Table 3.10) exhibited one aromatic proton as singlet at δ_{H} 7.14, two *ortho*-coupled ($J=9.7$ Hz) protons at δ_{H} 8.15 (H-4) and 6.27 (H-3) and one methoxyl signal at δ_{H} 4.26 (3H). Moreover, the presence of two aromatic protons displayed at δ_{H} 7.59 (1H, d, $J = 2.4$ Hz, H-2') and 7.02 (1H, d, $J = 2.4$ Hz, H-3') in the ¹H NMR spectrum constituted a furan ring. The ¹³C NMR spectrum (Figure 3.38, Table 3.10) presented a total of twelve carbons including a carbonyl at δ_{C} 161.6 (C-2), a methoxyl at δ_{C} 60.5, five methane carbons at δ_{C} 113.0 (C-3), 139.6 (C-4), 94.3 (C-8), 145.1 (C-2'), 105.3 (C-3') and five quaternary carbons at δ_{C} 113.1 (C-6), 158.7 (C-7), 153.0 (C-9), 149.9 (C-5), 106.8 (C-10). In the HMBC experiment (Figure 3.39), long range correlations were observed from H-2' to C-6, C-7 and from H-3 to C-5, C-6 C-7. In the HMBC experiment (Figure 3.44), a common ³*J* correlation from protons at δ_{H} 8.15 (H-4), 7.02 (H-3') and 4.26 (OMe) to an oxygen bearing quaternary carbon at δ_{C} 149.9 confirmed the presence of methoxyl group at C-5. The H-4 also showed ³*J* correlation to the carbonyl δ_{C} 161.6 (C-2) and another oxygenated quaternary at δ_{C} 153.0 (C-9) while the H-3 revealed ³*J* correlation to a quaternary at δ_{C} 106.8 (C-10). In addition, H-8 displayed HMBC connectivity to C-9 (by ²*J*), C-7 (δ_{C} 158.7; ²*J*), C-6 (δ_{C} 113.1; ³*J*) and C-10 (δ_{C} 106.8; ³*J*). Thus, compound **43** was identified as bergapten, a common furocoumarin present in the plant family, Rutaceae. The spectroscopic data of compound **43** were in a good agreement with respective published data of bergapten (Um *et al.*, 2010; O'Neill *et al.*, 2013; Dehghan *et al.*, 2017). Bergapten (**43**) was previously isolated from *Ruta* and *Citrus* species (Dreyer, 1966; Gonzalez *et al.*, 1977). The researchers also reported the cytotoxic activity of

bergapten against different cancer cell line including ovarian, prostate, lung and breast (De Amicis, 2015). The antimicrobial activity of bergapten (**43**) against *E. coli*, *S. aureus* and *H. pylori* was confirmed (Santoro *et al.*, 2016).



Figure 3.35: The HRESIMS spectrum of compound **43**

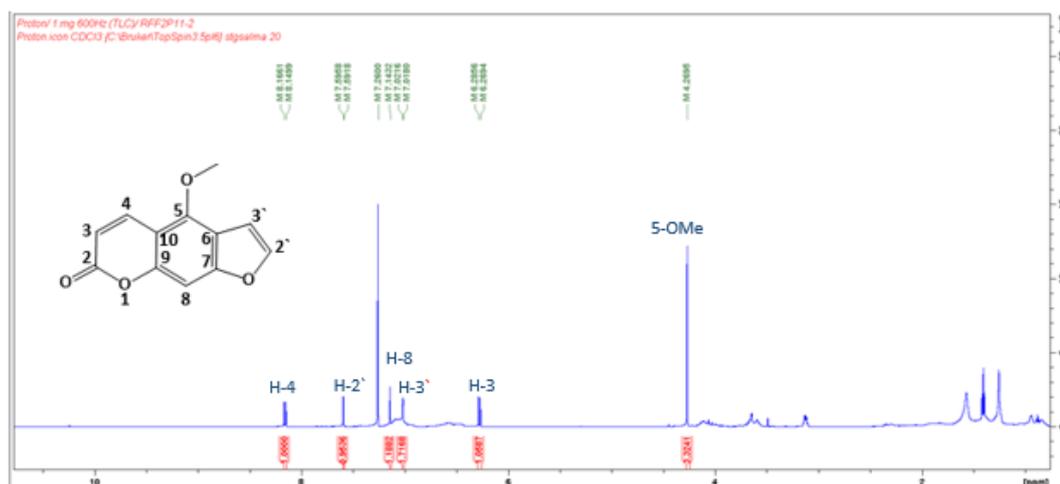


Figure 3.36: ¹H NMR (600 MHz, CDCl₃) spectrum of compound **43**

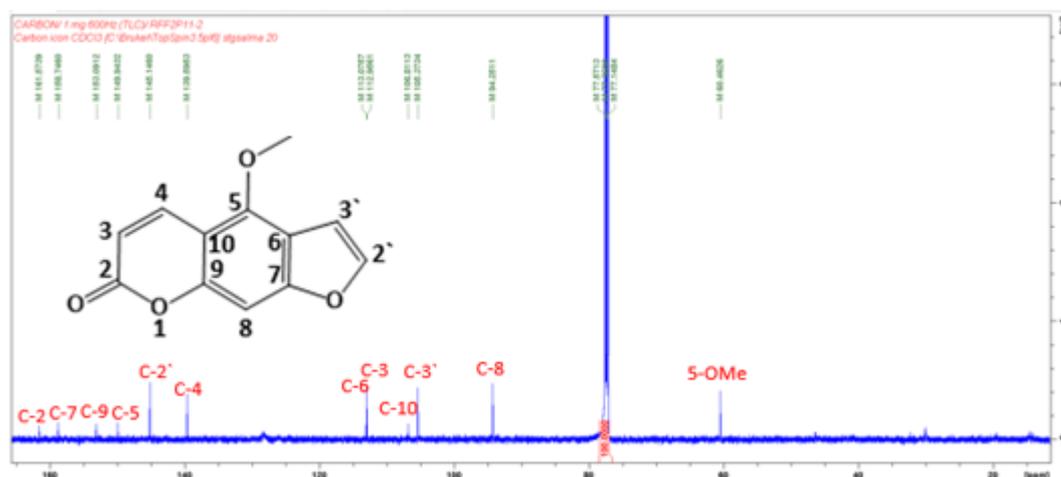


Figure 3.37: The ¹³C NMR (150 MHz, CDCl₃) spectrum of compound **43**

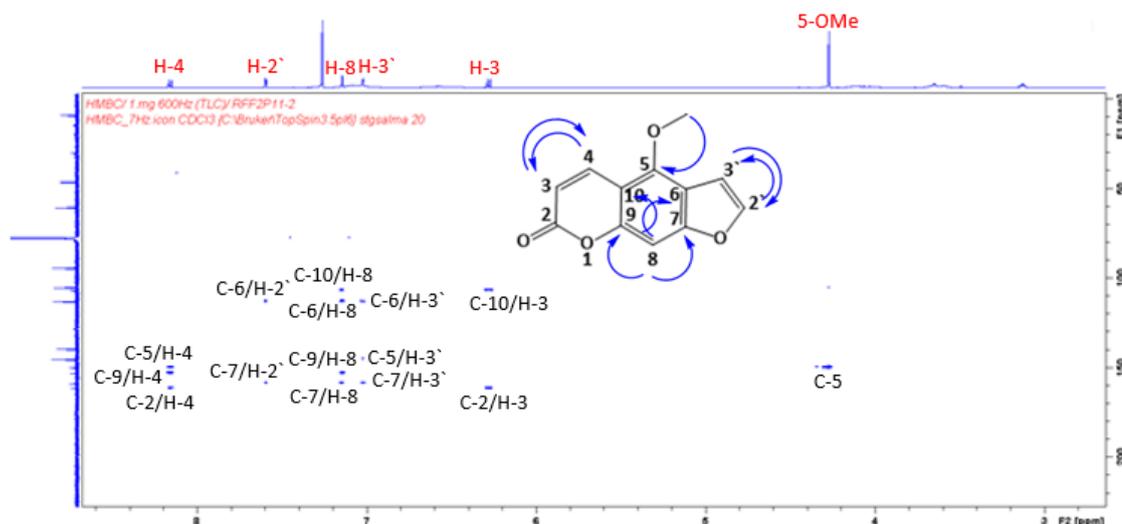


Figure 3.38: The HMBC correlations of compound **43**

3.1.9.6 Structure elucidation of isopimpinellin (**44**)

The compound **44** was isolated as colourless needles. The HRESIMS (Figure 3.40) suggested the empirical formula as $C_{13}H_{10}O_5$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 247.0604 (calculated 247.0606). The 1H NMR and ^{13}C NMR spectra of compound **44** are very similar to those of **43**. The 1H NMR spectrum (Figure 3.41, Table 3.10) displayed two sets of methoxy signals at δ_H 4.13 (3H) and 4.23 (3H) and four aromatic methines at δ_H 6.32 (1H, d, $J=9.78$ Hz), 8.28 (1H, d, $J=9.78$ Hz), 7.85 (1H, d, $J=2.34$ Hz) and 7.25 (1H, d, $J=2.34$ Hz). The DEPTQ spectrum (Figure 3.42, Table 3.10) revealed a total of thirteen carbons comprising one carbonyl at δ_C 162.8, six quaternary carbons, four aromatic methines at δ_C 113.3, 141.6, 147.2, 106.5 and two methoxy carbons at δ_C 61.6 and 62.2. In the HMBC experiment, the methoxyl signals at δ_H 4.22 and 4.13 revealed 3J correlation to the oxygen bearing quaternary carbons at δ_C 145.6 (C-5) and 192.6 (C-8), respectively and thereby confirmed their positions in the molecule. The H-3 exhibited 3J correlation to a quaternary carbon at δ_C 108.8 (C-10) while H-4 presented 3J correlation to C-5 and also the carbonyl group at δ_C 162.8 (C-2) and oxygenated quaternary carbon at δ_C 145.0 (C-9). Moreover, H-2' showed HMBC connectivity to C-3' (106.5 by 2J) and C-6 (116.5 by 3J) whilst H-3' exhibited HMBC correlation to C-5 (3J) and C-7

(151.8; by 3J). Thus, compound **44** identified as 5,8-dimethoxy-6,7-furanocoumarin or isopimpinellin. The spectroscopic data of compound **44** were in a good agreement with respective published data of isopimpinellin (Wu *et al.*, 2003; Shu Shan *et al.*, 2013). The antimicrobial activity of isopimpinellin (**44**) was reported to prevent the growth of *Cryptococcus neoformans* with IC_{50} 40 mg/mL (Ngunde Ngwendson *et al.*, 2003) and successively inhibited *Staphylococcus aureus*, *Escherichia coli*, *Pseudomonas aeruginosa* and *Candida albicans* with MIC 3.75, >7.5, >7.5 and 0.93, respectively (Golfakhrabadi *et al.*, 2016).

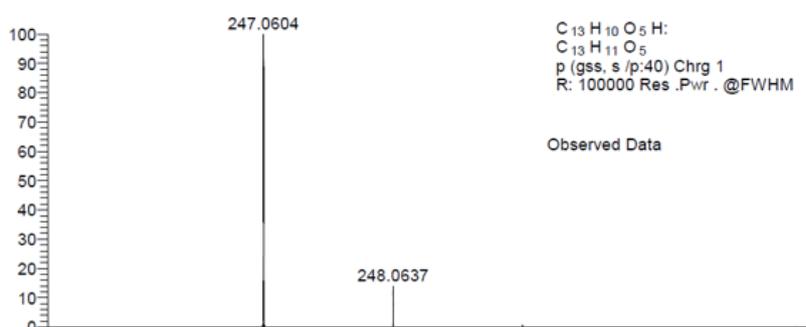


Figure 3.39: The HRESIMS spectrum of compound **44**

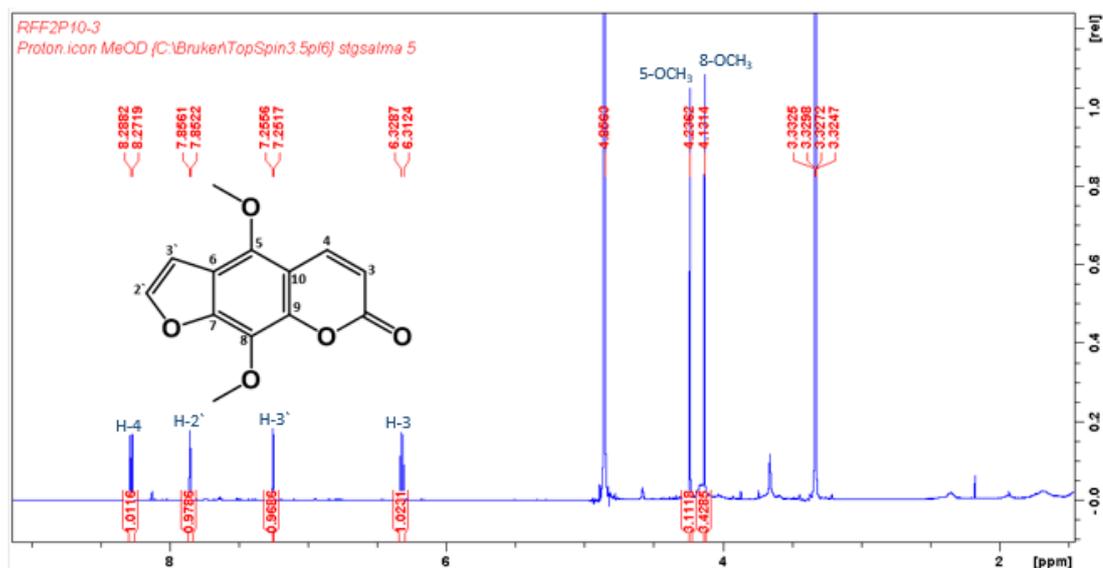


Figure 3.40: 1H NMR (600 MHz, CD_3OD) spectrum of compound **44**

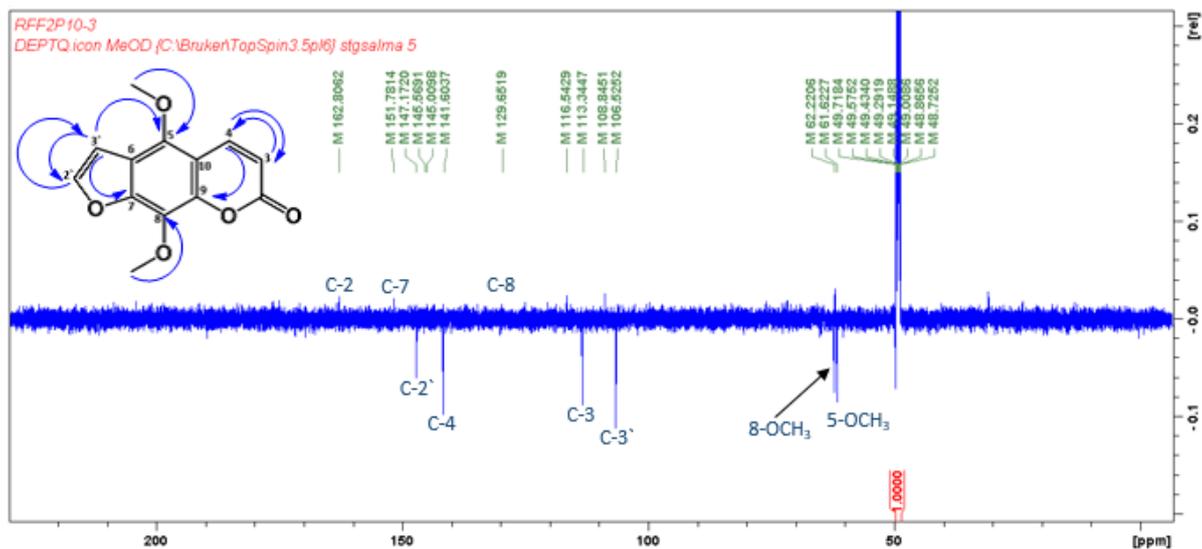


Figure 3.41: DEPT NMR (150 MHz, CD_3OD) spectrum of compound **44**

3.1.9.7 Structure elucidation of chalepin (**45**)

The compound **45** was isolated as white crystals. The HRESIMS (Figure 3.43) suggested the empirical formula as $C_{19}H_{22}O_4$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 315.1596 (calculated 315.1596). The 1H NMR spectra (Figure 3.44, Table 3.10) exhibited three sets of methyl signals at δ_H 1.22 (3H), 1.36 (3H) and 1.46 (2X 3H) and three aromatic methine signals at δ_H 7.47 (H-5), 7.19 (H-4) and 6.70 (H-9). Moreover, the 1H NMR presented three olefinic protons at δ_H 6.15 (1H, dd, $J=17.4, 10.7$ Hz, H-4'), 4.71 (1H, dd, $J=17.4, 6.00$ Hz, H-2), 5.08 (2H, dd, $J=3.5, 10.6$ Hz, H-4', H-5') and two protons multiplet at δ_H 3.20 (H-3). The ^{13}C NMR spectrum (Table 3.10) showed a total of nineteen carbons including carbonyl at δ_C 160.53 (C-7), four methyls, one methylene, seven quaternary and the remaining carbons methines. In the HMBC experiment, the methyl protons at δ_H 1.46 (H2', 3') showed 3J correlation to quaternary carbon at δ_C 131.2 (C-6) and methine carbon at δ_C 145.9 (C-4') while the methyl groups at δ_H 1.22 (H-2'') and 1.26 (H-3'') revealed 3J correlation to an oxymethine carbon at δ_C 91.2 (C-2). Based on these spectral data and their correlations in the HSQC and HMBC, the compound **45** was assigned as chalepin, a furanocoumarin. The spectroscopic data of compound **45** were in a good

agreement with respective published data of chalepin (Richardson *et al.*, 2016) which is widespread compound on *Ruta* species (Richardson *et al.*, 2016). Chalepin (**45**) was documented to exhibit excellent cytotoxicity against lung cancer cells with an IC₅₀ value 8.69 µg/ml and worked as an antimicrobial agent (Richardson *et al.*, 2016; Tamene and Endale, 2019).

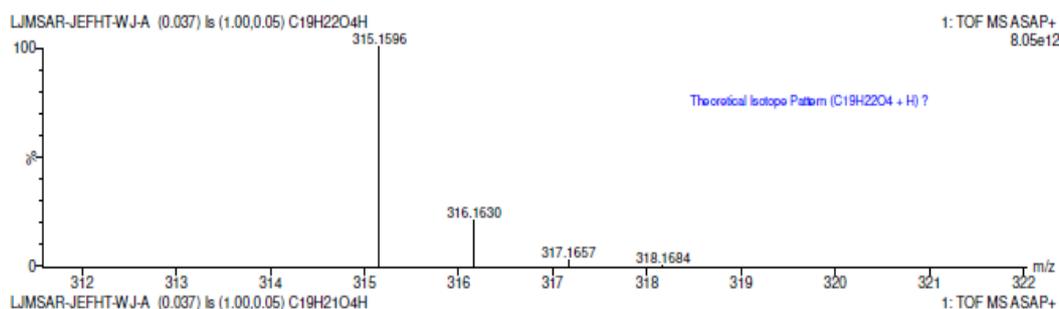


Figure 3.42: The HRESIMS spectrum of compound **45**

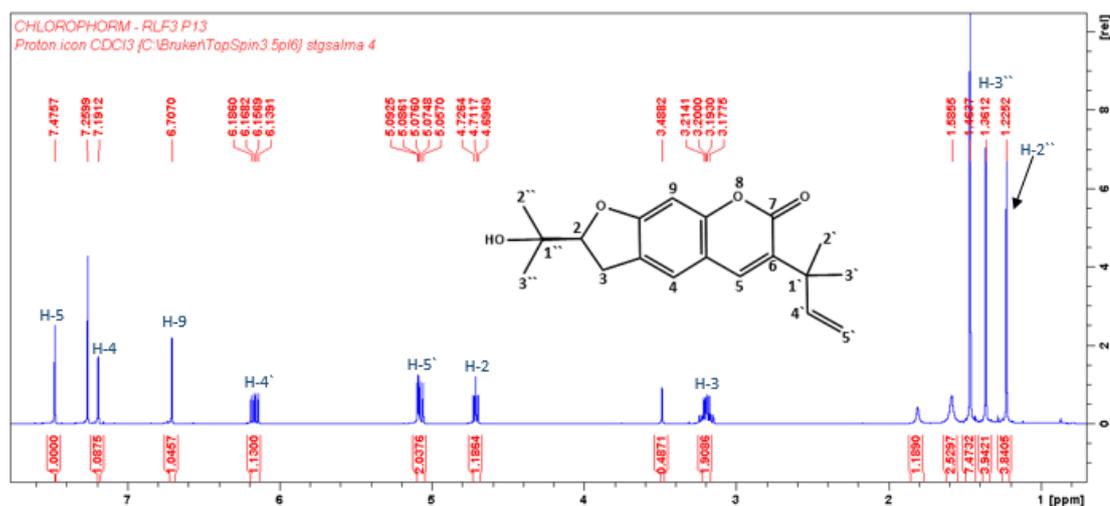


Figure 3.43: ¹H NMR (600 MHz, CDCl₃) spectrum of compound **45**

3.1.9.8 Structure elucidation of rutamarin (**47**)

The compound **47** was isolated as a colourless prism. The HRESIMS (Figure 3.45) suggested the empirical formula as C₂₁H₂₄O₅ and in the positive ion mode it showed [M+H]⁺ peak at *m/z* 357.1702 (calculated 357.1702). The 1D and 2D NMR spectra of compound **47** were similar to chalepin (**45**). The only difference was the presence of an additional signal at δ_H 1.98 (3H, H-5'') in the ¹H NMR spectrum of compound **47** (Figure 3.47, Table 3.10)

and extra peaks for the presence of an acetyl group (δ_C 21.4 for methyl and δ_C 170.6 carbonyl group in the ^{13}C NMR spectrum (Table 3.10). The observation of the cross peak in the HSQC spectrum approved the direct connectivity of proton at δ_H 1.98 (H-5'') to carbon at δ_C 21.4 (C-5''). In the HMBC experiment the methyl groups at δ_H 1.50 (H-2'') and 1.55 (H-3'') exposed 3J correlation to δ_C 170.6 (C-4''). The NMR and mass spectroscopic data as well as the correlation revealed in the HSQC and HMBC confirmed the identification of compound **47** as chalepin acetate or rutamarin (Wu *et al.*, 2003). The spectroscopic data of compound **47** were in a good agreement with respective published data of rutamarin (Wu *et al.*, 2003). The isolation of rutamarin (**47**) was confirmed from the Rutaceae (Adamska-Szewczyk *et al.*, 2016). Rutamarin was shown to inhibit HT29 cells (Colon Adenocarcinoma cell line) with IC_{50} 5.6 μM (Suhaimi *et al.*, 2017) and the antiviral activity of rutamarin was reported (Xu *et al.*, 2014). Rutamarin (**47**) improved glucose homeostasis in mice (Mancuso *et al.*, 2015).

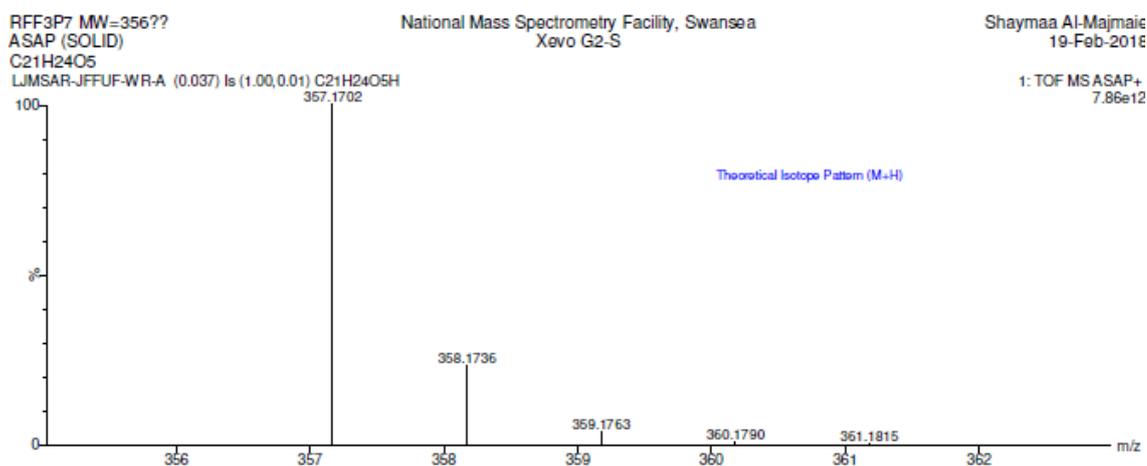


Figure 3.44: The HRESIMS spectrum of compound **47**

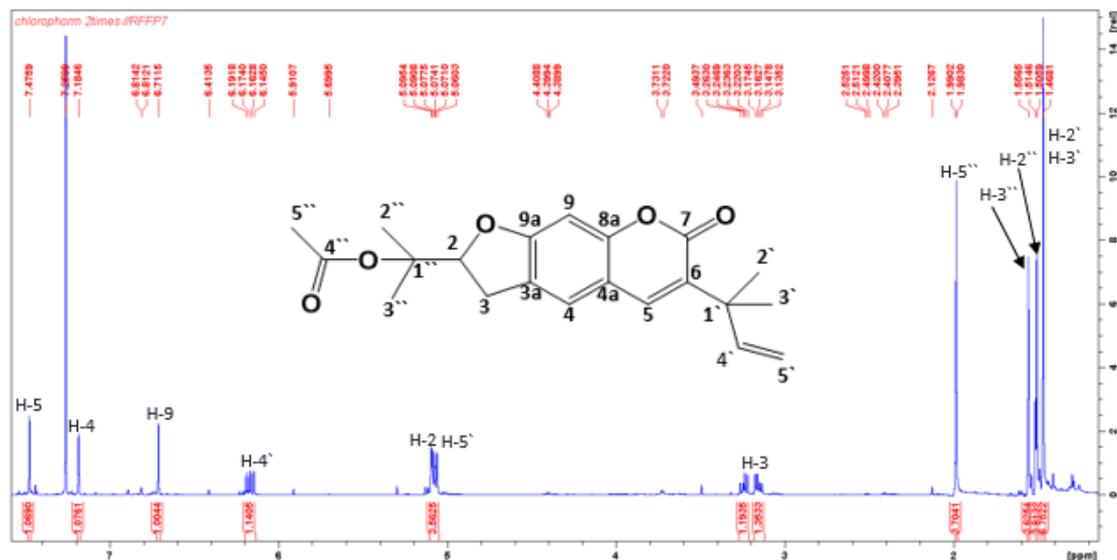


Figure 3.45: ¹H NMR (600 MHz, CDCl₃) spectrum of compound 47

3.1.9.9 Structure elucidation of chalepsin (46)

The compound **46** was isolated as white crystals. The HRESIMS (Figure 3.47) suggested the empirical formula as C₁₆H₁₄O₃ and in the positive ion mode it showed [M+H]⁺ peak at *m/z* 255.1021 (calculated 255.1021). The 1D and 2D NMR spectra of compound **46** were very similar to those of compounds **45** and **47** except for the absence of one side chain. The ¹H NMR spectra for compound **46** (Figure 3.48, Table 3.10) showed *ortho*-coupling of olefinic protons at δ_H 7.66 and 6.81 which confirmed the presence of a furan ring in the molecule. The ¹³C NMR spectra revealed a total of fourteen carbons including two sets of methyl signals at δ_C 26.5, five methine carbons at δ_C 106.7 (H-2), 146.9 (H-3), 119.8 (H-4), 138.6 (H-5) and 99.4 (H-9) and one carbonyl at δ_C 160.2 (C-7). The HSQC experiment revealed the direct connectivity from protons at δ_H 7.66 (H-2) and 6.81 (H-3) to carbons at δ_C 146.9 (C-2) and 106.7 (C-3), respectively. In the HMBC experiment, H-2 revealed ³*J* correlation to the quaternary carbons at δ_C 124.9 (C-4a) and 151.7 (C-9a) while H-3 showed ³*J* to olefinic carbon at δ_C 119.8 (C-4) and quaternary carbons at δ_C 151.7 (C-9a). Based on these spectral data and their correlation in the HSQC and HMBC experiments, the structure of compound **46** was confirmed as chalepsin. The spectroscopic data of compound **46** were in good agreement with respective published data of chalepsin (Wu *et al.*, 2003).

Chalepensin (**46**) was previously isolated from other *Ruta* species (San Miguel, 2003; Adamska-Szewczyk *et al.*, 2016). Chalepensin (**46**) was reported to reduce the activity of the primary enzyme responsible about the oxidation of nicotine and cotinine, which is cytochrome P450 (CYP) 2A69 (Ueng *et al.*, 2011). Chalepensin (**46**) showed significant cytotoxic activity against lung and blood cancer cells (Khlifi *et al.*, 2013; Wannas *et al.*, 2017). One of the traditional uses of *R. chalepensis* was curing tooth pain and the studies revealed that *Streptococcus mutans*, which is found in the cavity of the human mouth and is a major contributor to tooth decay is inhibited by chalepensin with MIC 7.8 $\mu\text{g/mL}$ (Gomez-Flores *et al.*, 2016).

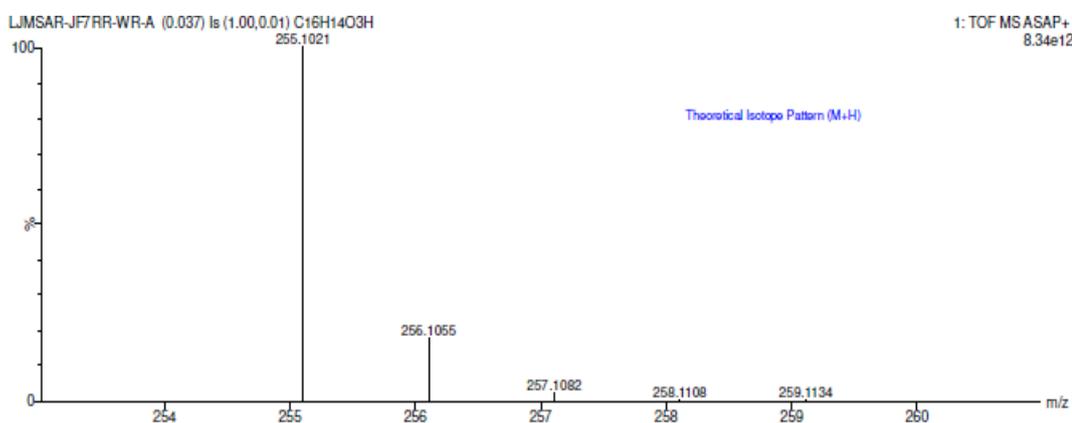


Figure 3.46: The HRESIMS spectrum of compound **46**

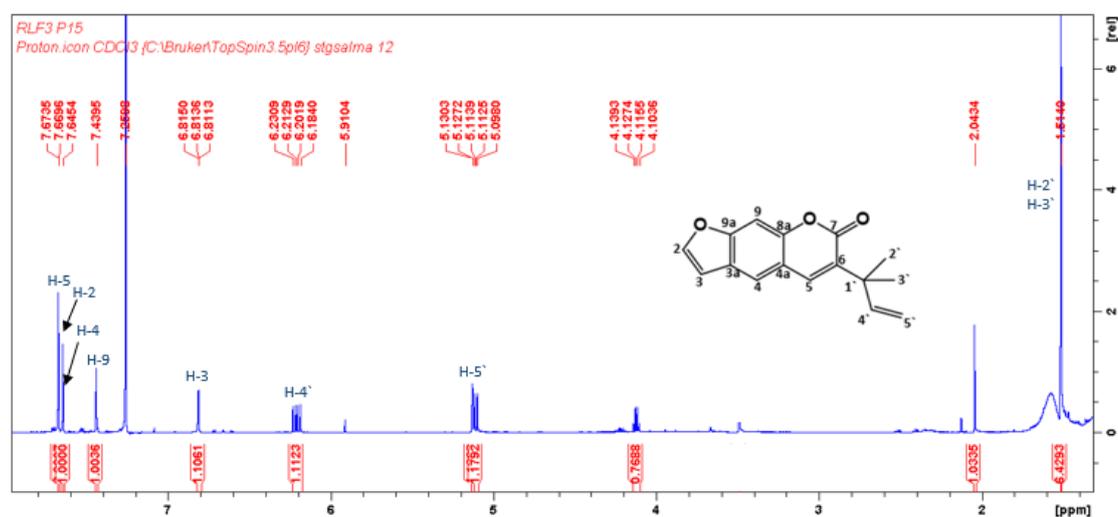


Figure 3.47: ¹H NMR (600 MHz, CDCl₃) spectrum of compound **46**

Table 3.10: ^1H NMR (600 MHz) and ^{13}C NMR data of compounds **43-47**

Position	δ_{H} (J in Hz)					δ_{C}				
	45	46	47	43	44	45	46	47	43	44
2	4.71 <i>dd</i> (17.4)	7.65 <i>d</i> (2.22)	5.07 <i>t</i>	-	-	91.2	146.9	88.3	161.6	162.8
3	3.20 <i>m</i>	6.81 <i>d</i> (2.22)	3.15 <i>m</i> , 3.24 <i>m</i>	6.27 <i>d</i> (9.7)	6.32 <i>d</i> (9.78)	29.9	106.7	29.7	113.0	113.3
3a	-	-	-	-	-	124.9	124.9	124.2	-	-
4	7.19 <i>s</i>	-	7.18 <i>s</i>	8.15 <i>d</i> (9.7)	8.28 <i>d</i> (9.78)	123.6	119.8	123.4	139.6	141.6
4a	-	-	-	-	-	113.5	116.3	112.4	-	-
5	7.47 <i>s</i>	7.64 <i>s</i>	7.47 <i>s</i>	-	-	138.4	138.6	138.3	150.0	145.6
6	-	-	-	-	-	131.2	133.5	131.3	113.1	116.5
7	160.53	-	-	-	-	160.5	160.2	160.5	158.7	151.8
8	-	-	-	7.14 <i>s</i>	-	-	-	-	94.2	129.6
8a	-	-	-	-	-	155.0	151.7	155.1	-	-
9	6.70 <i>s</i>	7.43 <i>s</i>	6.71 <i>s</i>	-	-	97.5	99.3	97.5	153.0	145.0
9a	-	-	-	-	-	162.6	156.2	162.8	-	-
10	-	-	-	-	-	-	-	-	106.8	108.8
1 [`]	-	-	-	-	-	40.6	40.9	40.8	-	-
2 [`]	1.46 <i>s</i>	1.51 <i>s</i>	6.17 <i>m</i>	7.59 <i>d</i> (2.4)	7.85 <i>d</i> (2.34)	26.4	26.5	26.1	145.1	174.1
3 [`]	1.46 <i>s</i>	1.51 <i>s</i>	5.08 <i>m</i>	7.02 <i>d</i> (2.4)	7.25 <i>d</i> (2.34)	26.4	26.5	26.1	105.3	106.5
4 [`]	6.1 <i>m</i>	6.20 <i>m</i>	1.46 <i>s</i>	-	-	145.9	145.8	146.0	-	-
5 [`]	5.08 <i>m</i>	5.11 <i>m</i>	1.46 <i>s</i>	-	-	112.4	112.7	112.0	-	-
1 ^{``}	-	-	-	-	-	72.0	-	82.2	-	-
2 ^{``}	1.22 <i>s</i>	-	1.50 <i>s</i>	-	-	24.5	-	22.3	-	-
3 ^{``}	1.36 <i>s</i>	-	1.55 <i>s</i>	-	-	24.5	-	22.6	-	-
4 ^{``}	-	-	-	-	-	-	-	170.6	-	-

5''	-	-	1.98 s	-	-	-	-	21.4	-	-
5- CH ₃	-	-	-	4.26	24.5	-	-	-	60.5	61.6
8- CH ₃	-	-	-	-	24.5	-	-	-	-	62.2

3.1.9.10 Structure elucidation of imperatorin (**104**)

The compound **104** was isolated as a colourless powder. The HRESIMS (Figure 3.49) suggested the empirical formula as C₁₆H₁₄O₄ and in the positive ion mode it showed [M+H]⁺ peak at *m/z* 271.0966 (calculated 271.0927). The ¹H NMR spectrum (Figure 3.50, Table 3.11) exhibited one aromatic proton at δ_H 7.57 and *ortho*-coupling (*J* = 9.5 Hz) protons at δ_H 8.03 and 6.38; and two olefinic protons at δ_H 7.89 (1H, *d*, *J* = 2.0 Hz), 6.96 (1H, *d*, *J* = 2.0 Hz) as a part of furan ring. The presence of an oxymethylene at δ_H 4.99, an olefinic triplet at δ_H 5.56 and two sets of methyl groups at δ_H 1.68 and 1.72 in the ¹H NMR spectrum confirmed the presence of prenyloxy group in the molecule. The DEPTQ spectrum (Figure 3.51, Table 3.11) displayed a total of sixteen carbons including a carbonyl at δ_C 162.9, two methyl signals at δ_C 26.1, 18.8, one methylene (δ_C 71.0), seven methane and six quaternary carbons. The HMBC experiment played a key role in the confirmation of the structure of compound **104** to be a furanocoumarin. In the HMBC experiment (Figure 3.52), H-5 (δ_H 8.03) showed ³*J* correlation to a methane carbon at δ_C 115.3 (C-4) and the carbonyl at δ_C 162.9 (C-7) while H-3 (δ_H 7.57) exposed ³*J* correlation to quaternary carbon at δ_C 150.2 (9a) and to a methine carbon at δ_C 115.3 (C-4). The H-1' (δ_H 4.98) revealed ³*J* correlation to carbons δ_C 132.5 (C-9) and 141.1 (C-3') and thereby confirmed the connectivity of the prenyloxy group to the molecule through C-9. Thus, compound **104** identified as 9-[(3-methyl-2-buten-1-yl)oxy]-7H-furo[3,2-*g*]chromen-7-one or imperatorin. The spectroscopic data of compound **104** were in a good agreement with respective published data of imperatorin (Shu Shan *et al.*, 2013). Compound **104** inhibited colon cancer cell growth (HT-29) with IC₅₀ value of 78 μM (Zheng *et al.*, 2016). Imperatorin (**104**) was reported to possess antimicrobial properties against *Escherichia coli*, *Aspergillus niger* and *Cladosporium cladosporioides* with MIC 1000 (μg/mL) and inhibited *Bacillus subtilis* with MIC 500 (μg/mL) (Kwon *et al.*, 2002).



Figure 3.48: The HRESIMS spectrum of compound 104

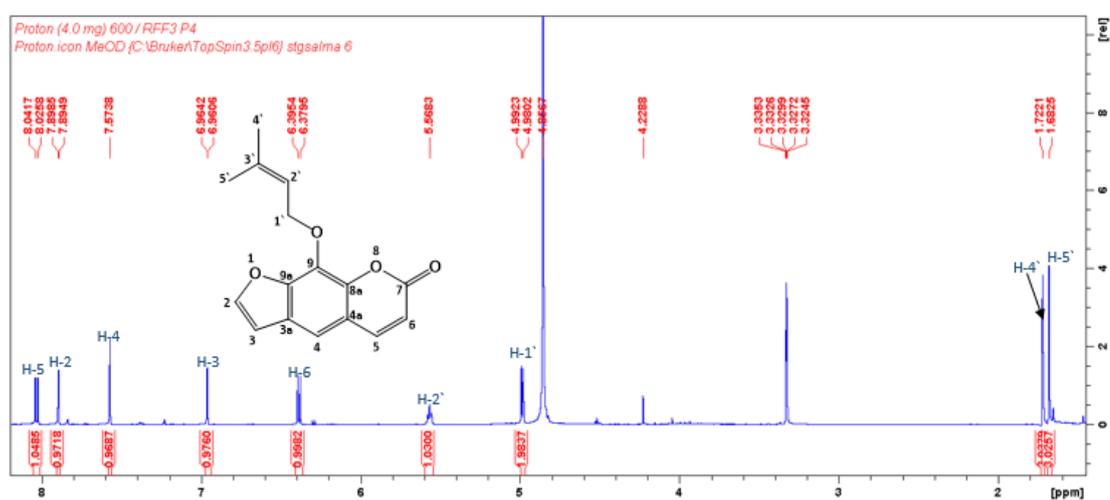


Figure 3.49: 1H NMR (600 MHz, CD_3OD) spectrum of compound 104

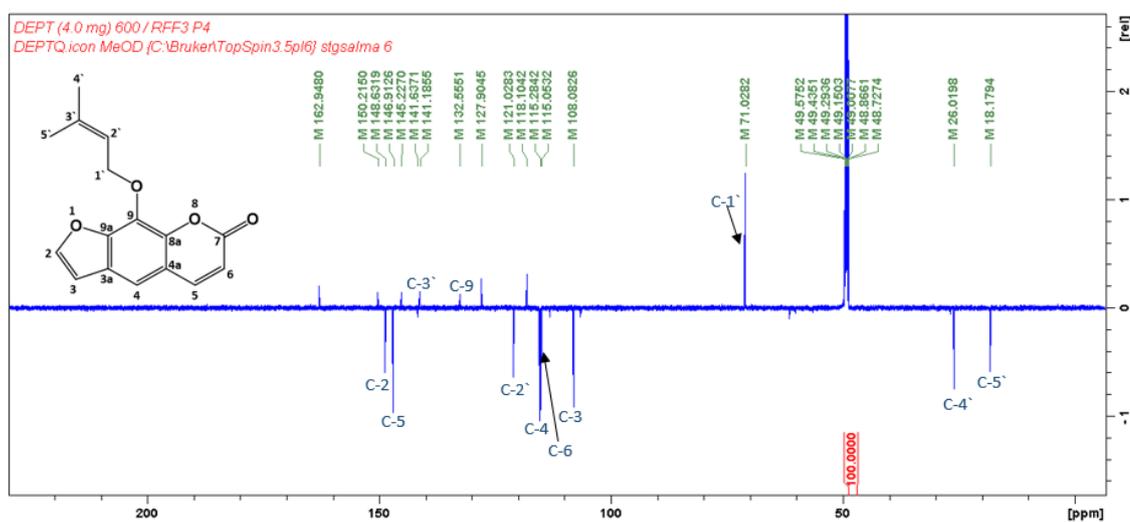


Figure 3.50: DEPT NMR (150 MHz, CD_3OD) spectrum of compound 104

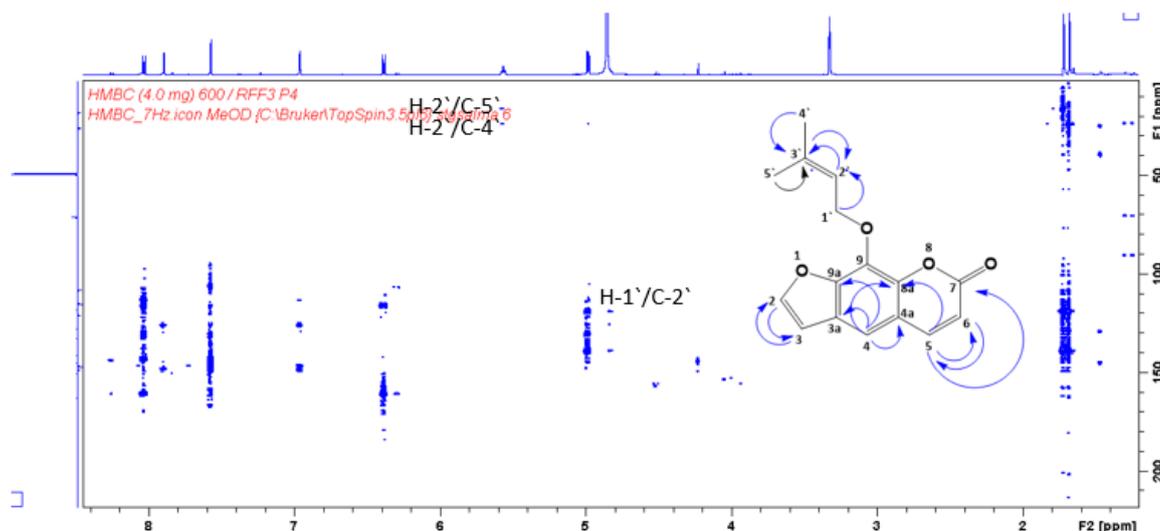


Figure 3.51: The HMBC correlations of compound **104**

Table 3.11: ^1H NMR (600 MHz) and ^{13}C NMR data of compound **104**

Position	(104) Chemical Shift δ (ppm), J in Hz		Position	(104) Chemical Shift δ (ppm), J in Hz	
	^1H	^{13}C		^1H	^{13}C
2	7.89 <i>d</i> (2.0)	148.6	8a	-	145.2
3	6.96 <i>d</i> (2.0)	108.1	9	-	132.5
3a	-	127.9	9a	-	150.2
4	7.57, <i>s</i>	115.3	1'	4.98 <i>t</i>	71.0
4a	-	118.1	2'	5.56 <i>t</i>	121.0
5	8.03 <i>d</i> (9.5)	146.9	3'	-	141.1
6	6.38 <i>d</i> (9.5)	115.0	4'	1.68 <i>s</i>	26.0
7	-	162.9	5'	1.72 <i>s</i>	18.8

3.1.9.11 Structure elucidation of arborinine (**107**)

The compound **107** was isolated as yellow needles. The HRESIMS (Figure 3.53) suggested the empirical formula as $\text{C}_{16}\text{H}_{15}\text{NO}_4$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 286.1074 (calculated 286.1079). The ^1H NMR spectrum (Figure 3.54, Table 3.12) exhibited a hydrogen bonded hydroxyl at δ_{H} 14.74, four aromatic protons as

ABCD pattern at δ_{H} 7.51 (d, $J= 8.7$ Hz), 7.72 (dd, $J= 8.7, 7.4$ Hz), 7.30 (dd, $J= 8.7, 7.4$ Hz), 8.45 (d, $J= 8.1$ Hz), another aromatic proton as singlet at δ_{H} 6.28, three sets of three protons singlets at δ_{H} 3.84 (NMe), 3.93 (OMe), 4.04 (OMe). The ^{13}C NMR spectrum (Figure 3.55, Table 3.12) displayed a total of sixteen carbons including a carbonyl at δ_{C} 181.21, five methine carbons, three sets of methyl carbon signals at δ_{C} 60.15, 56.36 and 34.47 and seven quaternary carbons. In the HMBC experiment (Figure 3.56) the methoxyl group at δ_{H} 3.93 and 4.04 revealed 3J correlations to quaternary carbons at δ_{C} 130.6 (C-2) and 168.1 (C-3), respectively. These correlations confirmed the position of OMe groups through C-2 and C-3 in the molecule. Moreover, 3J correlations from the methyl at δ_{H} 3.84 to quaternary carbons at δ_{C} 140.91 (C-11) and 142.39 (C-14) confirmed it to be the nitrogen bearing methyl group (NMe). Thus, compound **107** was identified as an acridone alkaloid, arborinine. The spectroscopic data of compound **107** were in a good agreement with respective published data of arborinine (Pal *et al.*, 2011). Compound **107** was previously isolated from *Ruta* species (Knölker, 2017; Kumar, 2018). Arborinine (**107**) was reported as an anticancer agent against HeLa (cervical cancer cells), MCF7 (breast cancer cell) and A431 cells (skin cancer) (Réthy *et al.*, 2007). The antimicrobial activity of arborinine (**107**) was confirmed against *Escherichia coli* and *Shigella dysenteriae* with zones of inhibition of 7 mm and 9 mm respectively. Moreover, arborinine (**107**) inhibited *Klebsiella aerogenes* and *Providencia stuartii* with MIC 256 $\mu\text{g}/\text{mL}$ and 512 $\mu\text{g}/\text{mL}$, respectively (Sohrab *et al.*, 2004; Fouotsa *et al.*, 2013).

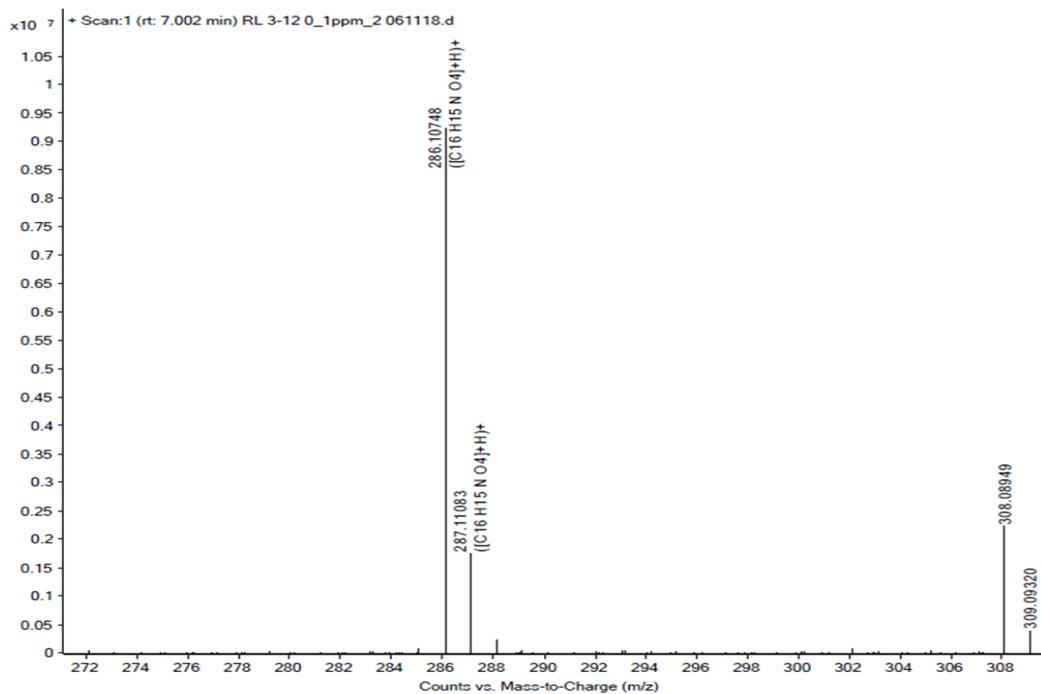


Figure 3.52: The HRESIMS spectrum of compound 107

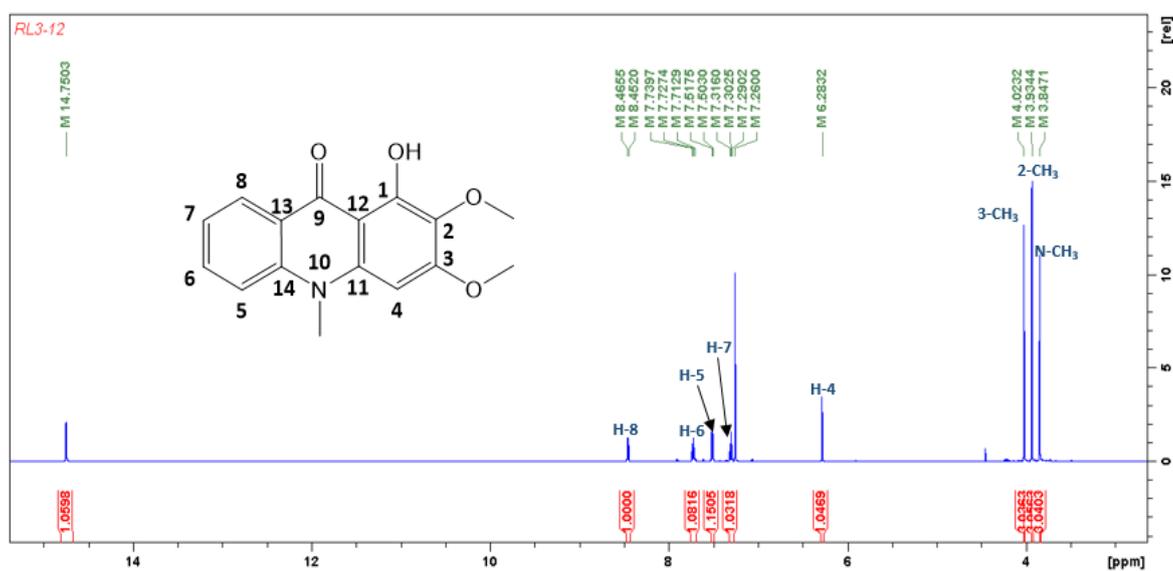


Figure 3.53: ¹H NMR (600 MHz, CHCl₃) spectrum of compound 107

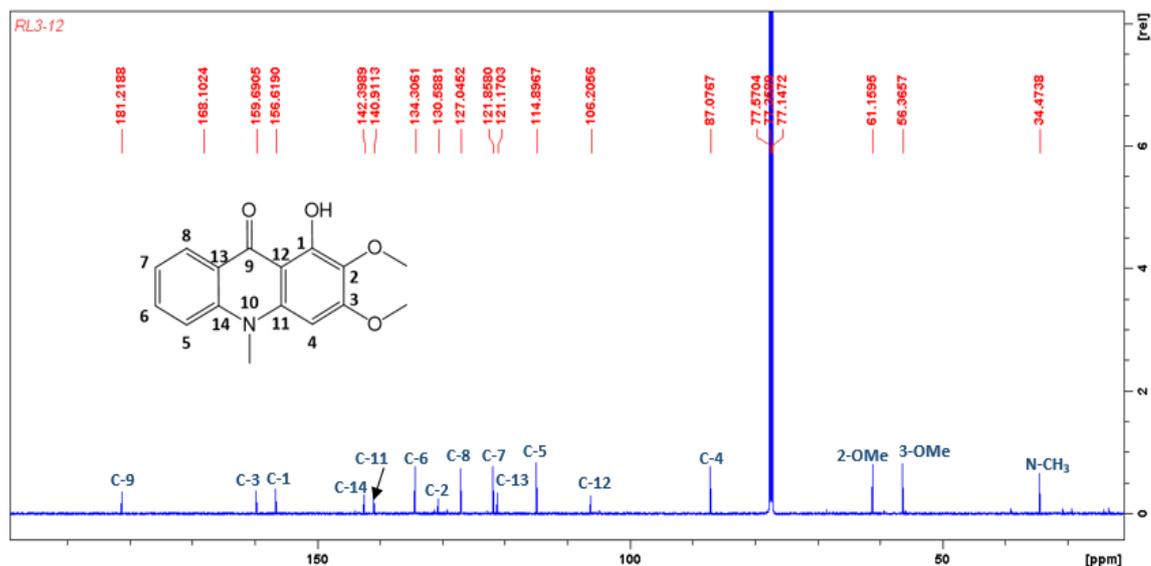


Figure 3.54: ^{13}C NMR (150 MHz, CHCl_3) spectrum of compound 107

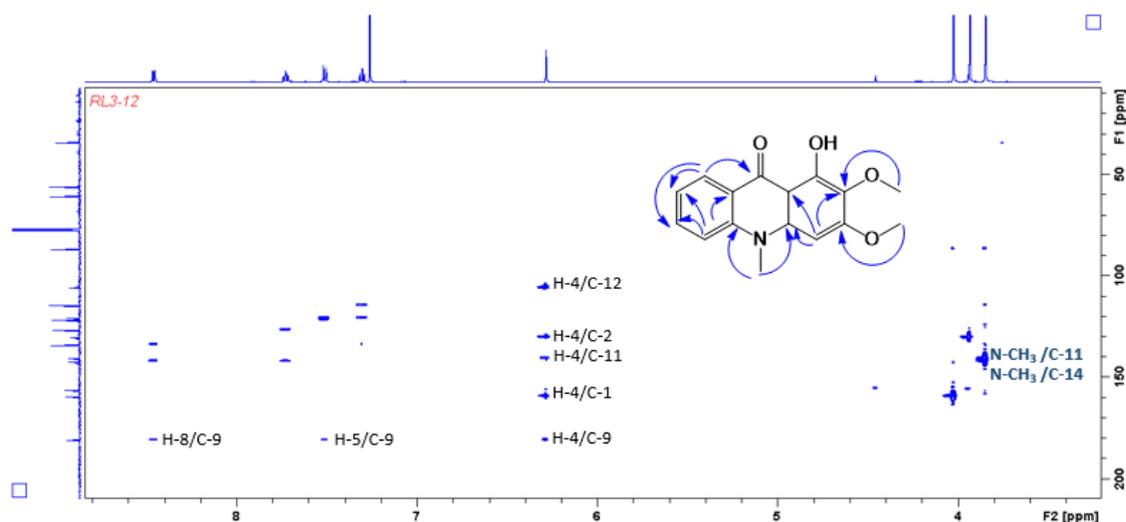


Figure 3.55: The HMBC correlations of compound 107

Table 3.12: ^1H NMR (600 MHz) and ^{13}C NMR data of compound 107

Position	(107) Chemical Shift δ (ppm), J in Hz		Position	(107) Chemical Shift δ (ppm), J in Hz	
	^1H	^{13}C		^1H	^{13}C
1	-	159.7	9	-	181.2
2	-	130.6	11	-	140.9
3	-	168.1	12	-	106.2
4	6.28 s	87.1	13	-	121.2
5	7.51 <i>d</i> (8.7)	114.9	14	-	142.4
6	7.72 <i>dd</i> (8.7, 7.38)	134.3	3-OCH ₃	4.02, s	56.4
7	7.30 <i>dd</i> (8.7, 7.38)	121.8	2-OCH ₃	3.93, s	61.1
8	8.45 <i>d</i> (8.1)	127.0	N-CH ₃	3.84, s	34.5

3.1.9.12 Structure elucidation of ribalinium (106)

The compound **106** was isolated as a dark brown amorphous powder. The HRESIMS suggested the empirical formula as $C_{16}H_{20}NO_4$ and in positive ion mode it showed $[M+H]^+$ peak at m/z 291.1401 (calculated 291.1470). The 1H NMR spectrum (Figure 3.57, Table 3.13) exhibited four sets of methyl signals including deshielded methyls at δ_H 4.07 and 4.51, three aromatic protons as ABX pattern δ_H 7.89 (d, $J = 8.9$ Hz), 7.47 (dd, $J = 8.9, 2.0$ Hz) and 7.57 (d, $J = 2.0$ Hz) and signals for a methylene (δ_H 3.90) and oxymethine (δ_H 5.22). The ^{13}C NMR spectrum (Figure 3.58) exhibited a total of sixteen carbons including seven quaternary carbons, three aromatic methines at δ_C 108.6 (C-5), 119.4 (C-8), 124.7 (C-7), one methylene at δ_C 30.2 (C-3'), one oxymethine at δ_C 95.4 (C-2') and four methyls which were characterized as a *N*-methyl at δ_C 34.9 (δ_H 4.1 from HSQC), two methyl at δ_C 25.0 (C-2''), 26.1 (C-3'') and a methoxy at δ_C 60.4 (4-OMe). In the HMBC experiment, the two sets of methyl protons at δ_H 1.32 (δ_C 25.0) and 1.47 (δ_C 26.1) which existed in *geminal* position were connected to an oxymethine carbon at δ_C 95.41 (C-2') and oxygen bearing quaternary carbon at δ_C 72.07 (C-1'''). The 3J correlations from methyl group at δ_H 4.77 to carbons at δ_C 166.5 (C-2) and 131.85 (C-9) confirmed this as *N*-methyl (NMe) Thus, compound **106** was identified as rabalinium quinoline alkaloids. The spectroscopic data of compound **106** were in a good agreement with published data of rabalinium (Gaston *et al.*, 1980; Pal *et al.*, 2011). Compound **106** was previously isolated from *Ruta graveolens* and reported here for the first time from *R. chalepensis*.

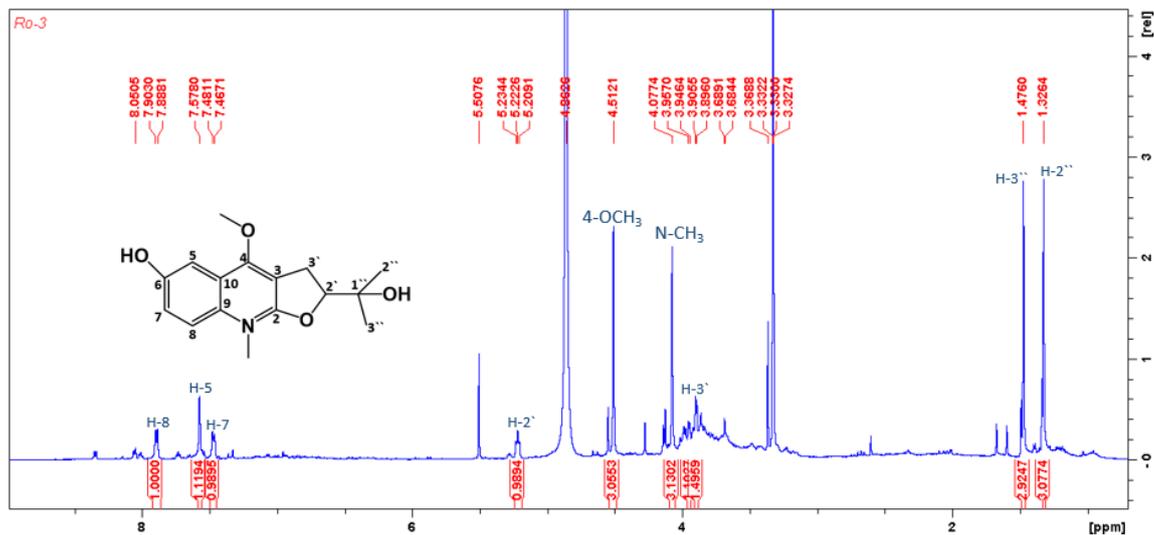


Figure 3.56: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 106

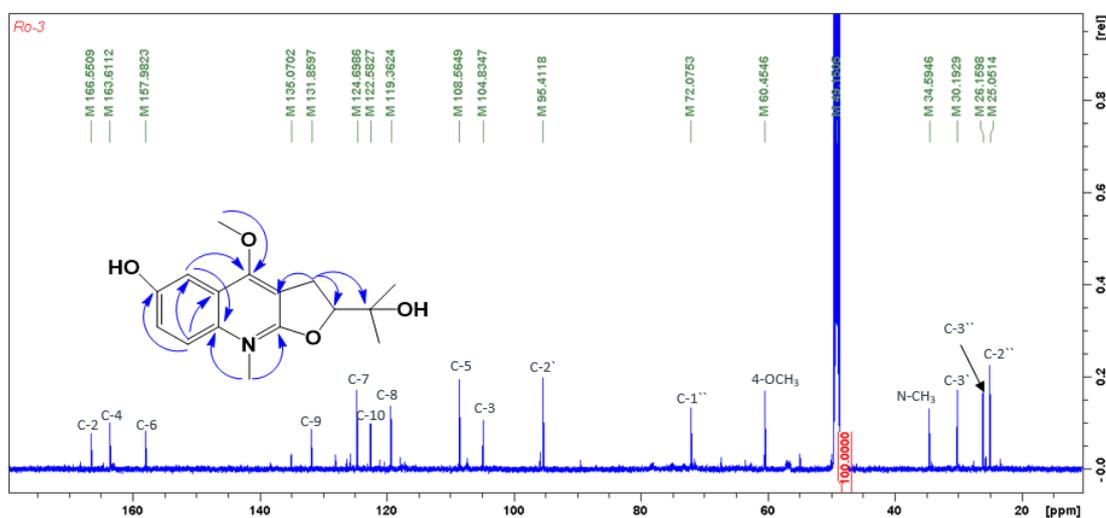


Figure 3.57: ^{13}C NMR (150 MHz, CHCl_3) spectrum of compound 106

Table 3.13: ^1H NMR (600 MHz) and ^{13}C NMR data of compound 106

Position	(106) Chemical Shift δ (ppm), J in Hz		Position	(106) Chemical Shift δ (ppm), J in Hz	
	^1H	^{13}C		^1H	^{13}C
2	-	166.5	10	-	122.6
3	-	104.8	2'	5.22 m	95.4
4	-	163.6	3'	3.90 m	30.2
5	7.57 <i>d</i> (2.00)	108.6	1''	-	72.1
6	-	158.0	2''	1.32 s	25.0
7	7.45 <i>dd</i> (8.94, 2.00)	124.7	3''	1.47 s	26.1
8	7.89 <i>d</i> (8.94)	119.4	4-OCH ₃	4.51 s	60.4
9	-	131.8	N-CH ₃	4.07 s	34.9

3.1.9.13 Structure elucidation of compound 3',6-disinapoylsucrose (108)

The compound **108** (Figure 3.58) was isolated as a white amorphous powder. The HRESIMS suggested the empirical formula as $C_{34}H_{42}O_{19}$ and in the negative ion mode it showed $[M+H]^+$ peak at m/z 753.2300 (calculated 753.2242). The 1H NMR spectrum (Figure 3.59) exhibited four methoxy groups presented by two signals at δ_H 3.71 (6H) and 3.88 (6H), four olefinic protons at δ_H 7.68 (1H, d, $J = 15.7$ Hz, H-7), 7.60 (1H, d, $J = 16.3$ Hz, H-7), 6.48 (1H, d, $J = 16$ Hz, H-8) and 6.46 (1H, d, $J = 16.6$ Hz, H-8), two aromatic methines at δ_H 6.94 (H-2) and 6.88 (H-6). The presence of an anomeric proton at δ_H 5.53 (1H, d, $J = 4.8$ Hz, H-1') suggested the existence of a sugar as part of the molecule. The ^{13}C NMR spectrum (Figure 3.60) revealed a total of thirty four carbons, twelve for sugars, two carbonyls at δ_C 169.2 and 168.4 (C-9) and four carbons of methoxyl groups. The HMBC showed 3J correlation from H-6' to C-9. Also, the key correlation observed from H-3'' to carbons C-8 (3J), C-9 (4J). Moreover, H-1' showed long range correlation to carbon at δ_C 115.8 (C-2''). Thus, compound **108** was identified as a sinapoyl glycoside. The data of compound **108** were in a good agreement with respective data for 3',6-disinapoylsucrose (Chen *et al.*, 2001; Wu *et al.*, 2014; Jin *et al.*, 2016). Sinapoyl glycosides (**108**) was isolated from *Ruta graveolens* and the research suggested it has **bioavailability** effect (Chen *et al.*, 2001; Dhale *et al.*, 2010; Chen *et al.*, 2013). Depending on available data, compound (**108**) isolated for the first time from *R. chalepensis*.

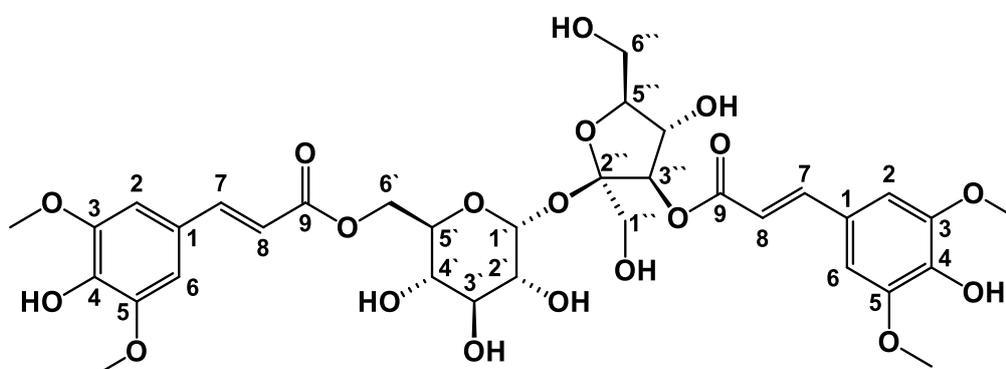


Figure 3.58: Compound 3',6-disinapoylsucrose (**108**) with numbering atoms

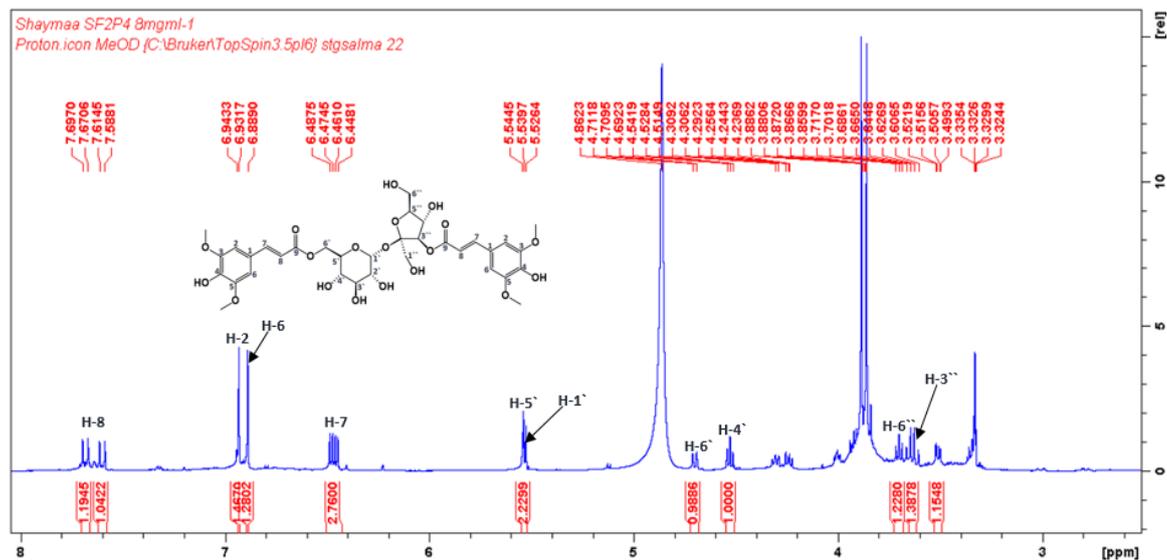


Figure 3.59: ^1H NMR (600 MHz, CDCl_3) spectrum of compound **108**

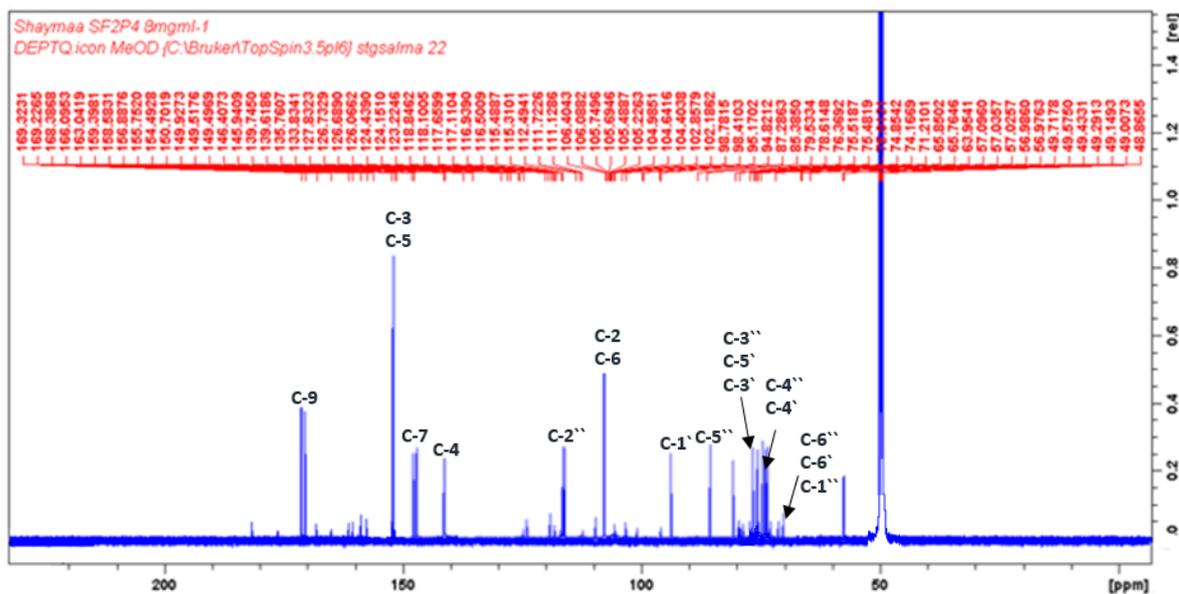


Figure 3.60: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **108**

3.1.9.14 Structure elucidation of graveoline (**105**)

The compound **105** was isolated as a yellow powder. The HRESIMS (Figure 3.62) suggested the empirical formula as $\text{C}_{17}\text{H}_{13}\text{NO}_3$ and in the positive ion mode, it showed $[\text{M}+\text{H}]^+$ peak at m/z 280.0868 (calculated 280.097369). The ^1H NMR spectrum (Figure 3.63, Table 3.14) exhibited a methyl signal at δ_{H} 3.60 (3H), a symmetrical ether at δ_{H} 6.01 and three aromatic protons as ABX pattern resonating at δ_{H} 6.82 (d, $J=1.6$ Hz), 6.84 (d, $J=8.0$ Hz), 6.85 (dd, $J=8.0, 2.1$ Hz). Moreover, four olefinic protons as ABCD pattern indicating

at δ_{H} 7.52 (*d*, $J=8.94$ Hz), 7.36 (*dd*, $J=8.60, 7.84$ Hz), 7.89 (*dd*, $J=7.84, 8.94$ Hz) and 8.42 (*d*, $J=8.60$ Hz) and singlet proton at δ_{H} 6.26. The ^{13}C NMR spectrum (Figure 3.64, Table 3.14) revealed a total of seventeen carbons including a carbonyl at δ_{C} 207.8, one methylenedioxy (δ_{C} 101.7), one methyl (δ_{C} 37.4), eight aromatic methines and five quaternary carbons. In the HMBC experiment, H-7' revealed 3J correlations to carbons at δ_{C} 108.9 (C-3') and 148.8 (C-4') while, H-2' and H-6 exhibited a long range correlation to carbon resonating at δ_{C} 154.6 (C-2). Moreover, a 3J correlation from methyl protons to the quaternary carbons at δ_{H} 154.6 (C-2) and 141.9 (C-8a) in the HMBC experiment confirmed the presence of a nitrogen atom with this methyl (NMe). Thus, compound **105** was identified to be graveoline alkaloid. The data of compound **105** were in a good agreement with respective published data of graveoline (**105**) (Bandatmakuru and Arava, 2018; Kamal *et al.*, 2018; Sampaio *et al.*, 2018). The antimicrobial activity of graveoline (**105**) has confirmed against *Staphylococcus aureus*, *Enterococcus faecalis* and *Escherichia coli* with MIC 1000, 500, 1000 $\mu\text{g/mL}$, respectively (Kamal *et al.*, 2018). Graveoline (**105**) promoted the apoptotic and autophagic cell death for skin melanoma cells, which indicted the anticancer activity (Ghosh *et al.*, 2014).

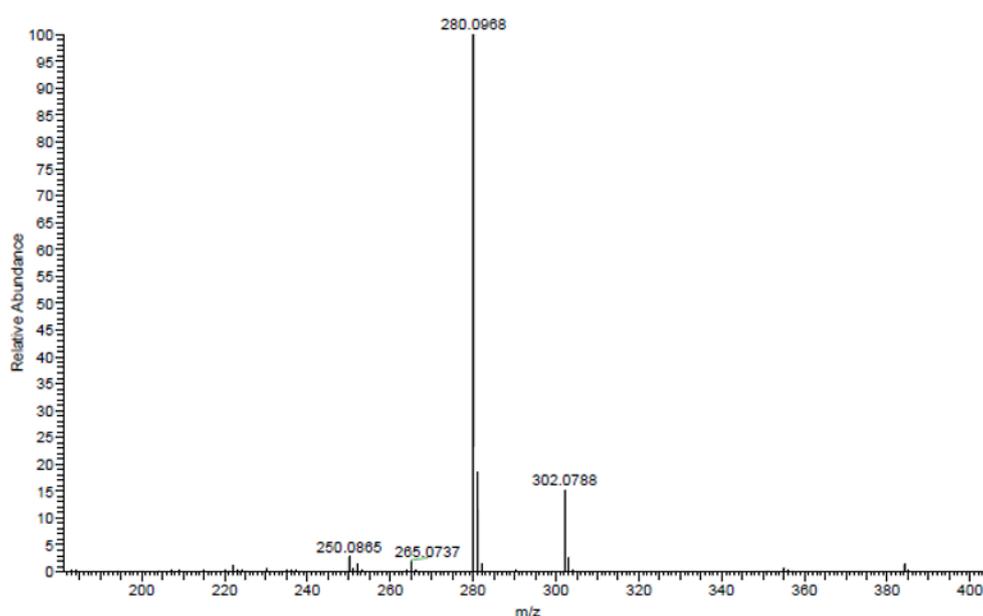


Figure 3.61: The HRESIMS spectrum of compound **105**

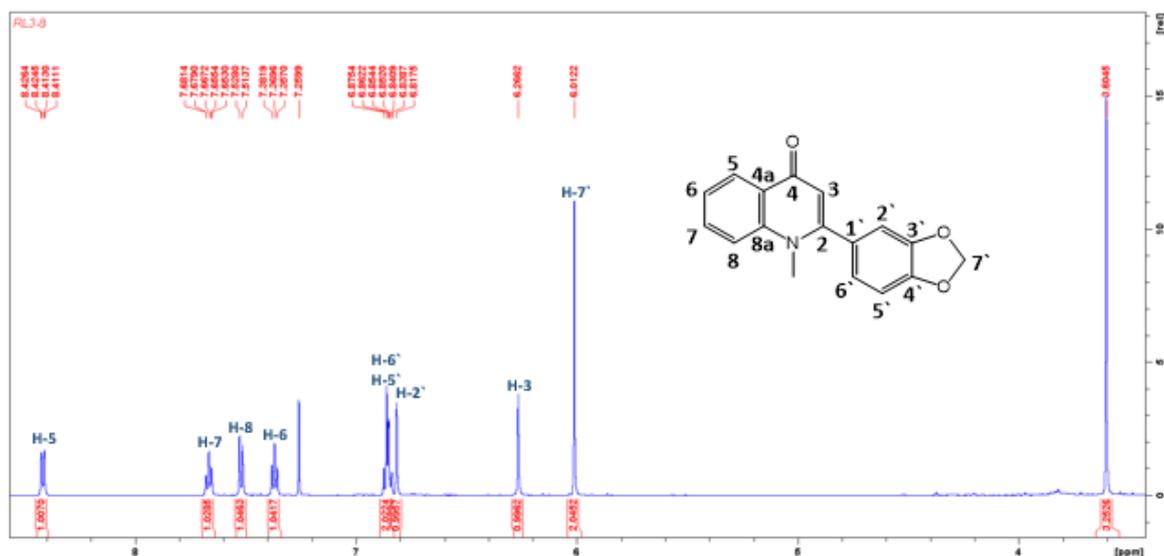


Figure 3.62: ^1H NMR (300 MHz, CDCl_3) spectrum of compound **105**

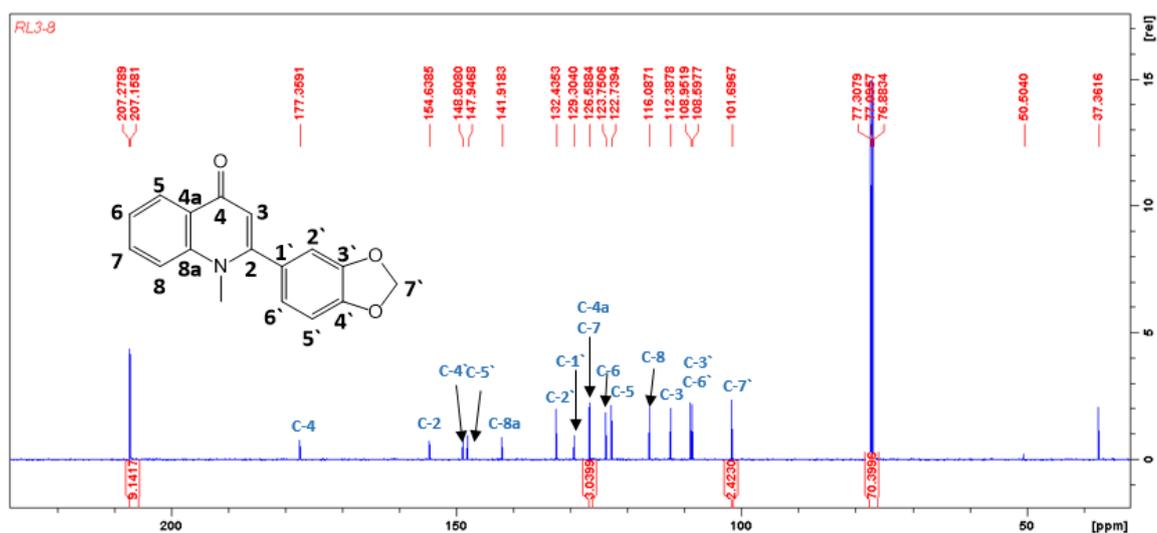


Figure 3.63: ^{13}C NMR (75 MHz, CDCl_3) spectrum of compound **105**

3.1.9.15 Structure elucidation of rutin (**48**)

The compound **48** was isolated as a yellow amorphous solid. The HRESIMS (Figure 3.65) suggested the empirical formula as $\text{C}_{27}\text{H}_{30}\text{O}_{16}$ and in the negative ion mode it showed $[\text{M}-\text{H}]^-$ peak at m/z 609.1461 (calculated 609.1455). The ^1H NMR spectrum (Figure 3.66, Table 3.15) exhibited three aromatic protons as ABX pattern resonating at δ_{H} 7.69 (d , $J = 2.0$ Hz), 6.89 (d , $J = 8.4$ Hz) and 7.65 (dd , $J = 2.0, 8.4$ Hz) and two meta-coupled protons at δ_{H} 6.23 (d , $J = 2.1$ Hz), 6.42 (d , $J = 2.1$ Hz). Moreover, the rutinosyl moiety was revealed by the presence of a glucose anomeric proton at δ_{H} 5.12 (d , $J = 7.7$ Hz), while rhamnose

anomeric proton resonated at δ_{H} 4.54 (*d*, $J=1.5$ Hz) and methyl protons of rhamnose at δ_{H} 1.14 (*d*, $J = 6.2$). The primary elucidation of the ^1H NMR spectrum of compound **48** expected this compound to be flavanone glycoside. The ^{13}C NMR spectrum showed a total of twenty seven carbon signals including carbonyl at δ_{C} 179.5, methoxyl and twelve carbons constituted the rutinosyl moiety. Moreover, five carbons of olefinic protons were recognised from HSQC resonated at δ_{C} 163.2 (C-5), 166.2 (C-7), 146.0 (C-3') and 150.4 (C-4') and the remaining are quaternary carbons. The HMBC experiment showed the long correlation from H-1''' (δ_{H} 4.54) carbons at δ_{C} 68.7 (C-6'') which confirmed the connectivity of glucose and rhamnose. Another 3J correlation from proton at δ_{H} 5.12 (H-1'') to oxygenated quaternary carbon at δ_{C} 135.7 (C-3) confirmed the linkage of sugar through C-3. Thus, compound **48** was identified as rutin. The spectroscopic data of compound **48** was in a good agreement with respective published data of rutin (R. Markham *et al.*, 1987; Li *et al.*, 2014a; Souravh Bais and Abrol, 2016). Rutin (**48**) was documented as an antioxidant agent (Terashima *et al.*, 2012) and showed activity against human lung and colon cancer cell lines by adjusting cells behaviour (Ben Sghaier *et al.*, 2016). The combination of rutin and sunscreens gave strong and safe protective activity on skin (Graziola *et al.*, 2016), which indicated compound **48** may be important in the cosmetic industry.

Table 3.14: ^1H NMR (300 MHz) and ^{13}C NMR data of compound **105**

Position	(105) Chemical Shift δ (ppm), J in Hz		Position	(105) Chemical Shift δ (ppm), J in Hz	
	^1H	^{13}C		^1H	^{13}C
2	-	154.6	1'	-	129.3
3	6.26 s	112.4	2'	6.82 <i>d</i> (1.6)	132.4
4	-	177.3	3'	-	108.9
4a	-	126.6	4'	-	148.8
5	8.42 <i>d</i> (8.60)	122.7	5'	6.84 <i>d</i> (8.0)	147.9
6	7.36 <i>dd</i> (8.60, 7.84)	123.75	6'	6.85 <i>dd</i> (1.6, 8.0)	108.6
7	7.89 <i>dd</i> (7.84, 8.94)	126.6	7'	6.01 s	101.7
8	7.52 <i>d</i> (8.94)	116.1	<i>N</i> -CH3	3.60 s	37.4
8a	-	141.9	-	-	-

respective published data of rutin 3'-methyl ether (R. Markham *et al.*, 1987; Li *et al.*, 2014a; Souravh Bais and Abrol, 2016). Based on available data, this is the first report for isolation of compound **101** from *Ruta chalepensis*.

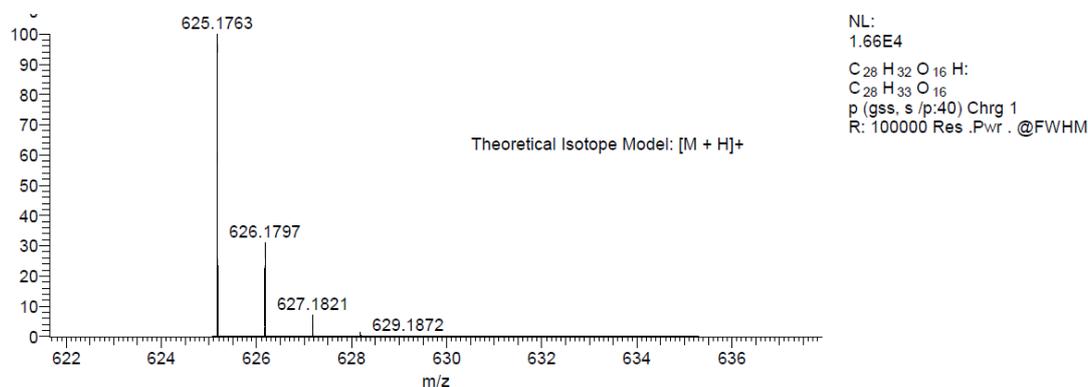


Figure 3.66: The HRESIMS of spectrum compound **101**

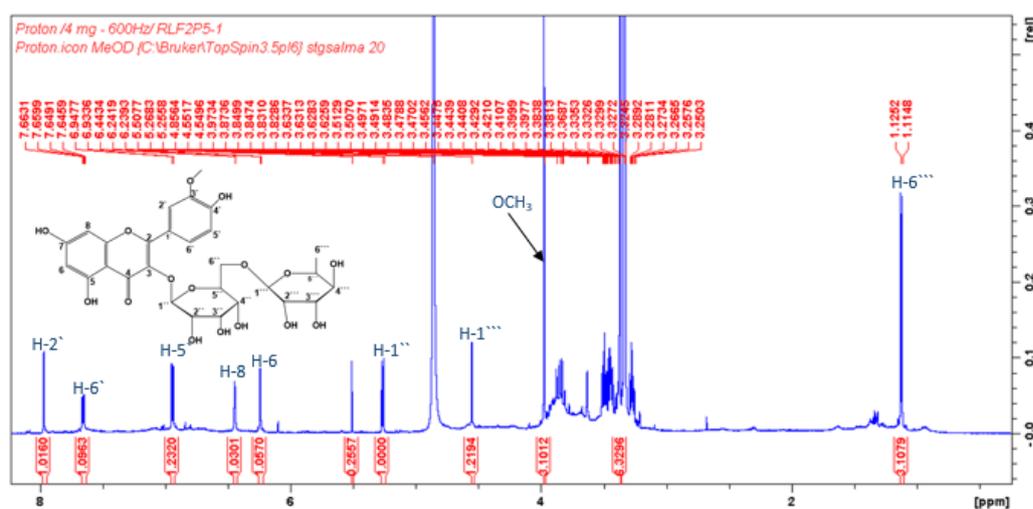


Figure 3.67: ^1H NMR (150 MHz, CD_3OD) spectrum of compound **101**

3.1.9.17 Structure elucidation of rutin-7,4'-dimethyl ether (**102**)

The compound **102** was isolated as a brown amorphous solid. The HRESIMS suggested the empirical formula as $\text{C}_{29}\text{H}_{34}\text{O}_{16}$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 639.1950 (calculated 639.1925). The 1D and 2D spectra of compound **102** were similar to compound **101**. The only difference was the presence two methoxy groups at δ_{H} 3.96 (3H) and 3.97 (3H) in the ^1H NMR spectrum of compound **102** (Figure 3.69, Table 3.15) instead of one methoxy in compound **101**. The ^{13}C NMR and HSQC (Table 3.16)

showed a cross link from oxygenated an methyl at δ_H 3.96, 3.97 to carbons at δ_C 57.4 and 56.1, respectively. The HMBC experiment showed 3J correlation from methoxyl groups at δ_H 3.96, 3.97 to carbons at δ_C 167.3 (C-7) and 151.0 (C-4') respectively, to confirm their position in the molecule. Thus, compound **102** was identified as dimethoxy rutin. The spectroscopic data of compound **102** were in a good agreement with respective published data of rutin 7, 4'-methyl ether (Matsuda *et al.*, 2002). Based on available data, this is the first report of the isolation of rutin-7,4'-dimethyl ether from *R. chalepensis*.

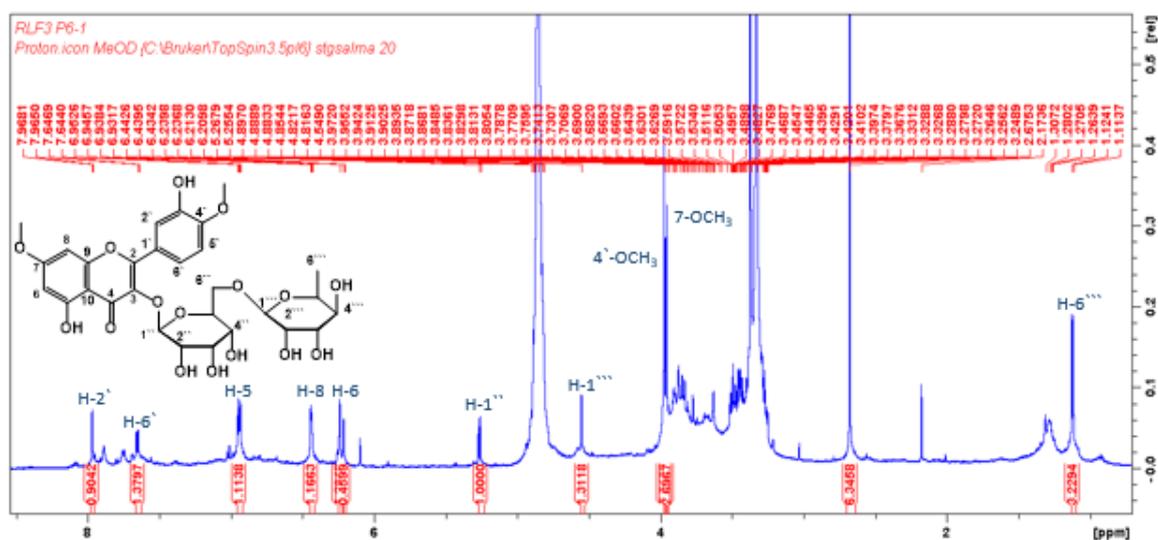


Figure 3.68: 1H NMR (300 MHz, $CDCl_3$) of compound **102**

3.1.9.18 Structure elucidation of 6-hydroxy-rutin 3'-7-dimethyl ether (**103**)

The compound **103** was isolated as a brown amorphous solid. The HRESIMS (Figure 3.70) suggested the empirical formula as $C_{29}H_{35}O_{17}$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 655.1870 (calculated 655.1874). The 1D and 2D spectrum were identical with data of compound **48**, **101** and **102** to confirm the main skeleton of compound **103** to be a flavonoid glucoside. In the 1H NMR spectrum (Figure 3.71, Table 3.15) of compound **103** there was a downfield 6H singlet at δ_H 3.97, and instead of the usual shielded aromatic *meta*-coupled proton signals for H-6 and H-8 in the flavonol skeleton, there was only a 1H singlet at δ_H 6.53 suggesting that one of those protons was substituted by an extra functional group, most likely a hydroxyl group based on the HRESIMS data. The DEPTQ spectrum of

compound **103** (Figure 3.72, Table 3.16) showed the presence of an additional oxygenated aromatic quaternary signal at δ_c 145.3, which could be assigned to C-6 or C-8. Whilst a COSY experiment of compound **103** established all major ^1H - ^1H scalar couplings, ^1H - ^{13}C direct (1J) couplings were obtained from an HSQC experiment confirming the assignment of all protonated carbon signals to their respective ^1H signals. An HMBC experiment displayed all major ^1H - ^{13}C long-range (2J and 3J) correlations (Figure 3.73). The rhamnose anomeric proton (H-1''') showed a 3J correlation to C-6''' of the glucose unit, and the glucose anomeric proton (H-1'') displayed a 3J correlation to C-3 of the flavonol aglycones, confirming the formation of the disaccharide, rutinose, and its connection to C-3 (as in compounds **48**, **101** and **102**). Similarly, 3J correlations from the methoxyl signal at δ_{H} 3.97 (6H) to C-7 and C-3' established methyl ether formation at C-7 and C-3'. The ^{13}C NMR (Table 3.16) signal at δ_c 96.4 confirmed that the hydroxylation was indeed on C-6 in compound **103**, and this fact was further corroborated from the ^1H - ^{13}C HMBC long-range correlations from H-8 signal to C-6 and C-10. Thus, the compound **103** was identified as 6-hydroxy-rutin 3',7-dimethyl ether (**103**), which to the best of our knowledge, is a new natural product.

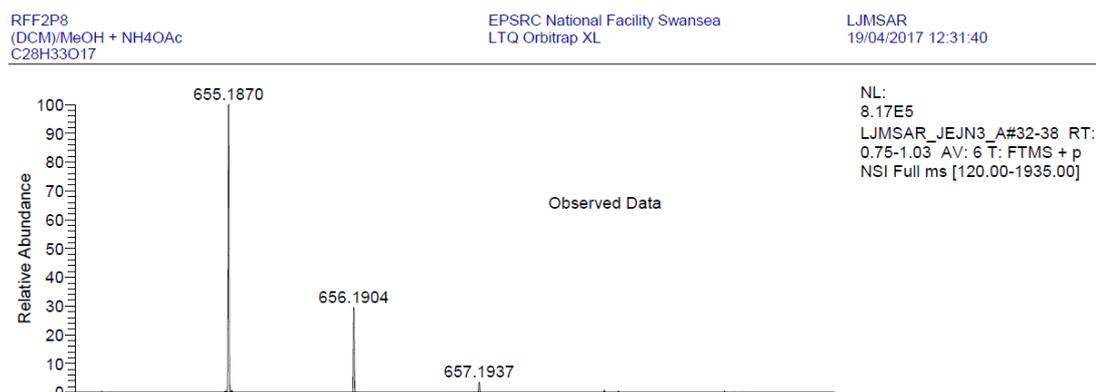


Figure 3.69: The HRESIMS of compound **103**

Table 3.15: ^1H NMR data for compounds **48,101-103** (at 600 MHz in CD_3OD , δ in ppm, J in Hz)

Position	48	101	102	103
6	6.23 <i>d</i> (2.1)	6.23 <i>d</i> (2.0)	6.23 <i>d</i> (2.0)	-
8	6.42 <i>d</i> (2.1)	6.44 <i>d</i> (2.0)	6.43 <i>d</i> (2.0)	6.53 <i>s</i>
2'	7.69 <i>d</i> (2.0)	7.96 <i>d</i> (2.0)	7.66 <i>d</i> (2.0)	8.00 <i>d</i> (2.0)
5'	6.89 <i>d</i> (8.4)	6.94 <i>d</i> (8.4)	6.93 <i>d</i> (8.4)	6.94 <i>d</i> (8.4)
6'	7.65 <i>dd</i> (2.0, 8.4)	7.65 <i>dd</i> (2.0, 8.4)	7.64 <i>d</i> (8.4)	7.80 <i>dd</i> (2.0, 8.4)
7-OMe	-	-	3.96 <i>s</i>	3.97 <i>s</i>
3'-OMe	-	3.97 <i>s</i>	-	3.97 <i>s</i>
4'-OMe	-	-	3.97 <i>s</i>	-
1''	5.12 <i>d</i> (7.7)	5.25 <i>d</i> (7.6)	5.25 <i>d</i> (7.5)	5.25 <i>d</i> , (7.4)
2''	3.49*	3.49*	3.49*	3.49*
3''	3.31*	3.32*	3.32*	3.31*
4''	3.29*	3.30*	3.28*	3.29*
5''	3.43*	3.46*	3.43*	3.43*
6''	3.41* 3.81 <i>m</i>	3.44* 3.84 <i>m</i>	3.44* 3.84 <i>m</i>	3.44* 3.84 <i>m</i>
1'''	4.54 <i>d</i> (1.5)	4.55 <i>d</i> (1.5)	4.54 <i>d</i> (1.4)	4.54 <i>d</i> (1.4)
2'''	3.65*	3.63*	3.63*	3.60*
3'''	3.55*	3.56*	3.53*	3.54*
4'''	3.30*	3.28*	3.28*	3.28*
5'''	3.46*	3.43*	3.43*	3.43*
6'''	1.14 <i>d</i> (6.2)	1.11 <i>d</i> (6.2)	1.12 <i>d</i> (6.2)	1.11 <i>d</i> (6.3)

* Overlapped peaks – confirmed from COSY, HSQC and HMBC experiments

3.1.9.19 Structure elucidation of hexadecane (109)

The compound **109** was isolated as white glittering bright particles solid. . The HRESIMS (Figure 3.74) suggested the empirical formula as $\text{C}_{16}\text{H}_{34}$ from the $[\text{M}+\text{Na}]^+$ peak at m/z 249.1562 (calculated 249.255820). The ^1H NMR spectrum (Figure 3.75) exhibited one huge peak at δ_{H} 1.10, typical of a number of methylenes of a long chain alkane. The DEPTQ spectrum (Figure 3.76) showed the presence of two-methyl groups at δ_{C} 14.10 and a number of methylene groups. The analysis of HMBC correlation along with 1D data observation suggested compound **109** to be a long chain alkane. Based on mass spectrometric and NMR spectroscopic data, compound **109** was identified as hexadecane, which was previously reported from *R. graveolens* (De Feo *et al.*, 2002; Diwan and Malpathak, 2011).

Table 3.16: ^{13}C NMR data for compounds **48**, **101-103** (at 150 MHz in CD_3OD)

Position	Chemical shift δ in ppm			
	48	101	102	103
2	159.5	159.0	159.4	159.3
3	135.7	135.6	135.2	135.3
4	179.5	179.5	180.0	180.0
5	163.2	163.2	161.4	153.2
6	100.0	100.1	98.0	127.8
7	166.2	166.2	167.3	155.5
8	95.0	95.0	92.8	96.4
9	158.5	158.7	158.4	145.3
10	105.7	105.9	104.2	106.0
1'	123.2	123.2	123.2	123.3
2'	117.8	114.7	114.7	114.8
3'	146.0	148.5	148.2	148.5
4'	150.4	151.0	151.0	151.0
5'	116.2	116.3	116.3	116.2
6'	123.7	124.2	124.7	124.7
7-OMe	-	-	57.4	57.2
3'-OMe	-	56.9	-	56.9
4'-OMe	-	-	56.1	-
1''	104.8	104.5	104.5	104.5
2''	75.9	76.1	76.3	76.1
3''	77.4	72.4	72.4	72.4
4''	71.5	77.5	77.5	77.5
5''	78.3	78.3	78.3	78.3
6''	68.7	68.1	68.1	68.7
1'''	102.5	102.7	102.6	102.7
2'''	72.2	72.2	72.2	72.2
3'''	72.4	71.8	71.8	71.8
4'''	74.1	74.0	74.0	74.0
5'''	69.8	69.9	69.98	69.9
6'''	18.0	18.0	18.0	18.0

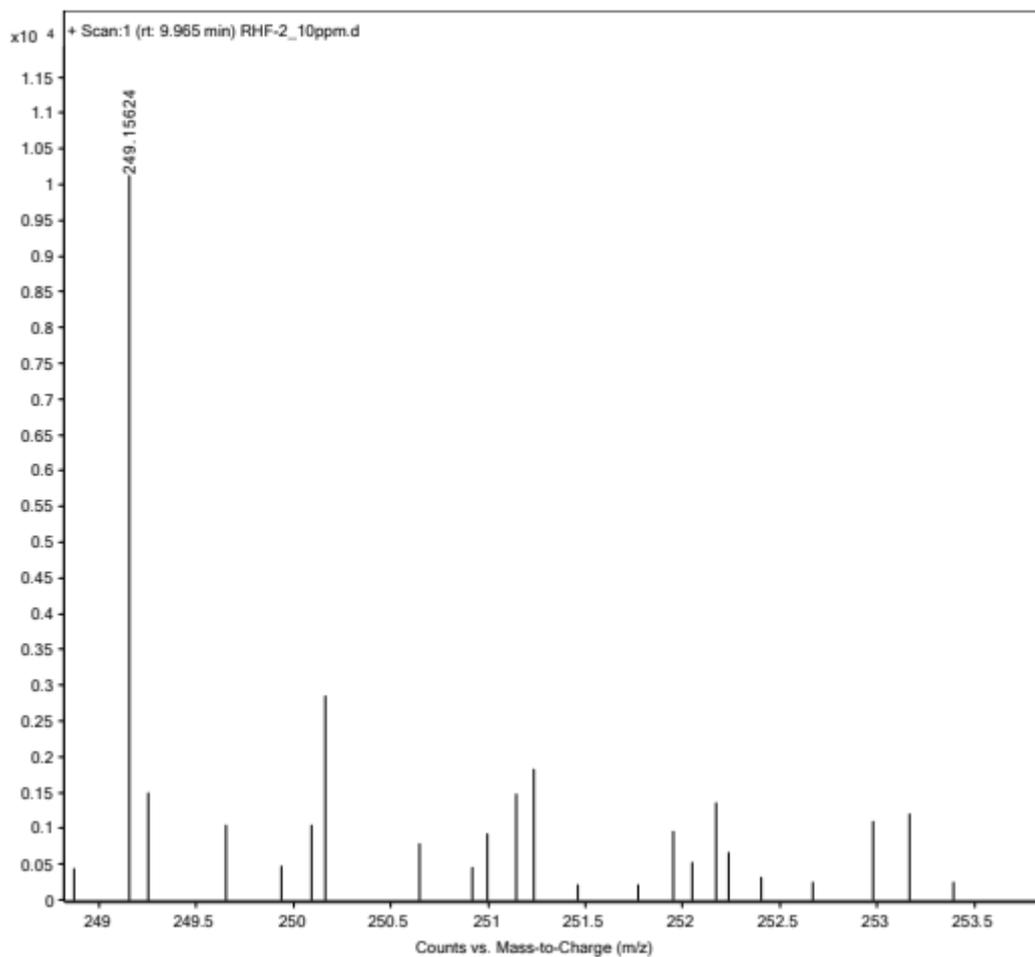


Figure 3.73: The HRESIMS of compound 109

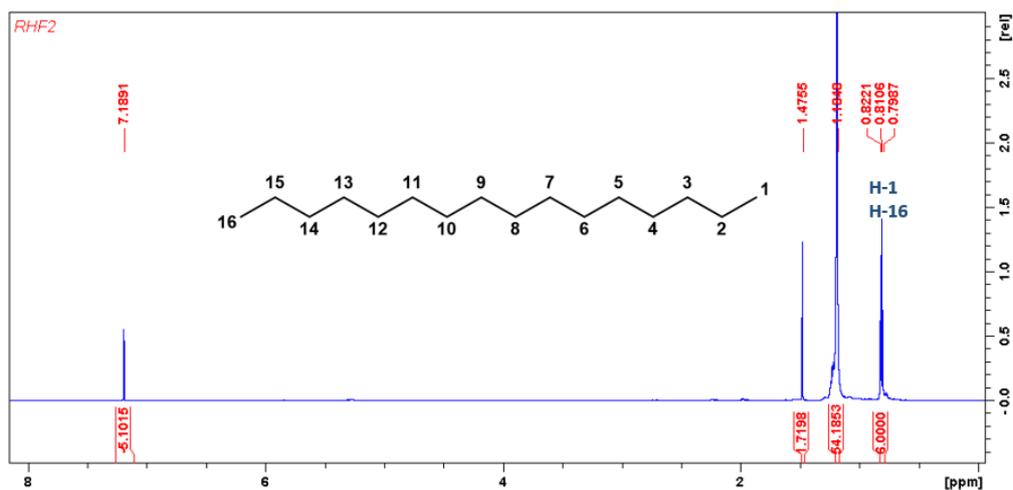


Figure 3.74: ¹H NMR (300 MHz, CDCl₃) spectrum of compound 109

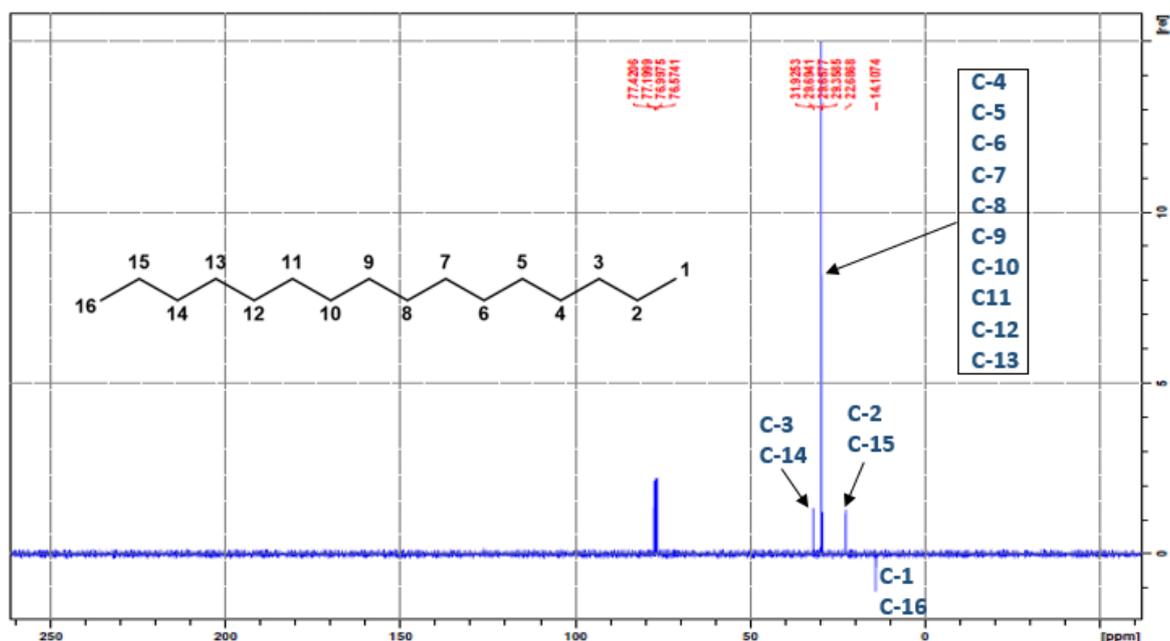


Figure 3.75: ^{13}C NMR (75 MHz, CD_3OD) spectrum of compound 109

3.1.10 Antimicrobial activity of the isolated compounds from *R. chalepensis*

The isolated compounds from *R. chalepensis* were screened to evaluate their antimicrobial activities and determine their minimum inhibition concentration (MIC) by serial dilution using microtitre assay. The antimicrobial method was performed on four bacteria species *S. aureus* (NCTC 12981), *E. coli* (NCTC 12241), *P. aeruginosa* (NCTC 12903), *M. luteus* (NCTC 7508) and one fungal strain *C. albicans* (ATCC 90028). The highest test concentration of compounds were 1mg/mL. The observed MIC values are outlined in Table 3.17.

The analyses of biological evaluation for the pure compounds revealed that all the tested compounds except compound 109 succeeded to constrain the growth of *C. albicans* and *M. luteus* with different concentrations. Out of fourteen compounds, only compounds 42 and 109 failed to prevent *S. aureus* growth, while five compounds inhibited the growth of *E. coli*. Moreover, five compounds (46, 47, 48, 101 and 107) presented significant inhibitory activities against *P. aeruginosa* with different MIC values. Four distinct compounds (46, 101, 103 and 107) inhibited all the tested strains, while there was no biological activity for compound 109 within the concentration range tested.

Different antibiotics, summarized in Table 3.3, were used as positive controls in the microtitre assay, these antibiotics have a wide inhibit any range for different bacterial strain such as: nalidixic acid, gentamycin, chloramphenicol and ciprofloxacin while, nystatin was used as a positive control to inhibit *C. albicans*. The nutrient broth, water and normal saline were used as negative controls in the antimicrobial tests.

Table 3.17: The MIC for isolated compounds from *R. chalepensis*

Compounds	All concentrations in (mg/mL)				
	<i>Staphylococcus aureus</i> (NCTC 12981)	<i>Escherichia coli</i> (NCTC 12241)	<i>Pseudomonas aeruginosa</i> (NCTC 12903)	<i>Micrococcus luteus</i> (NCTC 7508)	<i>Candida albicans</i> (ATCC 90028)
42	N/A	N/A	N/A	5x10 ⁻¹	5x10 ⁻¹
43	N/A	N/A	N/A	5x10 ⁻¹	5x10 ⁻¹
44	5x10 ⁻¹	N/A	N/A	6.25x10 ⁻²	5x10 ⁻¹
45	1.25x10 ⁻¹	5x10 ⁻¹	N/A	1.25x10 ⁻¹	1.25x10 ⁻¹
46	1.25x10 ⁻¹	1.25x10 ⁻¹	5x10 ⁻¹	2.5x10 ⁻¹	6.25x10 ⁻²
47	1.25x10 ⁻¹	N/A	5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹
48	2.5x10 ⁻¹	N/A	5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹
99	5x10 ⁻¹	N/A	N/A	5x10 ⁻¹	5x10 ⁻¹
101	2.5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹
103	2.5x10 ⁻¹	2.5x10 ⁻¹	2.5x10 ⁻¹	6.25x10 ⁻²	6.25x10 ⁻²
104	5x10 ⁻¹	N/A	N/A	2.5x10 ⁻¹	2.5x10 ⁻¹
105	5x10 ⁻¹	N/A	N/A	5x10 ⁻¹	5x10 ⁻¹
107	2.5x10 ⁻¹	5x10 ⁻¹	5x10 ⁻¹	1.25x10 ⁻¹	6.25x10 ⁻²
109	N/A	N/A	N/A	N/A	N/A

*N/A: No Activity

3.1.11 Discussion

The Rutaceae is one of the largest botanical families distributed worldwide, which including plants, herbs, shrubs and trees. In traditional medicine, the Rutaceae family is considered as an exhaustive pharmacy due to the successful indications for curing skin lesions, inflammatory diseases, intestines disorder, peptic ulcers, ear problems, respiratory infections, asthma and rheumatism and being a cardio tonic. In addition, the strong fragrance of the Rutaceae members led to the discovery of essential oils and led to use as a perfume and insect repellent. All these features attracted the researchers to discover and isolated

hundreds of compounds from this family (San Miguel, 2003a; Orwa *et al.*, 2008; Arbab *et al.*, 2012).

The phytochemical studies on the Rutaceae family were rich with many secondary metabolites such as alkaloids, coumarins, flavonoids, glycosides and tannins. The biological research confirmed the antimicrobial activity, antimalarial activity, anti-inflammatory effect and there is a positive significant result on treating cancer cell. The scientific reviews discovered the antioxidant effect of the Rutaceae members that led to using these materials as antioxidant agents in the cosmetics industry (Gunatilaka *et al.*, 1994; Viljoen *et al.*, 2006; Supabphol and Tangjitjareonkun, 2014)

The present study included one of the important members of the Rutaceae family, *Ruta chalepensis*. The current research screened the antimicrobial activity of different parts (fruit, stem, leaves and roots) of the Iraqi genus *R. chalepensis*. Microtitre assay was the biological assay used in this study and performed on two Gram-positive bacteria *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981); two Gram-negative *Escherichia coli* (NCTC 12241); *Pseudomonas aeruginosa* (NCTC 12903) and *Candida albicans* (ATCC 90028), which is considered one of the most common pathogenic fungi. The primarily screening of three extracts (*n*-hexane, DCM, MeOH) for every part (fruit, stem, leaves and roots) of *R. chalepensis* revealed that all extract successively inhibited the microbial strain with different MIC values and the most active extract was *n*-hexane fruits against *C. albicans* with MIC 1.56×10^{-2} mg/mL. The root extracts showed a weak effect with the maximum concentration tested which was 10 mg/mL.

Nineteen compounds were isolated from different plant parts and active fractions including eight alkaloids γ -fagarine (**42**), arborinine (**107**), graveoline (**105**), imperatorin (**104**), isokokusaginine (**100**), kokusaginine (**99**), ribalinium (**106**) and skimmianine (**41**). Five coumarins bergapten (**43**), chalepin (**45**), chalepensisin (**46**), isopimpinellin (**44**) and rutamarin (**47**); one sinapoyl glycoside, 3', 6-disinapoylsucrose (**108**) and one alkane hexadecane (**109**). Four glycosylated flavonoids rutin (**48**), rutin 3'-methyl ether (**101**), rutin

7,4' - dimethyl ether (**102**) and 6-Hydroxy-rutin 3' -7-dimethyl ether (**103**) which to the best of our knowledge, is a new natural product.

Fourteen of the isolated compounds were tested against the microbes used in this work. In general, all alkaloids showed significant antimicrobial activity. Previous studies have reported finding and isolating alkaloids from *Ruta* spp. Compound **107** successively inhibited all strains tested with high MIC value 1.25×10^{-1} , 6.25×10^{-2} mg/mL against *M. luteus* and *C. albicans*, respectively. Compound **107** is an acridone alkaloid with a wide spectrum of activity, which has a hydroxyl on position 1; methyl on position 10 and two oxygenated methyl groups on position 2 and 3. The high activity of compound **107** maybe belonging to the variety of substitution attached to this compound in addition for the presence of a nitrogen atom.

The coumarin compounds showed considerable antimicrobial activity and particularly compound **46** against *C. albicans* with MIC 6.25×10^{-2} mg/mL. The analysis of data of microbiological test led to realise that there was a structure-activity relationship, the coumarins with substituted 3,3-dimethylbut-1-ene (chalepin **45**, chalepensis **46**, rutamarin **47**) showed a remarkable inhibitory activity greater than other coumarins (bergapten **43**, isopimpinellin **44**). Among the most active three coumarins **45**, **46**, **47**, compound chalepensis **46** was the most active one.

The furanocoumarin chalepensis (**46**) occurs in many medicinal plants of the Rutaceae family, which have a wide range of pharmacological activities such as anti-inflammatory, antifertility, antiplatelet aggregation and anticancer activities (Lo *et al.*, 2012; Quintanilla-Licea *et al.*, 2014; Nakano *et al.*, 2017).

The hexadecane (**109**) was obtained after, the vacuum liquid chromatography (VLC) of the *n*-hexane leaf extracts. The biological experiment showed compound **109** did not have any inhibitory activity within the tested concentration range (1mg/ mL).

Flavonoids, one of the biggest class of natural products, play vital roles for plants and have many potential pharmaceutical proprieties. Flavonoids are associated with a number of claimed health benefits because of their free radical-scavenging properties. Research has shown the antimicrobial activity, anti-inflammatory activity, antioxidant effect, antimutagenic and anticarcinogenic activity for flavonoids (Rice-Evans *et al.*, 1996; Pietta, 2000; Cushnie and Lamb, 2005; Mabry *et al.*, 2012). Four glycosylated flavonoids **48**, **101**, **102**, **103** were isolated from different parts of *R. chalepensis*. Rutin (**48**), one of the most well known compounds, which has different biological activities, was the first flavonoid isolated on this work. Rutin consists of three benzene rings A, B, C connecting to a rutinosyl moiety. All the examined glycosylated flavonoid revealed significant inhibited activity against microbe tested practically compound **103** with MIC 6.25×10^{-2} mg/mL against *M. luteus* and *C. albicans*. The 1D and 2D spectra of compound **103** were similar to rutin (**48**) with additional two oxygenated methyl and extra hydroxyl on position 6 which identified it as new compound discovered for the first time. The primary testing to determine the mechanism of action for the tested compounds revealed all the tested compounds were bacteriostatic.

3.2 The Citrus

Citrus fruits are a major commodity in the world. *Citrus* species are used as traditional medicinal herbs in several countries. *Citrus* fruits are good sources of nutrition, mineral elements, and secondary metabolites (Davenport, 1990; Spiegel-Roy and Goldschmidt, 1996; Ladanyia and Ladaniya, 2010). This work includes a study of the phytochemicals and antimicrobial activity of *Citrus sinensis* (leaves and peels) and *Citrus grandis* (leaves) against *Escherichia coli* (NCTC 12241), *Pseudomonas aeruginosa* (NCTC 12903), *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981) and *Candida albicans* (ATCC 90028).

3.2.1 Extraction

The Soxhlet apparatus was utilized for extraction of leaves and peels of *Citrus sinensis* and leaves of *Citrus grandis* using three solvents consecutively, *n*-hexane, dichloromethane (DCM) and methanol (MeOH). The highest percentage yield of extraction for *Citrus* was 7.63% for methanolic leaves extract of *C. sinensis* (SLM) and the lowest extraction percentage was 0.42% for the peels methanolic extract of *C. sinensis* (PLM). The percentage yield of extracts summarized in Table 3.18

Table 3.18: Percentage yield of *Citrus*

<i>Citrus</i> species	<i>Citrus</i> parts	Weights (gm)	Extract type	% Yield
<i>Citrus sinensis</i>	Leaves	64.86	<i>n</i> -hexane (SLH)	0.60
			DCM (SLD)	0.75
			MeOH (SLM)	7.63
	Peels	131.13	<i>n</i> -hexane (PLH)	2.13
			DCM (PLD)	3.77
			MeOH (PLM)	0.42
<i>Citrus grandis</i>	Leaves	117.00	<i>n</i> -hexane (GLH)	1.69
			DCM (GLD)	1.15
			MeOH (GLM)	4.41

3.2.2 Preliminary analytical TLC screening

The first analysis for the development TLC plates of the *n*-hexane and DCM extracts of *Citrus* were performed following method **H** (Figure 3.75). The developed TLC plates were viewed under short (254 nm) and long (366 nm) were UV light followed by spraying with anisaldehyde reagent and then heating at 100°C for 5 min to reveal different coloured spots.

3.2.3 Analytical HPLC screening of the MeOH extracts

The three MeOH *Citrus* extracts (10 mg/mL) were subjected to analytical -HPLC. The leaf MeOH extract (Figure 3.76) of *C. sinensis* were successively analysed using method **Q** while, the method **L** was followed to analyze the MeOH extract of *C. sinensis* peel (Figure 3.77). The MeOH extract of *C. grandis* leaves were (Figure 3.78) loaded on to analytical HPLC using method **R**.

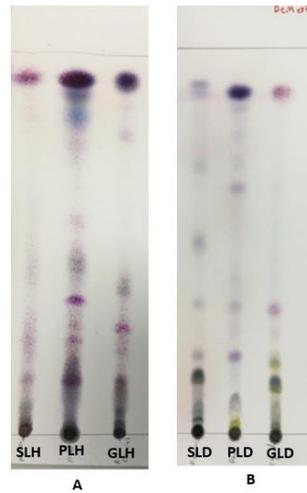


Figure 3.76: The TLC plates for Citrus extracts

A: *n*-hexane extracts B: DCM extracts

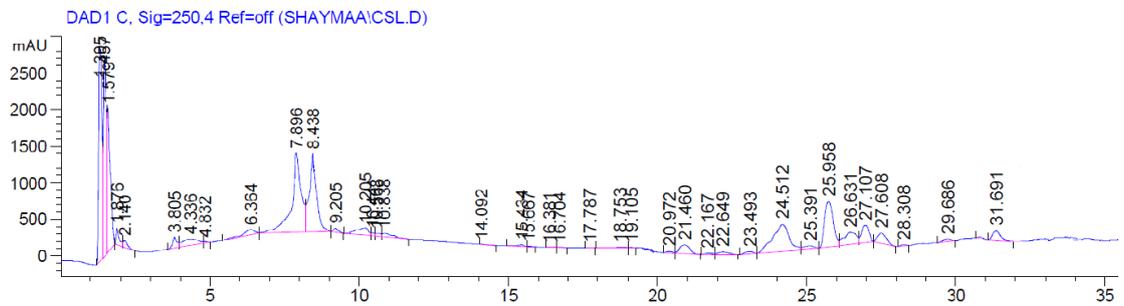


Figure 3.77: Analytical HPLC chromatogram of MeOH extract of *C. sinensis* leaves

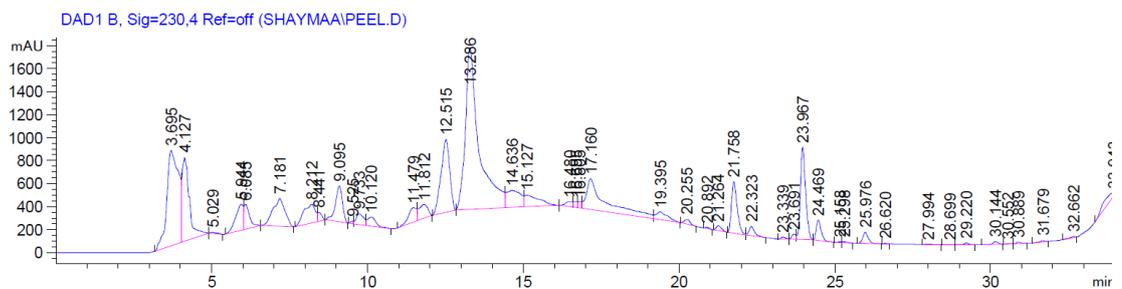


Figure 3.78: Analytical HPLC chromatogram of MeOH extract of *C. sinensis* peel

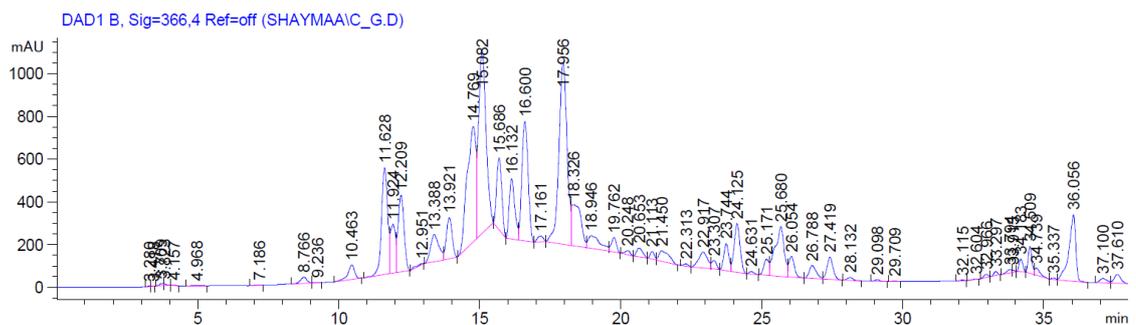


Figure 3.79: Analytical HPLC chromatogram of MeOH extract of *C. grandis*

3.2.4 The antimicrobial screening for *Citrus sinensis* and *Citrus grandis*

All the six extracts of *C. sinensis* and three extracts of *C. grandis* with maximum concentrate 10 mg/mL were investigated for their antimicrobial proprieties against *Escherichia coli* (NCTC 12241), *Pseudomonas aeruginosa* (NCTC 12903), *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981) and *Candida albicans* (ATCC 90028) using the modified microtitre plate assay (Sarker *et al.*, 2007). The findings (Table 3.19) suggested that the peel DCM extract had potent antimicrobial activity against *M. luteus* with a MIC value 4.88×10^{-3} mg/mL. In general, the three DCM extract inhibited all the microbes used in this study with MIC 6.25×10^{-1} mg/mL and less. The *n-hexane* extracts failed to present any antimicrobial effect within the maximum concentration (10 mg/mL) used in this study. Whereas, the peel MeOH extract showed moderate antimicrobial activity against *M. luteus* and *C. albicans* and the MIC values was 3.12×10^{-1} , 6.25×10^{-1} mg/mL, respectively.

Table 3.19: MIC values (mg/mL) of *Citrus* extracts

Bacteria and fungi	Extract	<i>Citrus sinensis</i>		<i>Citrus grandis</i>
		Leaves	Peels	Leaves
Gram-negative				
<i>Escherichia coli</i>	<i>n</i> -Hexane	2.5	5	5
	DCM	6.25×10^{-1}	3.12×10^{-1}	3.12×10^{-1}
	MeOH	2.5	1.25	1.25
<i>Pseudomonas aeruginosa</i>	<i>n</i> -Hexane	N/A	5	-
	DCM	5	6.25×10^{-1}	3.12×10^{-1}
	MeOH	5	1.25	5
Gram-positive				
<i>Micrococcus luteus</i>	<i>n</i> -Hexane	5	1.56×10^{-1}	6.25×10^{-1}
	DCM	3.12×10^{-1}	4.88×10^{-3}	3.125×10^{-1}
	MeOH	1.25	3.12×10^{-1}	6.25×10^{-1}
<i>Staphylococcus aureus</i>	<i>n</i> -Hexane	5	-	-
	DCM	6.25×10^{-1}	6.25×10^{-1}	6.25×10^{-1}
	MeOH	2.5	5	5
Pathogenic fungi				
<i>Candida albicans</i>	<i>n</i> -Hexane	N/A	2.5	5
	DCM	3.12×10^{-1}	1.56×10^{-1}	6.25×10^{-1}
	MeOH	5	6.25×10^{-1}	2.5

*N/A: No Activity

3.2.5 Chromatographic fractionation of the extracts

Based on the result of the primary screening of *Citrus* crude extracts and following bioassay guided the MIC value 6.25×10^{-1} mg/mL was selected as the minimum threshold concentration for the future work. Further fractionation for the most active crude extract involved using vacuum liquid chromatography fractionation (VLC) and solid phase extraction (SPE) as described in the experimental section.

3.2.5.1 Vacuum liquid chromatography fractionation (VLC)

The DCM extracts *Citrus* were subjected to VLC over silica gel using the method C and fractionated to eight fractions. Table 3.20 summarizes the yields of all fractions. The DCM fractions obtained were analysed by the TLC using method F (Figure 3.80).

3.2.5.2 Solid-phase extraction (SPE)

The MeOH extract of *C. sinensis* peels was fractionated by SPE using method D to obtain four fractions and their yield is summarized in Table 3.20.

3.2.6 Screening of *Citrus* fractions for antimicrobial activity

All the obtained fractions were screened for their antimicrobial activity against *Escherichia coli* (NCTC 12241), *Pseudomonas aeruginosa* (NCTC 12903), *Micrococcus luteus* (NCTC 7508), *Staphylococcus aureus* (NCTC 12981) and *Candida albicans* (ATCC 90028). The results are summarized in Table 3.21 and Table 3.22.

The highest activity of the DCM fractions in leaves of *C. sinensis* and *C. grandis* were shown by fraction 7 (F7). The DCM fractions of the *C. sinensis* peel showed a significant antimicrobial activity and specifically fraction 6 (F6) and fraction 7 (F7) against *M. luteus* and *C. albicans* with MIC 1.56×10^{-2} . In addition, fraction 3 (F3) of the methanolic extract of *C. sinensis* peel gave considerable activity against all microbial strains used in this study. Fraction 2 (F2) showed strong activity against *M. luteus* with MIC 6.25×10^{-1} . The phytochemical and antimicrobial work was continued with fractions which were found to be active.

Table 3.20: The yield of *Citrus* fractions

Plant extracts	Type of extract	Weight of extract (gm)	Fractions yield %							
			F1	F2	F3	F4	F5	F6	F7	F8
<i>C. sinensis</i> leaves	DCM	2.5142	22.63	3.28	38.41	23.30	7.07	3.97	4.86	9.19
<i>C. sinensis</i> peels	DCM	2.1655	4.42	2.31	8.08	15.09	6.30	6.86	19.71	15.64
	MeOH	1.8210	48.68	6.74	3.14	2.67	-	-	-	-
<i>C. grandis</i> Leaves	DCM	2.2044	7.34	6.35	45.86	10.24	4.94	2.54	4.81	5.34

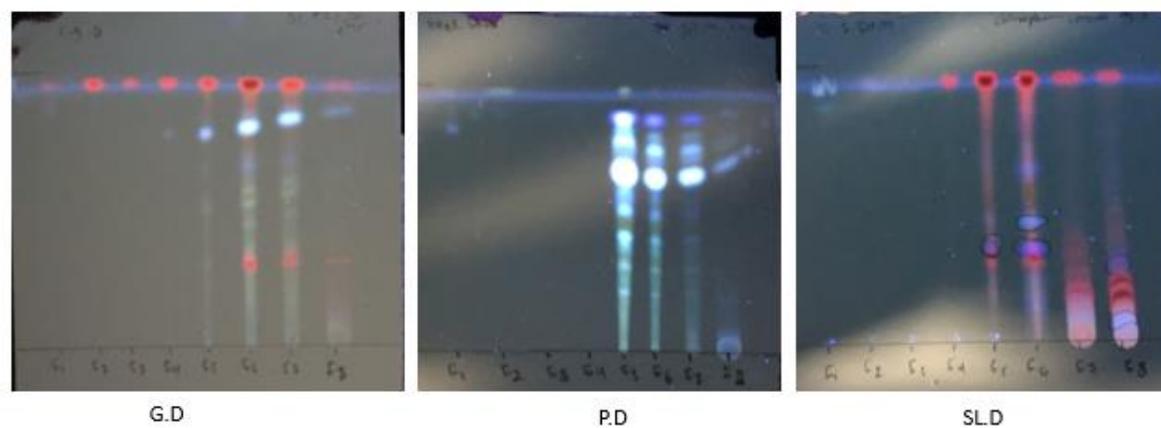


Figure 3.80: The analysis TLC for Citrus DCM fractions

G.D: *C. grandis* leaves; **P.D:** *C. sinensis* Peel; **SL.D:** *C. sinensis* leaves

Table 3.21: The MIC values (mg/mL) of DCM fractions for *Citrus*

Plant name and part	Extract type	Microbes	Fractions								
			F1	F2	F3	F4	F5	F6	F7	F8	
<i>C. sinensis</i> leaves	DCM	<i>E. coli</i>	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
		<i>P. aeruginosa</i>	N/A	N/A	N/A	N/A	N/A	1	1	N/A	
		<i>M. luteus</i>	N/A	N/A	N/A	N/A	N/A	N/A	6.25×10^{-2}	N/A	
		<i>S. aureus</i>	N/A	N/A	N/A	N/A	N/A	1	N/A	N/A	
		<i>C. albicans</i>	N/A	N/A	N/A	N/A	N/A	N/A	6.25×10^{-2}	1	
<i>C. sinensis</i> peels	DCM	<i>E. coli</i>	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	N/A	N/A	
		<i>P. aeruginosa</i>	N/A	N/A	N/A	N/A	1	5×10^{-1}	1	N/A	
		<i>M. luteus</i>	N/A	N/A	N/A	5×10^{-1}	5×10^{-1}	7.81×10^{-3}	1.56×10^{-2}	N/A	
		<i>S. aureus</i>	N/A	N/A	N/A	N/A	N/A	1	5×10^{-1}	N/A	
		<i>C. albicans</i>	N/A	N/A	N/A	5×10^{-1}	5×10^{-1}	7.81×10^{-3}	1.56×10^{-2}	N/A	
<i>C. grandis</i> leaves	DCM	<i>E. coli</i>	N/A	N/A	N/A	N/A	N/A	1	6.25×10^{-2}	N/A	
		<i>P. aeruginosa</i>	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	1.25×10^{-1}	N/A	
		<i>M. luteus</i>	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	1.56×10^{-2}	N/A	
		<i>S. aureus</i>	N/A	N/A	N/A	N/A	N/A	1	2.5×10^{-1}	N/A	
		<i>C. albicans</i>	N/A	N/A	N/A	N/A	N/A	5×10^{-1}	3.12×10^{-2}	N/A	

*N/A: No Activity

Table 3.22: The MIC (mg/mL) of MeOH extract of *C. sinensis* peels

Microbial name	Fractions			
	F1	F2	F3	F4
<i>E. coli</i>	N/A	1	2.5×10^{-1}	N/A
<i>P. aeruginosa</i>	N/A	N/A	5×10^{-1}	N/A
<i>M. luteus</i>	5×10^{-1}	6.25×10^{-1}	3.12×10^{-2}	5×10^{-1}
<i>S. aureus</i>	N/A	1	1.25×10^{-1}	N/A
<i>C. albicans</i>	5×10^{-1}	5×10^{-1}	1.56×10^{-2}	5×10^{-1}

*N/A: No Activity

3.2.7 Phytochemistry of *Citrus* species

The active fractions of DCM and methanolic extracts of *Citrus*, which exhibited a significant antimicrobial activity, were subjected to preparative-HPLC to separate the phytochemical constituents. The chromatographic separation and spectroscopic technique (NMR and HRESIMS) afforded isolation and characterization of fifteen compounds (Figure 3.81). The spectral data of known compounds were compared with the documented data. They were classified as hesperidin (**50**), rutin (**48**), narirutin (**51**), rutin 3`-methyl ether (**101**), 3-methoxynobiletin (**110**), nobiletin (**57**), sinensetin (**111**), 6,7,8,3',4'-pentamethoxyflavone (**112**), demethylnobiletin (**113**), 5-desmethylinensetin (**114**), cirsilineol (**115**), tangeritin (**116**), tetramethylscutellarein (**117**), salvigenin (**118**) and Marcitrus (**119**) .

All fourteen compounds isolated from the *Citrus* were identified as flavonoids. The literature depicts the flavonoids as one of the important medical natural product treatment material distributed in most the plant species, especially in the Rutaceae family (Havsteen, 2002). To the best of our knowledge, this is the first report on the phytochemical studies on leaves of the Iraqi genus *C. grandis* and compound **119** is reported here for the first time as a natural product.

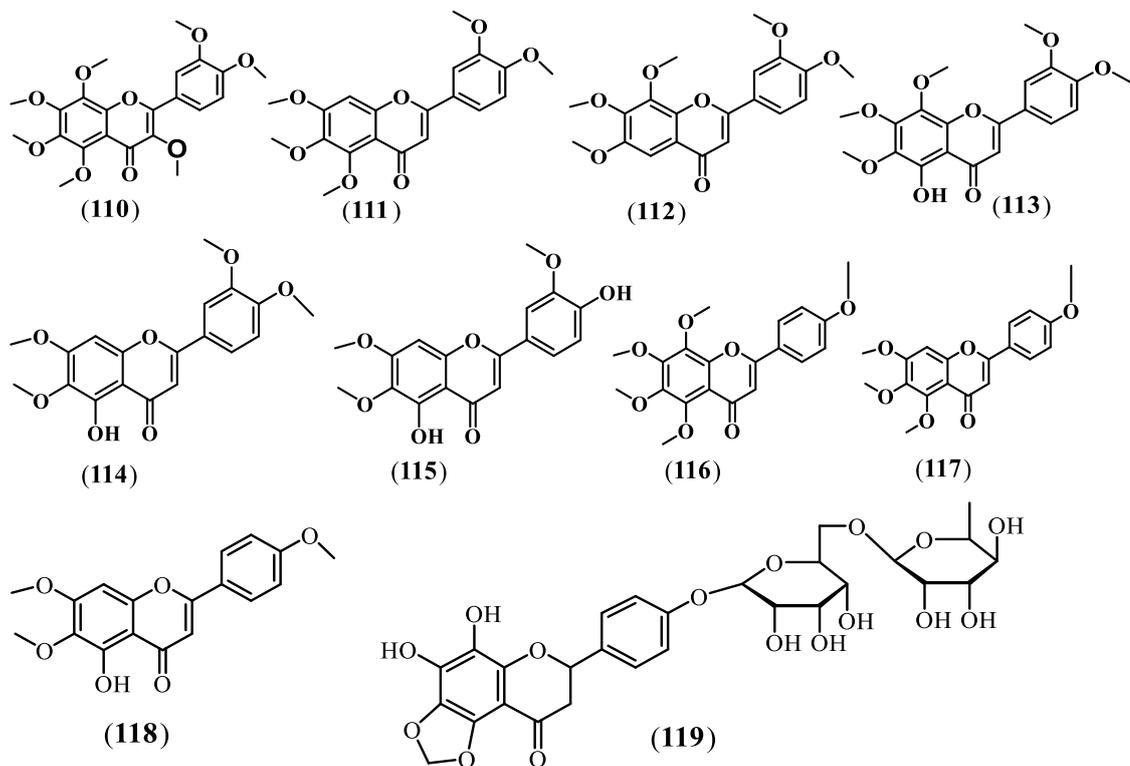


Figure 3.81: Isolated compounds from Citrus

3-methoxynobiletin (**110**), sinensetin (**111**), 6,7,8,3',4'-pentamethoxyflavone (**112**), demethylnobiletin (**113**), 5-desmethylinensetin (**114**), cirsilineol (**115**), tangeritin (**116**), tetramethylscutellarein (**117**), salvigenin (**118**) and marcitrus (**119**)

3.2.8 Isolation compounds from active fractions of Citrus

Based on the results of screening for the antimicrobial activity of the fractions, all the most active fractions were subjected to different chromatographic methods to isolate the active compounds.

3.2.8.1 Citrus grandis leaves DCM extract fraction 7

The fraction 7 of *C. grandis* leaves DCM extract was subjected to prep-HPLC (Agilent) using method T for 35 min. The volume of injection was 350 μ L and the flow rate was 10 mL/min (Figure 3.82). The separation process obtained six compounds, 6,7,8,3',4'-pentamethoxyflavone (**112**, 0.5 mg) (Harborne, 2013), sinensetin (**111**, 1.5 mg) (Han *et al.*, 2010), cirsilineol (**115**, 0.9 mg) (Hammoud *et al.*, 2012), nobiletin (**57**, 2.3 mg) (Li *et al.*, 2018), 5-desmethylinensetin (**114**, 0.7 mg) (Alarif *et al.*, 2013) and hesperidin (**50**, 1.0 mg) (Chiba *et al.*, 2003).

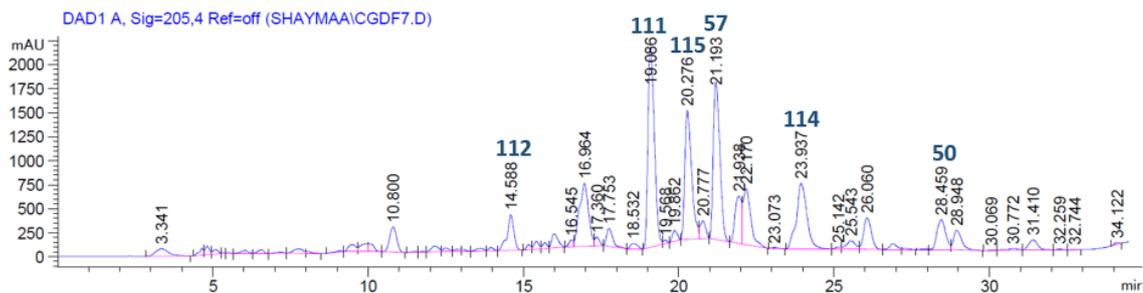


Figure 3.82: Preparative-HPLC chromatogram of isolated compounds from *Citrus grandis* leaves DCM extract fraction 7, using method **T**

3.2.8.2 *Citrus sinensis* leaves DCM extract fraction 7

The method **N** in prep-HPLC for 35 min was used to isolate the active compounds from fraction 7 of *C. sinensis* leaves (Figure 3.83). The injection volume was 250 μ L and the flow rate was 10 ml/min. The separation process afforded five compounds. They were identified as nobiletin (**57**, 1.0 mg) (Li *et al.*, 2018), 5-desmethylinensetin (**114**, 0.3 mg) (Alarif *et al.*, 2013), 6,7,8,3',4'-pentamethoxyl-flavone (**112**, 0.4 mg) (Harborne, 2013), cirsilineol (**115**, 0.3 mg) (Hammoud *et al.*, 2012) and hesperidin (**50**, 0.7 mg) (Chiba *et al.*, 2003).

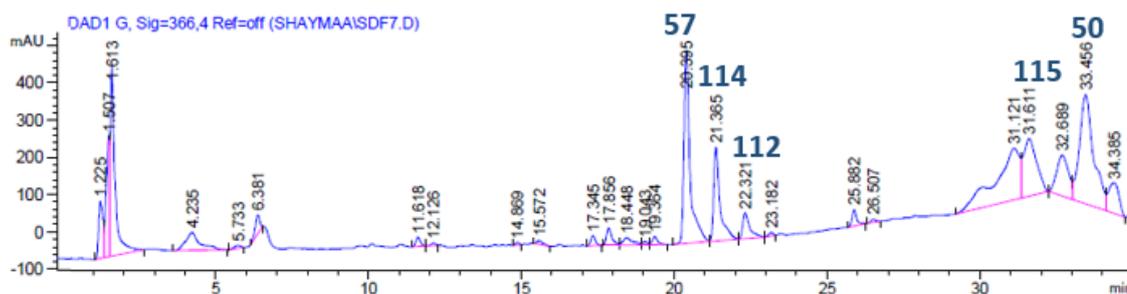


Figure 3.83: Preparative-HPLC chromatogram of isolated compounds from *Citrus sinensis* leaves DCM extract fraction 7, using method **N**

3.2.8.3 *Citrus sinensis* peels DCM extract fraction 6

The fraction 6 of the *C. sinensis* peels DCM extract was subjected to prep-HPLC (Agilent) following method **S** for 40 min. The volume of injection was 200 μ L and the flow rate was 8 mL/min (Figure 3.84). The isolation procedure produced six compounds,

sinensetin (**111**, 1.3 mg) (Han *et al.*, 2010), nobiletin (**57**, 1.9 mg) (Li *et al.*, 2018), tetramethylscutellarein (**117**, 1.0 mg) (Li *et al.*, 2018), 3-methoxynobiletin (**110**, 0.7 mg) (Li *et al.*, 2006), tangeritin (**116**, 0.7 mg) (Hamdan *et al.*, 2011) and demethylnobiletin (**113**, 0.4 mg) (Wang *et al.*, 2005).

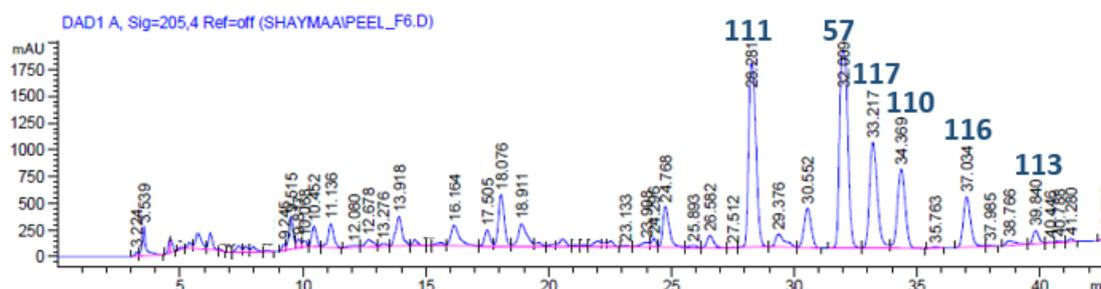


Figure 3.84: Preparative-HPLC chromatogram of isolated compounds from *Citrus sinensis* peel DCM extract fraction 6, using method **S**

3.2.8.4 *Citrus sinensis* peels DCM extract fraction 7

The VLC fraction 7 of the DCM extract of *C. sinensis* peels was separated by prep-HPLC (Agilent) using method **T** (Figure 3.85). The injection volume was 100 μ L and the flow rate was 8 mL/min. The separation process gave seven compounds, rutin (**48**, 0.3 mg) (Kamel *et al.*, 2014), narirutin (**51**, 0.4 mg) (Abu-Gharbieh and Shehab, 2017), sinensetin (**111**, 1.3 mg) (Han *et al.*, 2010), nobiletin (**57**, 1.8 mg) (Li *et al.*, 2018), tetramethylscutellarein (**117**, 1.1 mg) (Li *et al.*, 2018), 3-Methoxynobiletin (**110**, 0.8 mg) (Li *et al.*, 2006) and tangeritin (**116**, 0.4 mg) (Hamdan *et al.*, 2011).

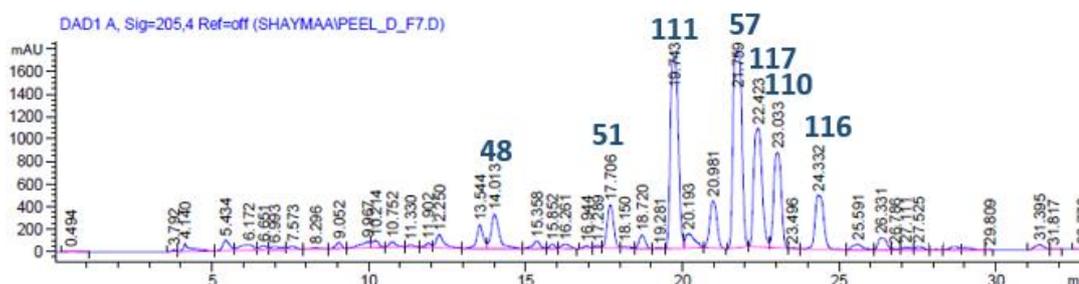


Figure 3.85: Preparative-HPLC chromatogram of isolated compounds from *Citrus sinensis* peel DCM extract fraction 7, using method **T**

3.2.8.5 *Citrus sinensis* peel MeOH extract fraction 2

The method **R** for 35 min on prep-HPLC was used to isolate the active compounds from fraction 2 of the *C. sinensis* peel MeOH extracts (Figure 3.86). The injection volume was 100 μ L and the flow rate was 10 mL/min. The separation process afforded five compounds identified as rutin (**48**, 0.3 mg) (Kamel *et al.*, 2014), Marcitrus (**119**) (0.5 mg), rutin 3`-methyl ether (**101**, 0.3 mg), narirutin (**51**, 1.2 mg) (Abu-Gharbieh and Shehab, 2017) and hesperidin (**50**, 2.3 mg) (Chiba *et al.*, 2003).

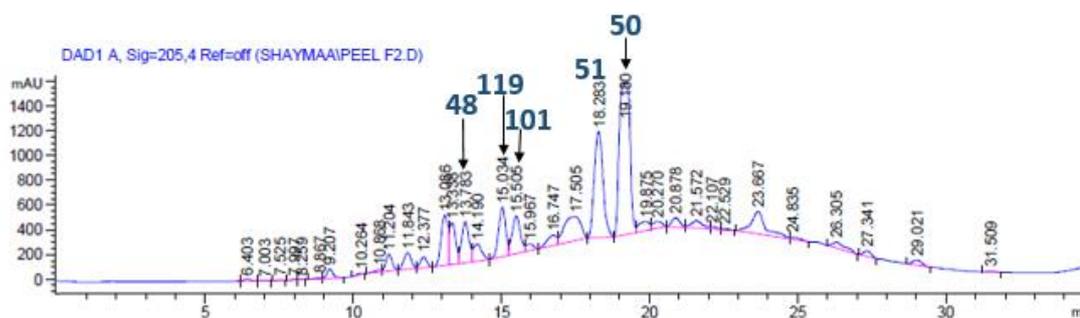


Figure 3.86: Preparative-HPLC chromatogram of isolated compounds from *Citrus sinensis* peel MeOH extract fraction 2, using method **R**

3.2.8.6 *Citrus sinensis* peels MeOH extract fraction 3

The method **R** for 35 min on prep-HPLC was used to isolate the active compounds from fraction 3 of the methanolic extract of *C. sinensis* peel (Figure 3.87). The injection volume was 200 μ L and the flow rate was 10 mL/min. The separation process revealed four compounds, characterized as rutin (**48**, 1.2 mg) (Kamel *et al.*, 2014), tangeritin (**116**, 0.9 mg) (Hamdan *et al.*, 2011), 3-methoxynobiletin (**110**, 2.1 mg), (Li *et al.*, 2006) and nobiletin (**57**, 0.5 mg) (Li *et al.*, 2018).

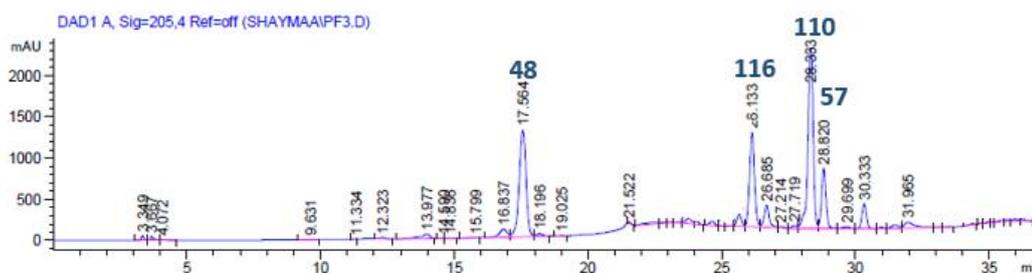


Figure 3.87: Preparative-HPLC chromatogram of isolated compounds from *Citrus sinensis* peel MeOH extract fraction 3, using method **R**

3.2.9 Characterisation and structure elucidation of isolated compounds.

3.2.9.1 Structure elucidation of 3-methoxynobiletin (110)

The compound **110** was isolated as dark yellow needles. The HRESIMS (Figure 3.88) suggested the empirical formula as $C_{22}H_{24}O_9$ and in the positive ion mode it showed peak $[M+H]^+$ at m/z 433.1577 (calculated 432.142035). The 1H NMR spectrum (Figure 3.89, Table 3.23) exhibited three aromatic protons as ABX pattern resonating at δ_H 7.01 (d, $J=8.52$ Hz), 7.84 (dd, $J= 8.7, 2.1$ Hz), 7.80 (d, $J= 2.1$ Hz) and seven methoxy groups at δ_H 3.88 (3H), 3.94 (3H), 3.96 (3H), 4.00 (3H) and 4.09 (3H). The ^{13}C NMR spectrum (Figure 3.90, Table 3.23) showed a total of twenty two carbons including a carbonyl at δ_C 174.3 (C-4), seven methoxyl carbons, three aromatic methines and the remaining as quaternary carbons. In the HMBC experiment (Figure 3.91), 3J correlations were observed from methoxyl at δ_H 3.88, 3.94, 3.96, 4.00 and 4.09 to the oxygenated quaternary carbons at 141.1 (C-3), 144.2 (C-6), 151.4 (C-4'), 138.6 (C-8) and 151.7 (C-7) respectively, while the methoxyl at δ_H 3.97 (2X 3H) revealed 3J correlation to oxygen bearing quaternary carbons at δ_C 148.6 (C-5) and 149.1 (C-3'). Moreover, H-6' showed long range correlation to δ_C 111.3 (C-2') and 151.4 (C-4') and the H-5 exhibited 3J correlation to δ_C 149.1 (C-3') and 123.8 (C-1'). The NMR spectroscopic and mass spectrometric data as well as the correlation revealed in the HSQC and HMBC confirmed the identification of compound **110** as 3,5,6,7,8,3',4'-heptamethoxyflavone, a poly methoxy flavonoid. The spectroscopic data of compound **110** were in a good agreement with respective published data of 3,5,6,7,8,3',4'-heptamethoxyflavone (Li *et al.*, 2014b; Owis, 2019) isolated from *C. sinensis* peel (Li *et al.*, 2006).

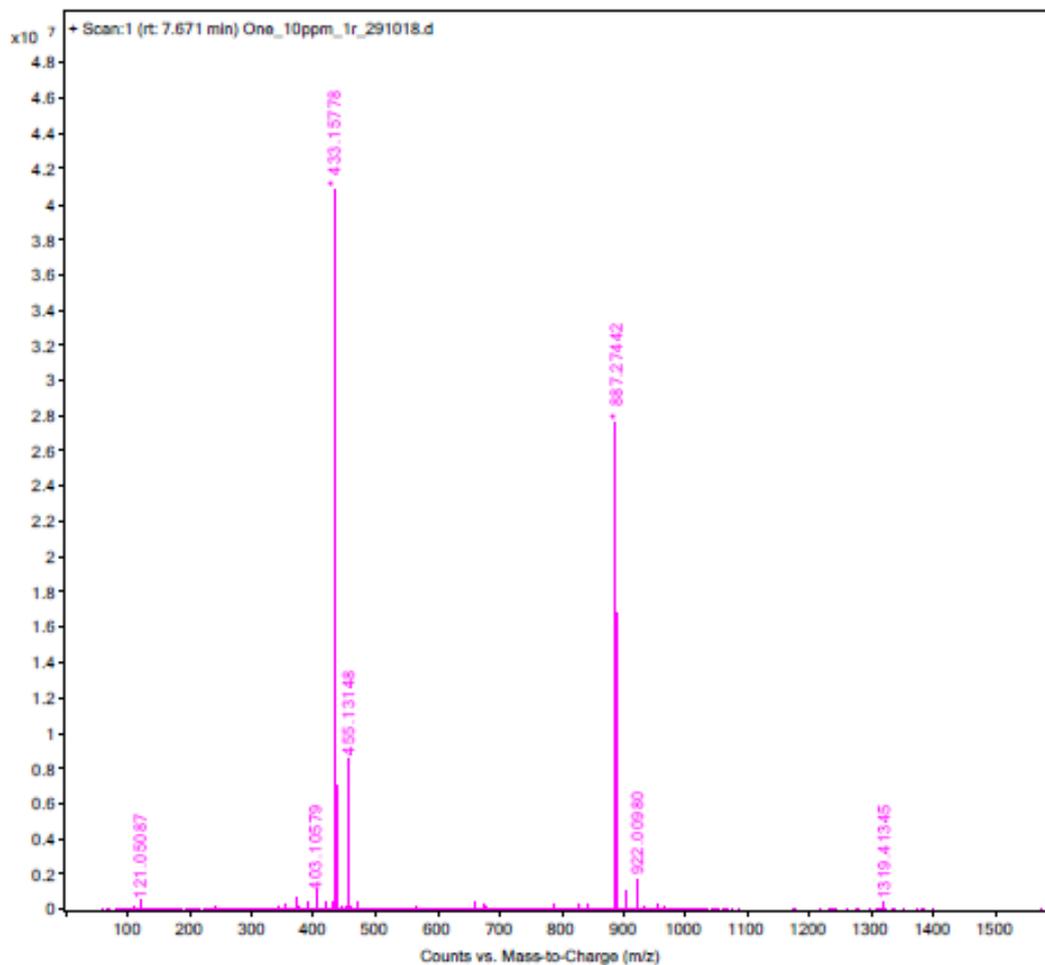


Figure 3.88: The HRESIMS spectrum of compound 110

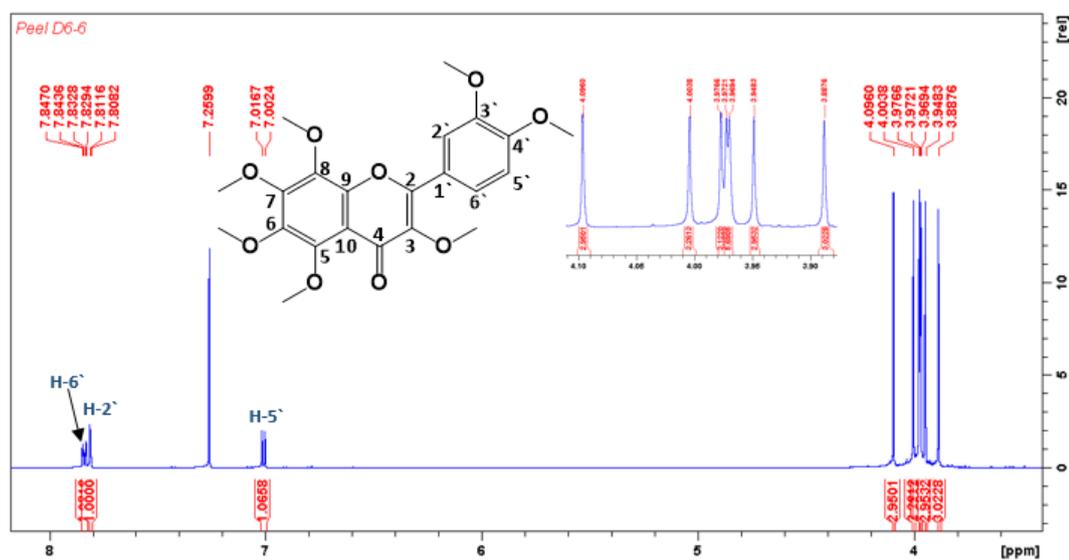


Figure 3.89: ¹H NMR (600 MHz, CDCl₃) spectrum of compound 110

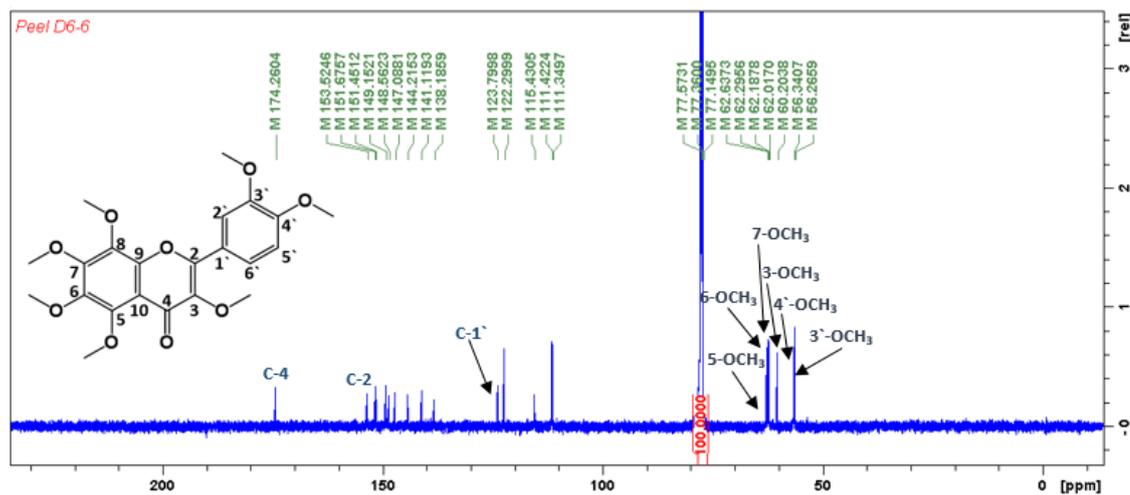


Figure 3.90: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **110**

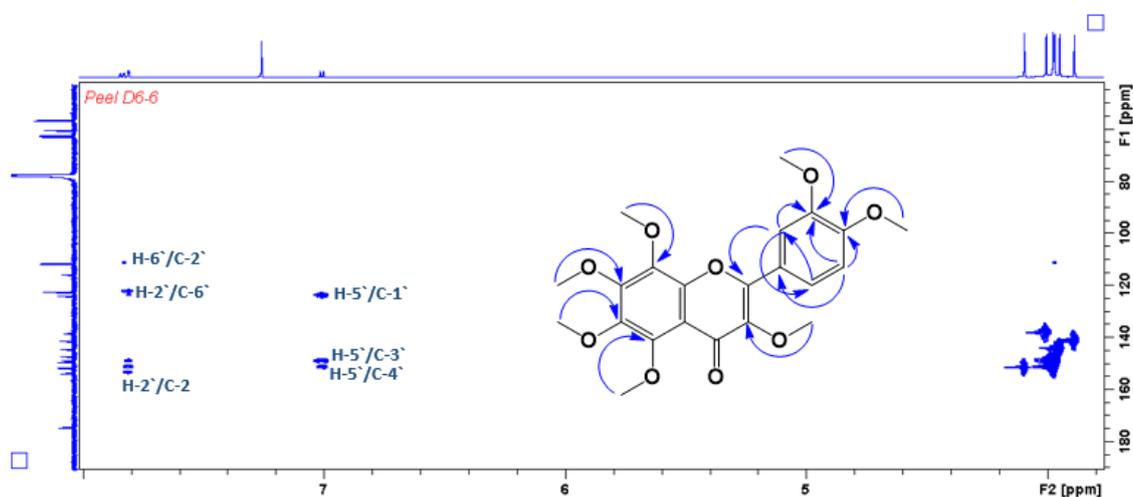


Figure 3.91: The HMBC correlation of compound **110**

3.2.9.2 Structure elucidation of nobiletin (**57**)

The compound **57** was isolated as a sharp white needle crystal. The HRESIMS (Figure 3.92) suggested the empirical formula as $\text{C}_{21}\text{H}_{22}\text{O}_8$ and in positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 403.1388 (calculated 403.1392). The 1D and 2D NMR spectra of compound **57** were similar to compound **110**. The main differences were the presence of an additional olefinic proton at δ_{H} 6.64 (H-3) and six methoxyl signals instead of seven in the ^1H NMR spectrum (Figure 3.93, Table 3.23). The DEPTQ NMR spectrum (Figure 3.94, Table 3.23) exhibited a total of twenty-one carbons including one carbonyl group at δ_{C} 177.8, six methoxyls, four methines and eleven quaternary carbons. In the HMBC experiment (Figure 3.95), long range correlations were observed from the olefinic proton at δ_{H} 6.64 (H-

3) to C-2 (δ_C 161.4, 2J), C-4 (δ_C 177.7, 2J), C-10 (δ_C 115.1, 3J) and C-1' (δ_C 124.2, 3J) which confirmed its position through C-3 in flavonoid molecule. Moreover, the methoxyl signals at δ_H 4.10 (3H), 4.02 (3H), 3.96 (H) and 3.97 (3H) obtained 3J correlation to carbons δ_C 152.2 (C-7), 138.1 (C-8), 149.1 (C-3') and 151.4 (C-4') while the methoxyl at δ_H 3.95 (2X 3H) revealed long-range correlation to carbons δ_C 149.6 (C-5) and 144.4 (C-6). Thus, compound **57** identified as 5,6,7,8,3',4'-hexamethoxyflavone or nobiletin. The spectroscopic data of compound **57** were in a good agreement with respective published data of nobiletin (Li *et al.*, 2018). Compound **57** has been reported to reduce skin and urinary inflammation (Murakami *et al.*, 2000; Li *et al.*, 2007) and it has an antidepressant effect (Wu *et al.*, 2015).

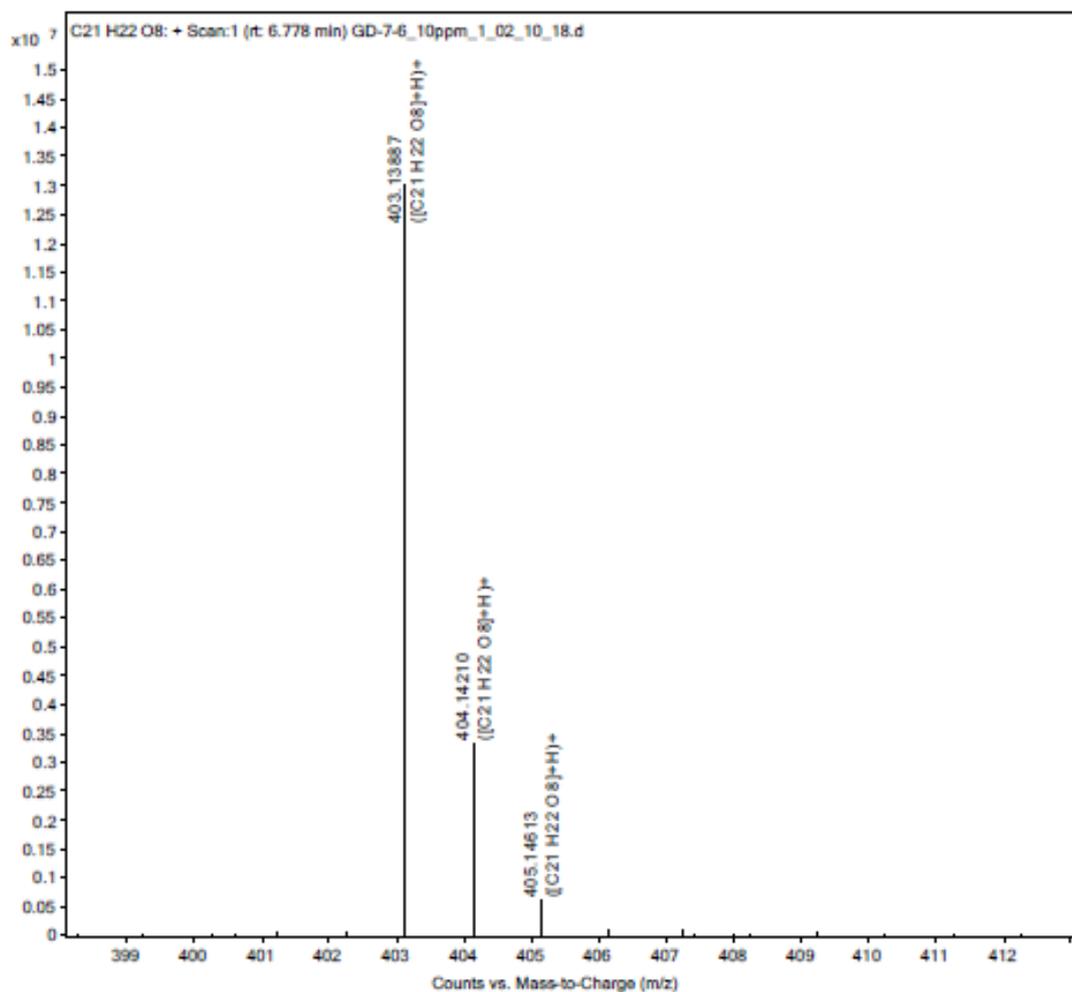


Figure 3.92: The HRESIMS spectrum of compound **57**

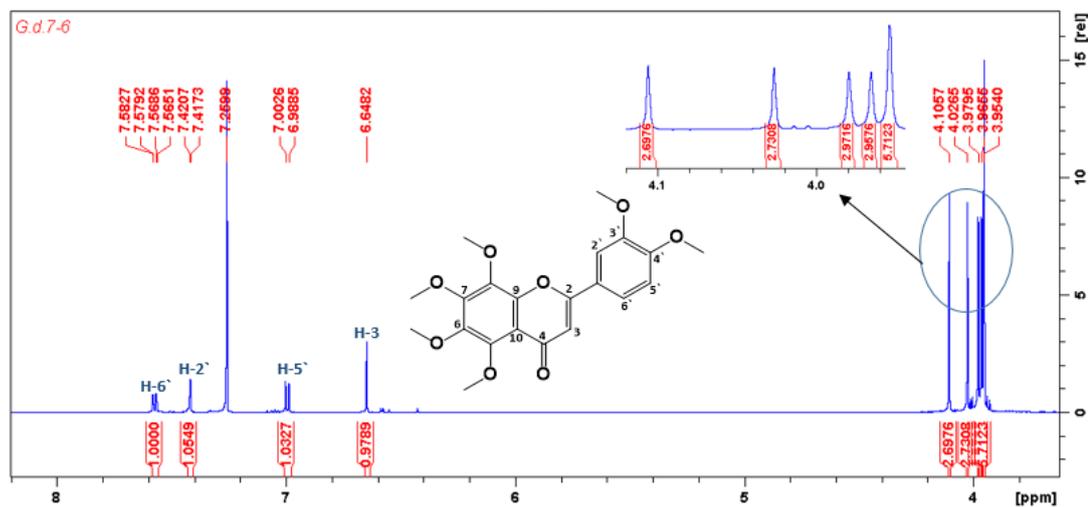


Figure 3.93: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 57

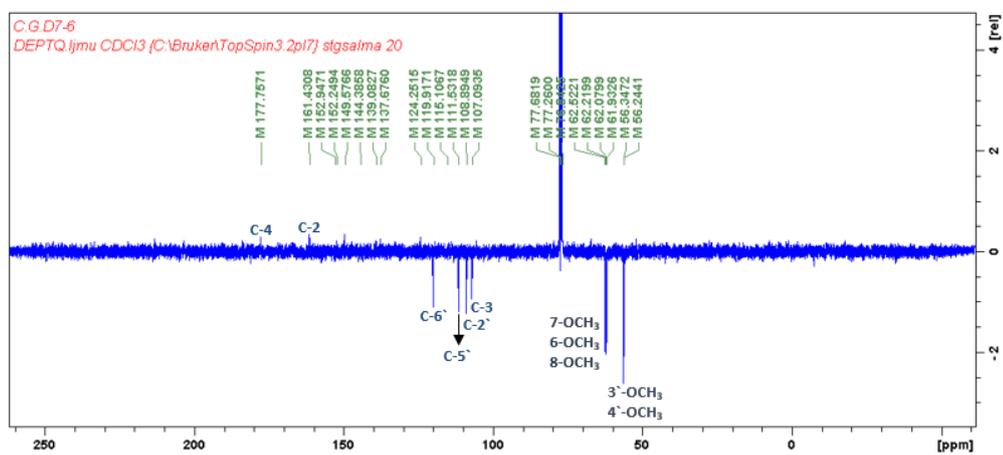


Figure 3.94: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound 57

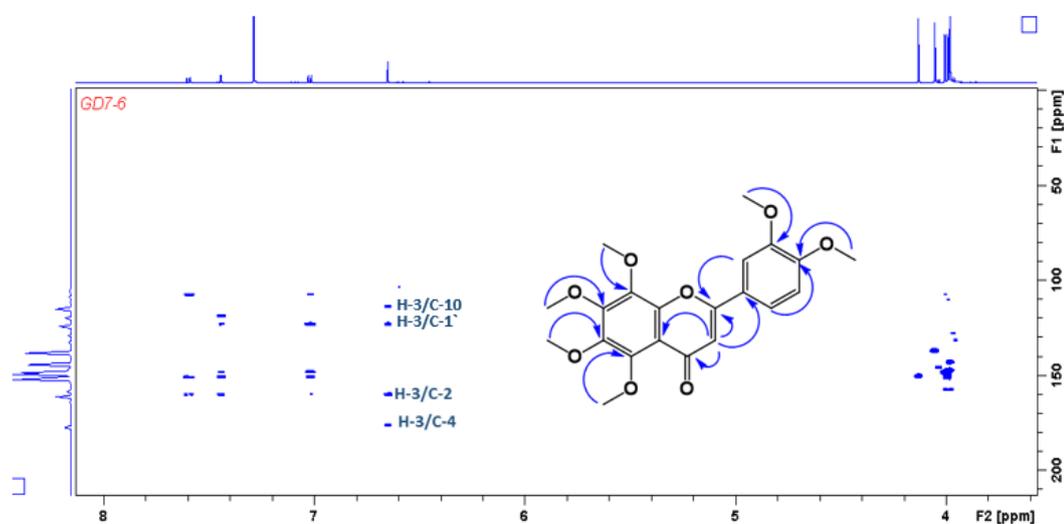


Figure 3.95: The HMBC correlation compound 57

3.2.9.3 Structure elucidation of sinensetin (111)

The compound **111** was isolated as a yellow powder. The HRESIMS (Figure 3.96) suggested the empirical formula as $C_{20}H_{20}O_7$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 373.1284 (calculated 373.1287). The 1D and 2D NMR spectra of compound **111** were very similar to compound **57**. The main differences were the presence of one extra aromatic methine proton at δ_H 6.60 and 6.79 and the presence of five methoxyl signals instead of six in the 1H NMR spectrum (Figure 3.97, Table 3.23). The ^{13}C NMR spectrum (Figure 3.98, Table 3.23) revealed a total of twenty carbons five methoxyl groups, five methine carbons and ten quaternary carbons including a deshielded carbonyl at δ_C 177.70 (C-4). In the HMBC experiment (Figure 3.99), H-3 showed long range correlations to carbons at 161.7 (C-2, 2J), 177.7 (C-4, 2J), 115.1 (C-10, 3J), 124.2 (C-1', 3J) while the aromatic methine proton at 6.79 (H-8) revealed 3J correlations 115.1 (C-10), 140.79 (C-6) and 161.7 (C-2). These correlations confirmed the positions of two olefinic protons in the flavonoid molecule. Moreover, the methoxyl signals at δ_H 3.92 (3H), 3.96 (3H) and δ_C 3.98 (3H) obtained 3J correlation to carbons 152.3 (C-5), 158.1 (C-7) and 149.7 (C-3') while the methoxyl at 3.99 (2 x 3H) revealed long range correlations to carbons 140.8 (C-6) and 152.9 (C-4'). Thus, compound **111** identified as 5,6,7,3',4'-pentamethoxyflavone. The data of compound **111** were in a good agreement with respective published data of sinensetin (Han *et al.*, 2010). Compound **111** has been reported to possess antioxidant (Akowuah *et al.*, 2004), antidepressant activities (Wu *et al.*, 2015) and worked as an angiogenesis agent (Lam *et al.*, 2012).

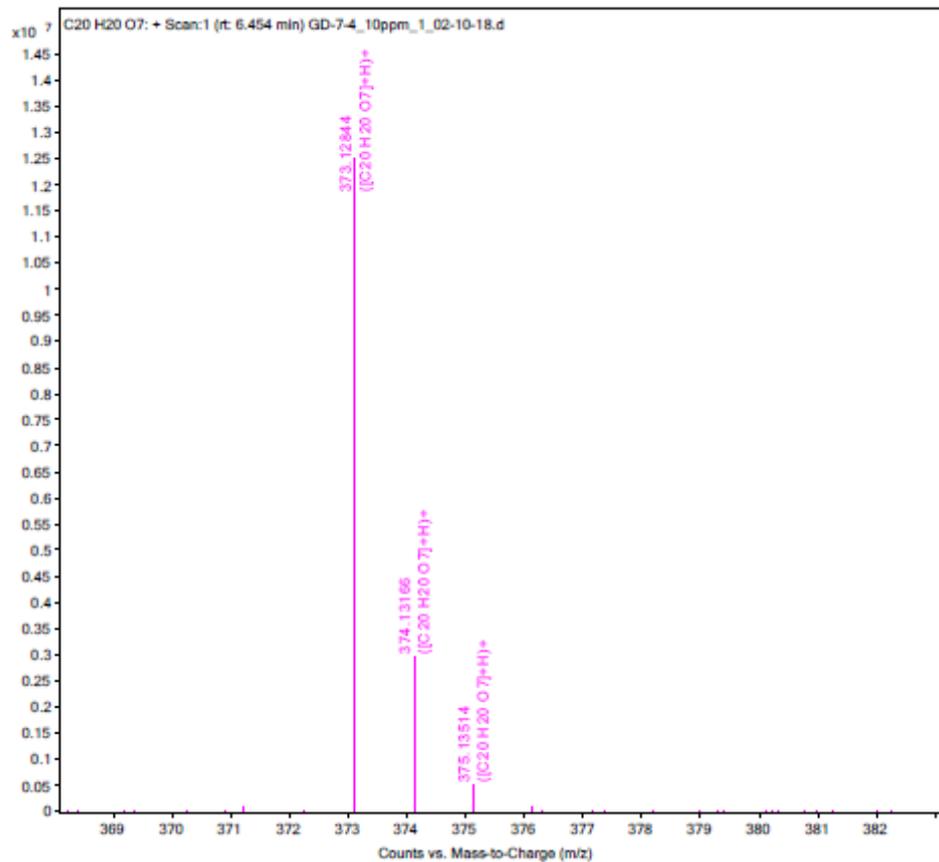


Figure 3.96: The HRESIMS of compound 111

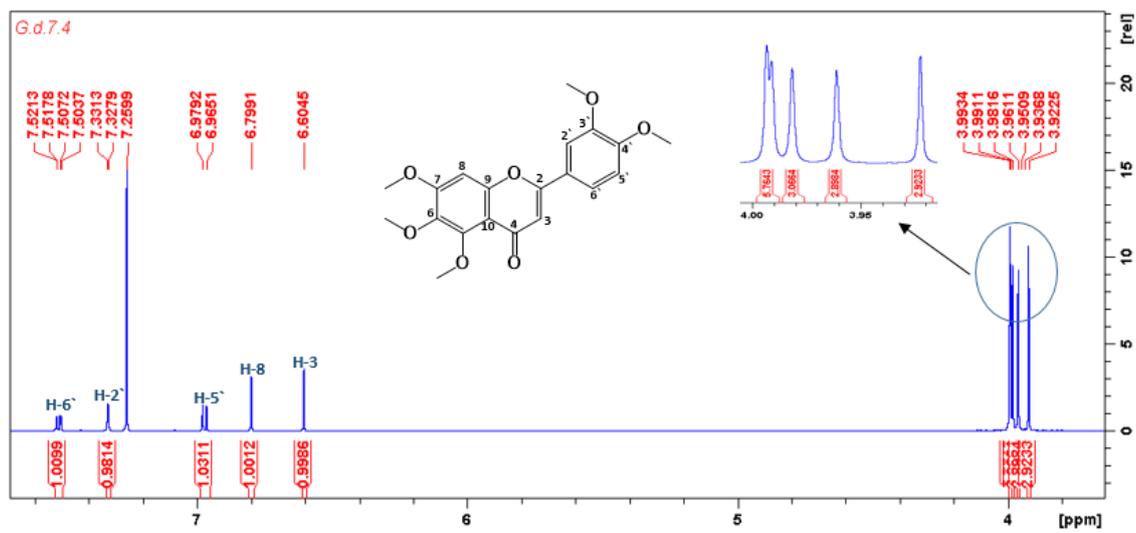


Figure 3.97: ¹H NMR (600 MHz, CDCl₃) of compound 111

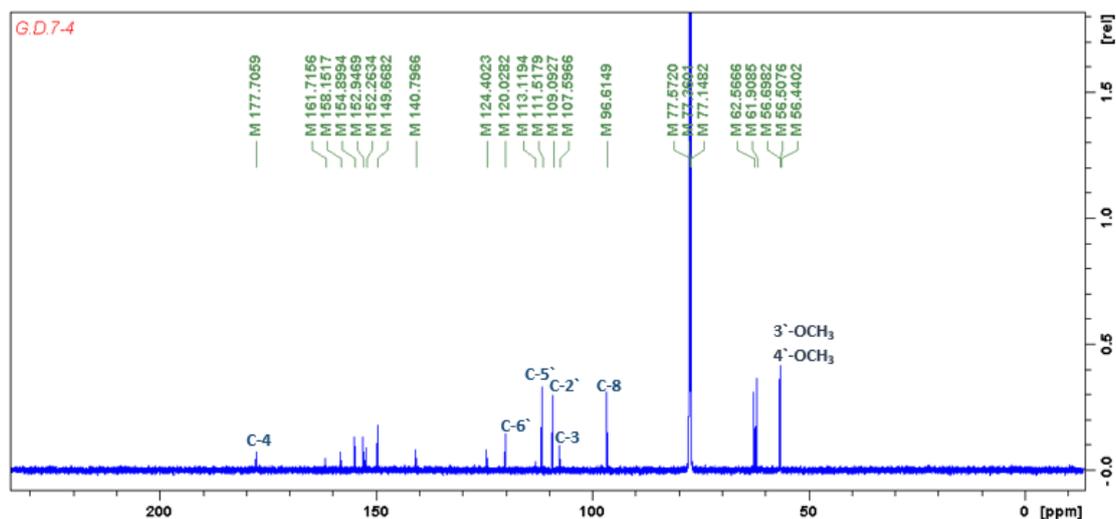


Figure 3.98: ^{13}C NMR (150 MHz, CDCl_3) of compound **111**

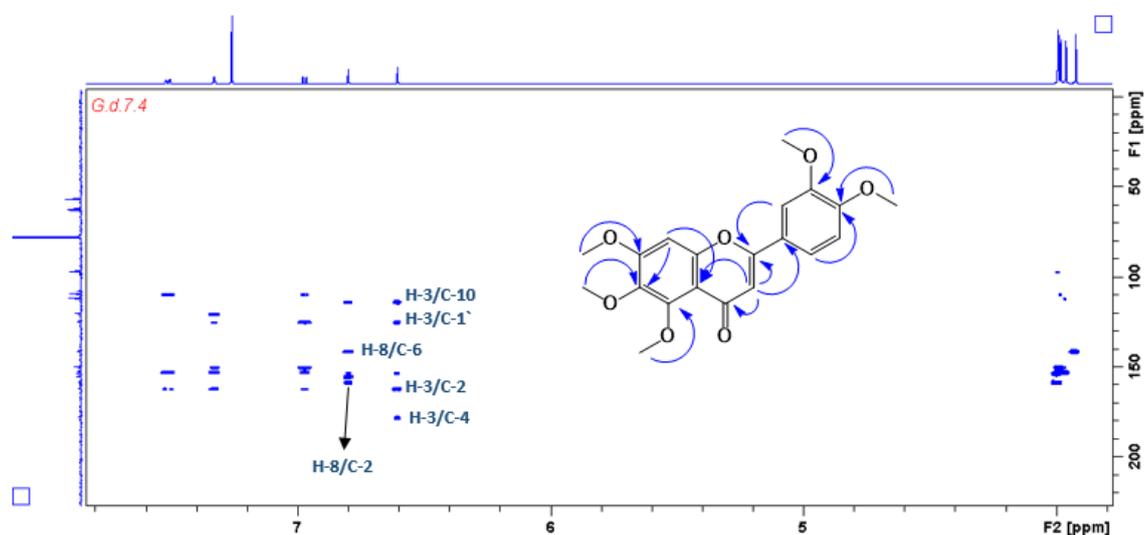


Figure 3.99: The HMBC correlation compound **111**

3.2.9.4 Structure elucidation of 6, 7, 8, 3',4'-pentamethoxyflavone (**112**)

The compound **112** was isolated as deep yellow crystals. The HRESIMS (Figure 3.100) suggested the empirical formula as $\text{C}_{20}\text{H}_{20}\text{O}_7$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 373.1317 (calculated 373.1287). The ^1H NMR spectrum (Figure 3.101, Table 3.23), ^{13}C NMR spectrum (Figure 3.102, Table 3.23) and 2D experiments of compound **112** were identical to compound **111**. The only difference was the position of an aromatic methine at δ_{H} 6.80 in the ^1H NMR spectrum. In the HMBC experiment (Figure 3.103), the methoxyl signals at δ_{H} 3.92 (3H), 3.96 (3H) and 3.98 (3H) obtained 3J correlation to carbons δ_{C} 140.7 (C-8), 153.0 (C-3') and 149.7 (C-4') while the methoxyl at δ_{H} 3.99 (2X

³H) revealed long range correlations to carbons at δ_C 125.2 (C-6) and 158.0 (C-7). Moreover, the aromatic proton at δ_H 6.80 showed long range correlations to δ_C 177.6 (C-4, ³J), 158.0 (C-7, ³J), 143.0 (C-9, ³J) and 113.2 (C-10, ²J) and confirmed its identity as H-5 in a flavonoid molecule. Thus, compound **112** was identified as pentamethoxy flavone. The data of compound **112** were in good agreement with respective published data for 6,7,8,3',4'-pentamethoxyflavone (Han *et al.*, 2010).

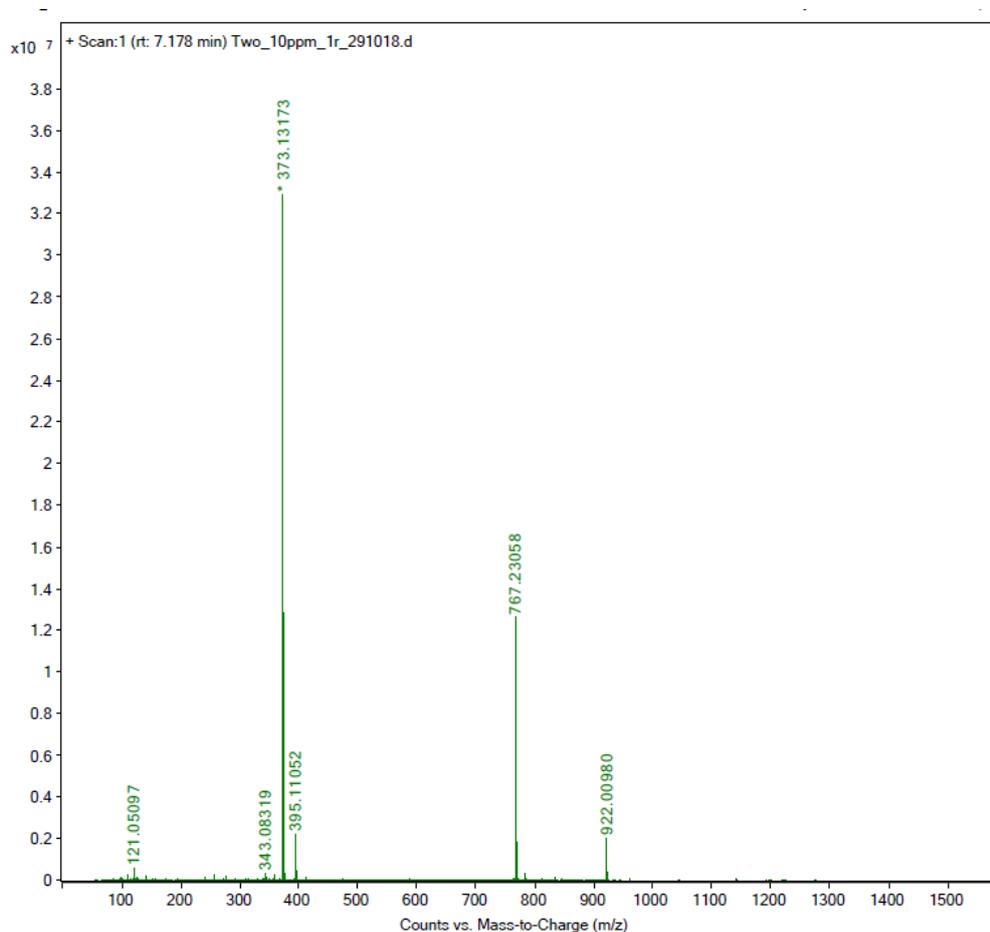


Figure 3.100: The HRESIMS spectrum of compound **112**

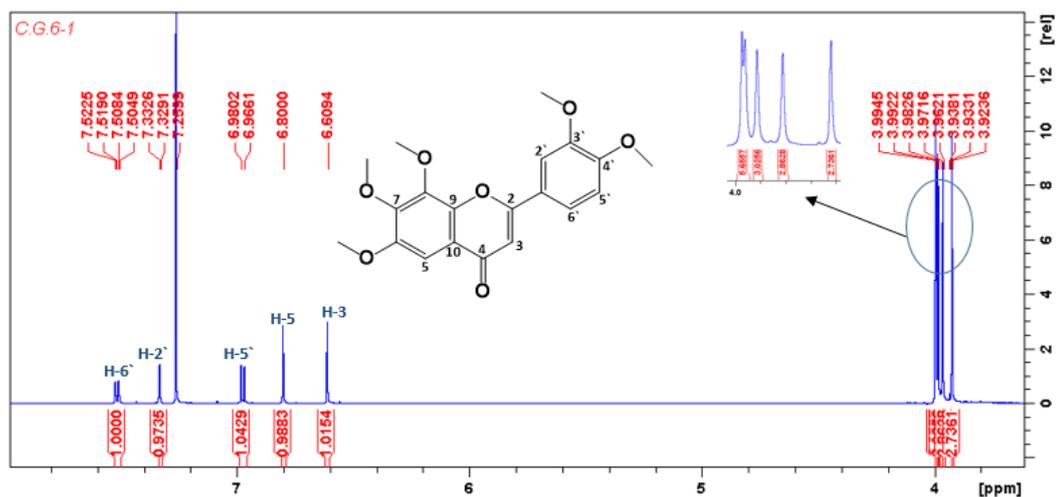


Figure 3.101: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 112

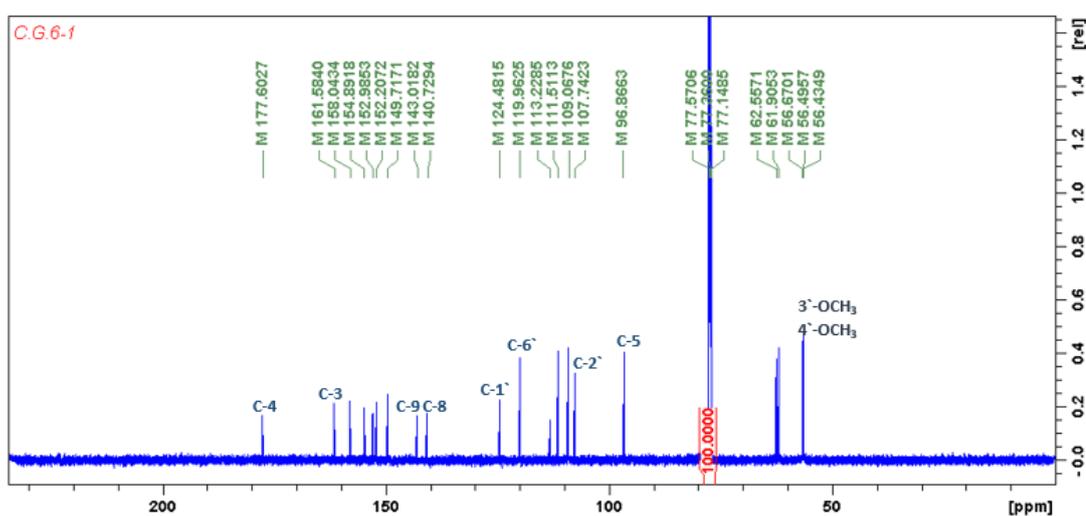


Figure 3.102: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound 112

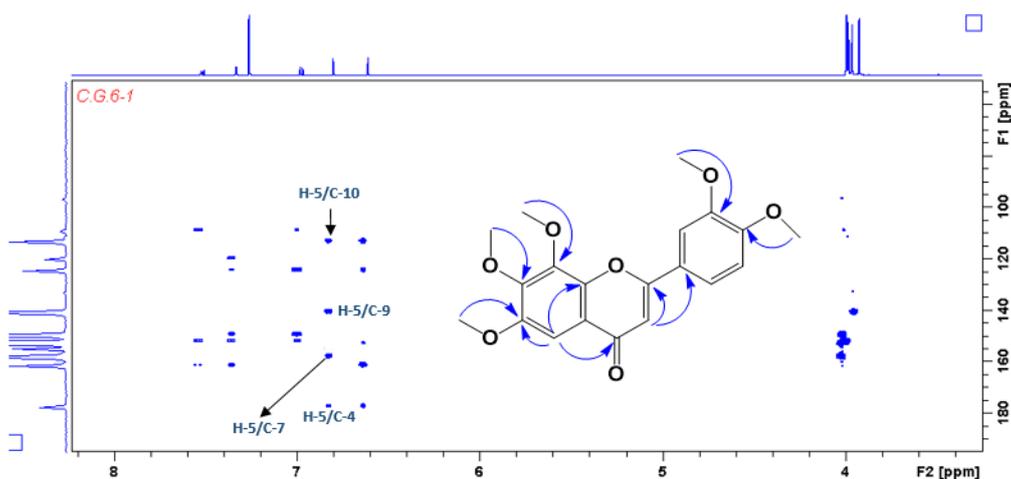


Figure 3.103: The HMBC correlation of compound 112

Table 3.23: ¹H NMR (600 MHz) and ¹³C NMR data of compounds **57**, **110**, **111** and **112**

Position	Chemical Shift δ_{H} (ppm), <i>J</i> in Hz				Chemical Shift δ_{C} (ppm), <i>J</i> in Hz			
	110	57	111	112	110	57	111	112
2	-	-	-	-	153.5	161.4	161.7	161.6
3	-	6.64 s	6.60 s	6.60 s	141.1	107.1	107.6	107.7
4	-	-	-	-	174.3	177.7	177.7	177.6
5	-	-	-	6.97 s	148.6	149.6	152.3	96.9
6	-	-	-	-	144.2	144.4	140.8	125.2
7	-	-	-	-	151.7	152.2	158.1	158.0
8	-	-	6.97,s	-	138,6	138.1	96.6	140.7
9	-	-	-	-	147.1	148.7	154.9	143.0
10	-	-	-	-	115.4	115.1	113.1	113.2
1'	-	-	-	-	123.8	124.2	124.4	124.5
2'	7.81 <i>d</i> (2.04)	7.41 <i>d</i> (2.1)	7.33 <i>d</i> (2.1)	7.33 <i>d</i> (2.1)	111.3	108.9	109.1	109.1
3'	-	-	-	-	149.1	149.6	149.7	153.0
4'	-	-	-	-	151.4	152.9	152.9	149.7
5'	7.01 <i>d</i> (8.52)	7.00 <i>d</i> (8.52)	6.96 <i>d</i> (8.4)	6.96 <i>d</i> (8.4)	111.4	111.5	111.5	111.5
6'	7.83 <i>dd</i> (2.04, 8.52)	7.57 <i>dd</i> (8.52, 2.1)	7.51 <i>dd</i> (8.4, 2.1)	7.51 <i>dd</i> (8.4, 2.1)	122.3	119.9	120.0	120.0
3-OCH ₃	3.88 s	-	-	-	60.2	-	-	-
5-OCH ₃	3.97 s	3.95 s	3.92 s	-	62.6	62.5	61.9	-
6-OCH ₃	3.94 s	3.95 s	3.99 s	3.99 s	62.2	62.2	62.6	62.5
7-OCH ₃	4.09 s	4.10 s	3.96 s	3.99 s	62.0	61.9	56.5	61.9
8-OCH ₃	4.00 s	4.02 s	-	3.92 s	62.3	62.1	-	56.7
3'-OCH ₃	3.97 s	3.96 s	3.98 s	3.96 s	56.34	56.3	56.4	56.5
4'-3	3.96 s	3.97 s	3.99 s	3.98 s	56.26	56.2	56.7	56.4

3.2.9.5 Structure elucidation of demethylnobiletin (**113**)

The compound **113** was isolated as a dark yellow amorphous solid. The HRESIMS (Figure 3.104) suggested the empirical formula as $C_{20}H_{20}O_8$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 389.1230 (calculate 389.1236). The 1H NMR spectrum (Figure 3.105, Table 3.24) exhibited three aromatic protons as ABX pattern resonating at δ_H 7.00 (d, $J=8.46$ Hz), 7.59 (dd, $J=2.1, 8.46$ Hz) and 7.43 (d, $J=2.1$ Hz), aromatic methine at δ_H 6.60, hydrogen bonded hydroxyl signal at δ_H 12.75 and five methoxy groups at δ_H 4.11 (3H), 3.95 (3H), 3.98 (3H), 3.97 (3H) and 3.98 (3H). The ^{13}C NMR spectrum (Figure 3.106, Table 3.24) showed a total of twenty carbons including a carbonyl at δ_C 183.3 (C-4), five methoxyl carbons, four aromatic methines and the remaining as quaternary carbons. In the HMBC experiment, 3J correlations were observed from methoxyl at δ_H 4.11, 3.95, 3.98, 3.97 and 3.98 to the oxygenated quaternary carbons at δ_C 137.2 (C-6), 153.0 (C-7), 131.2 (C-8), 149.7 (C-3') and 152.1 (C-4') respectively, while H-3 showed long range correlations to δ_C 164.7 (C-2, 2J), 183.3 (C-4, 2J), 108.2 (C-10, 3J) and 124.1 (C-1'). Moreover, the OH group at δ_H 12.75 exhibited along range correlations to carbons at δ_C 153.6 (C-5; by 2J), 137.2 (C-6; by 3J) and 108.2 (C-10; by 3J) and these correlations confirmed the position of the hydroxyl group through C-5. Thus, compound **113** identified as 5-hydroxy-3',4',6,7,8-pentamethoxyflavone. The spectroscopic data of compound **113** were in good agreement with respective published data of demethylnobiletin (Wang *et al.*, 2005). According to Bas *et al.* (2007), compound **113** might have an effect on cell infiltration and induce e of the anti-inflammatory cytokine, interleukin-10. Moreover, demethylnobiletin (**113**) was reported to reduce the apoptosis of T cells, which is a type of lymphocyte (a subtype of white blood cell).

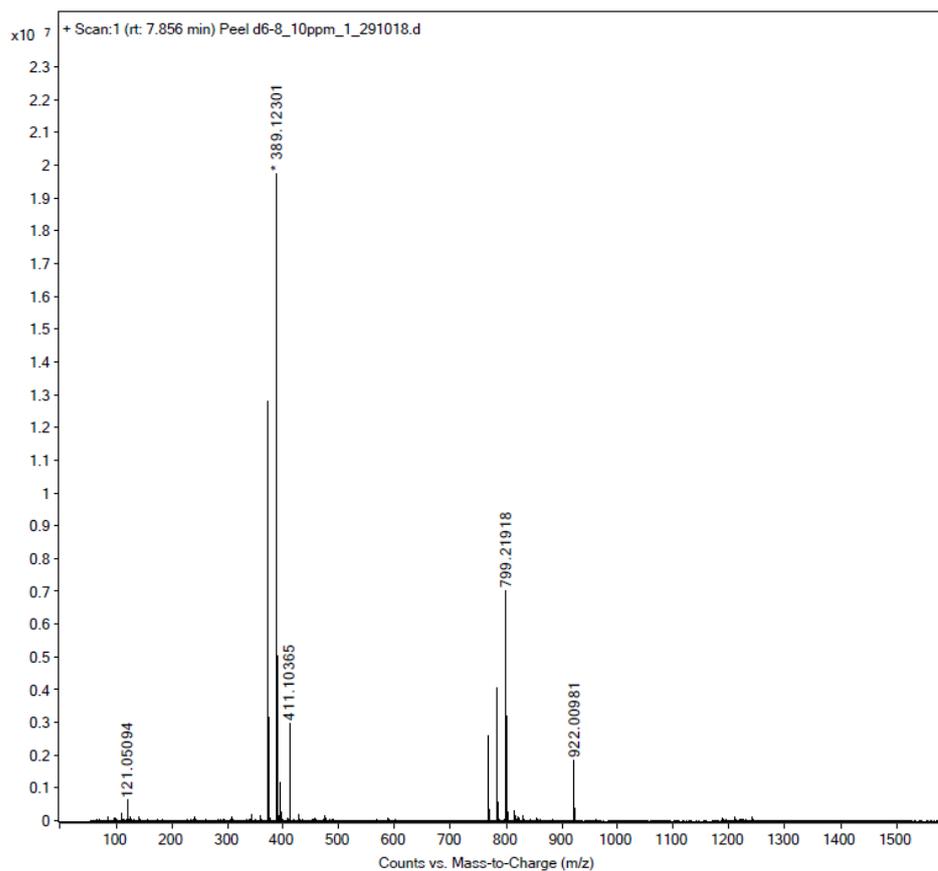


Figure 3.104: The HRESIMS spectrum of compound 113

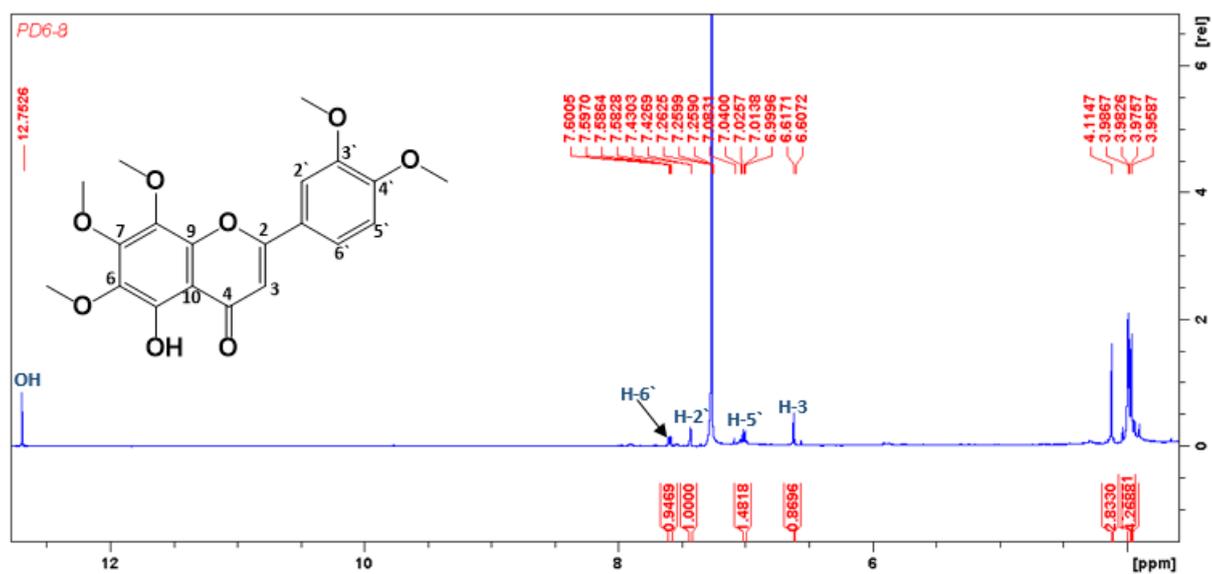


Figure 3.105: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 113

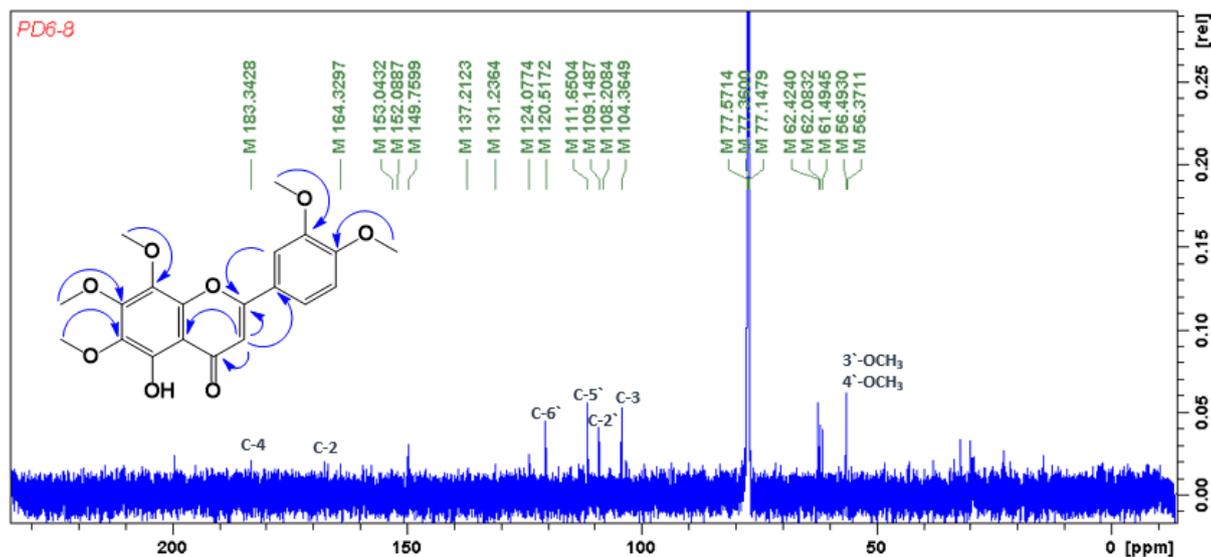


Figure 3.106: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **113**

3.2.9.6 Structure elucidation of 5-desmethylinensetin (**114**)

The compound **114** was isolated as a dark yellow amorphous solid. The HRESIMS (Figure 3.107) suggested the empirical formula as $\text{C}_{19}\text{H}_{18}\text{O}_7$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 359.1141 (calculated 359.1130). The 1D and 2D NMR spectra of compound **114** were very similar to compound **113**. The main differences were the presence of an additional olefinic proton at δ_{H} 6.55 (H-8) and four methoxyl signals instead of five in the ^1H NMR spectrum (Figure 3.108, Table 3.24). The ^{13}C NMR spectrum (Figure 3.109, Table 3.24) revealed a total of nineteen carbons including carbonyl at δ_{C} 183.0 (C-4), four methoxyl carbons, five aromatic methines and the remaining as quaternary carbons. The HMBC experiment (Figure 3.110) showed long range correlations from H-8 to quaternary carbons at δ_{C} 133.6 (C-6, 3J), 159.1 (C-7, 2J) and, 106.5 (C-10, 3J) and 3J correlations were observed from the methoxyl at δ_{H} 3.98 (3H), 3.93 (3H), 3.97 (3H), 3.99 and 3.98 to the oxygenated quaternary carbons at δ_{C} 159.1 (C-7), 91.0 (C-8), 152.7 (C-4') and 149.7 (C-3'), respectively. Thus, compound **114** was identified as 5-hydroxy-3',4',6,7-tetramethoxyflavone. The data of compound **114** were in good agreement with respective published data of 5-desmethylinensetin (Alarif *et al.*, 2013). Compound **114** is documented

as an antiphrastic agent because it inhibited *Trypanosoma cruzi* at 0.4 mg/mL concentration (Beer *et al.*, 2016) and revealed significant activity against (HL-60) leukaemia cell proliferation with induction of apoptosis (Owis, 2019).

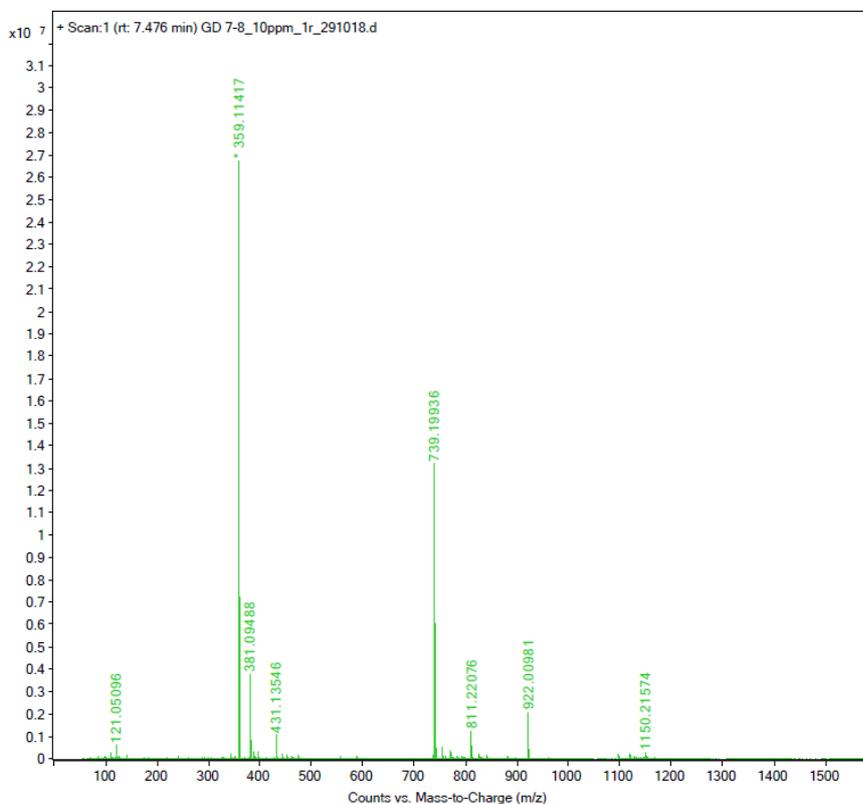


Figure 3.107: The HRESIMS spectrum of compound 114

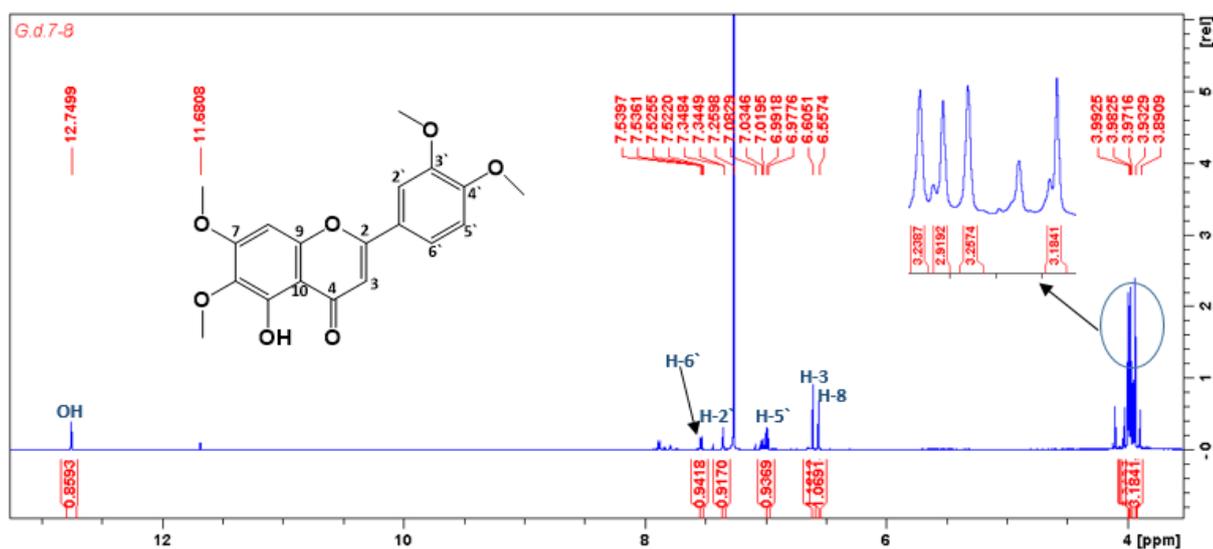


Figure 3.108: ¹H NMR (600 MHz, CDCl₃) spectrum of compound 114

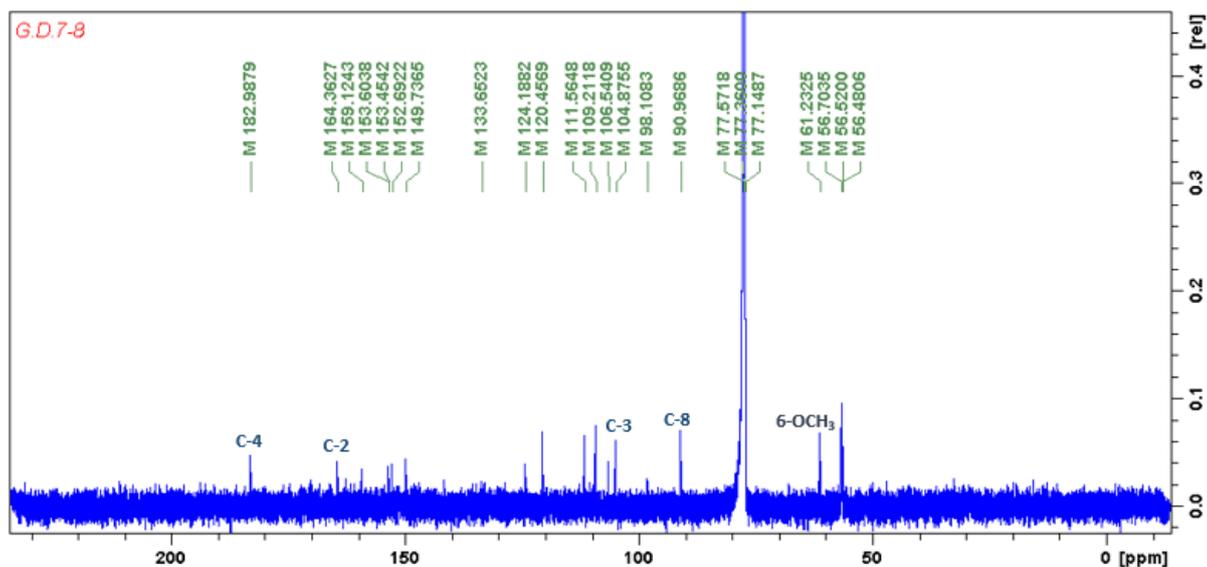


Figure 3.109: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **114**

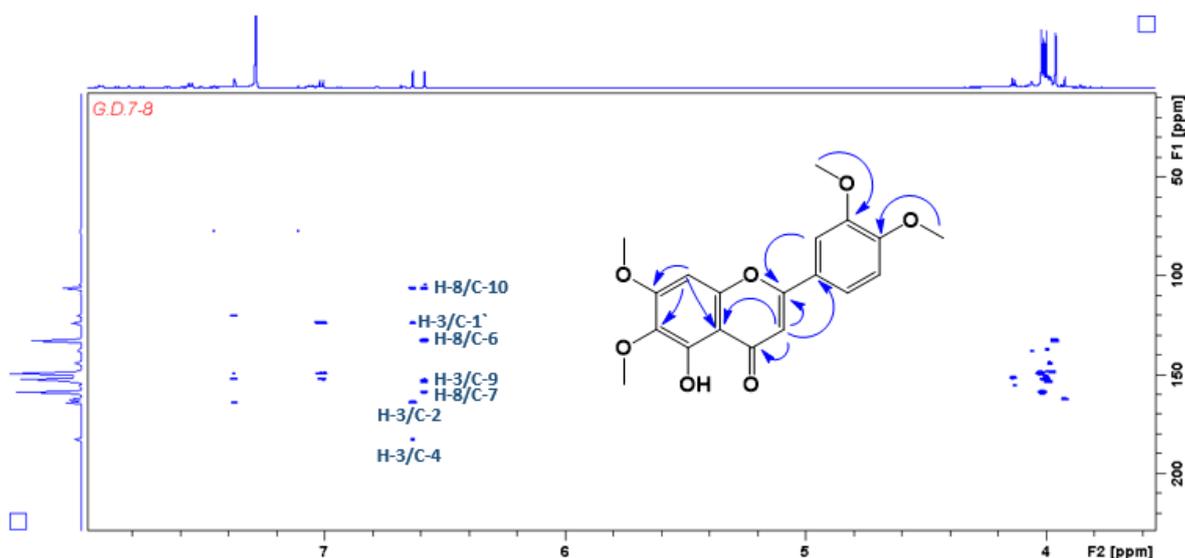


Figure 3.110: The HMBC correlation compound **114**

3.2.9.7 Structure elucidation of cirsilineol (**115**)

The compound **115** was isolated as a dark yellow amorphous solid. The HRESIMS (Figure 3.111) suggested the empirical formula as $\text{C}_{18}\text{H}_{16}\text{O}_7$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 345.0974 (calculated 345.0974). The 1D and 2D NMR spectra of compound **115** were very similar to compound **114**. The main differences were the presence of one extra hydroxyl group at δ_{H} 5.99 and the presence of three methoxyl signals instead of four in the ^1H NMR spectrum (Figure 3.112, Table 3.24). The ^{13}C NMR spectrum

(Figure 3.113, Table 3.24) revealed a total of eighteen carbons including deshelled carbonyl at δ_C 182.9 (C-4), three methoxyl groups, five methine carbons and the remains were quaternary carbons. The HMBC (Figure 3.114) showed the long range correlation from OH (δ_H 5.99) to carbons at δ_C 147.2 (C-3', 3J), 149.6 (C-4', 2J) and 115.4 (C-5', 3J) while the OH group at δ_H 12.75 showed a correlation to carbons at δ_C 153.6 (C-5, 2J), 133.6 (C-6, 3J), 106.5 (C-10, 3J) these correlations confirmed the position of hydroxyl groups in the molecule. Moreover, 3J correlations were observed from methoxyl at δ_H 3.93 (3H), 3.97 (3H) and 4.01 (3H) to the oxygenated quaternary carbons at δ_C 133.1 (C-6), 159.1 (C-7) and 147.2 (C-3') respectively. Thus, compound **115** identified as 4',5-dihydroxy-3',6,7-trimethoxyflavone or cirsilineol. The spectroscopic data of compound **115** were in good agreement with respective published data for cirsilineol (Hammoud *et al.*, 2012). Compound **115** is reported to have anticancer activity against human ovarian cancer (Caov-3, Skov-3) and prostate cancer cell PC3 (Sheng *et al.*, 2008).

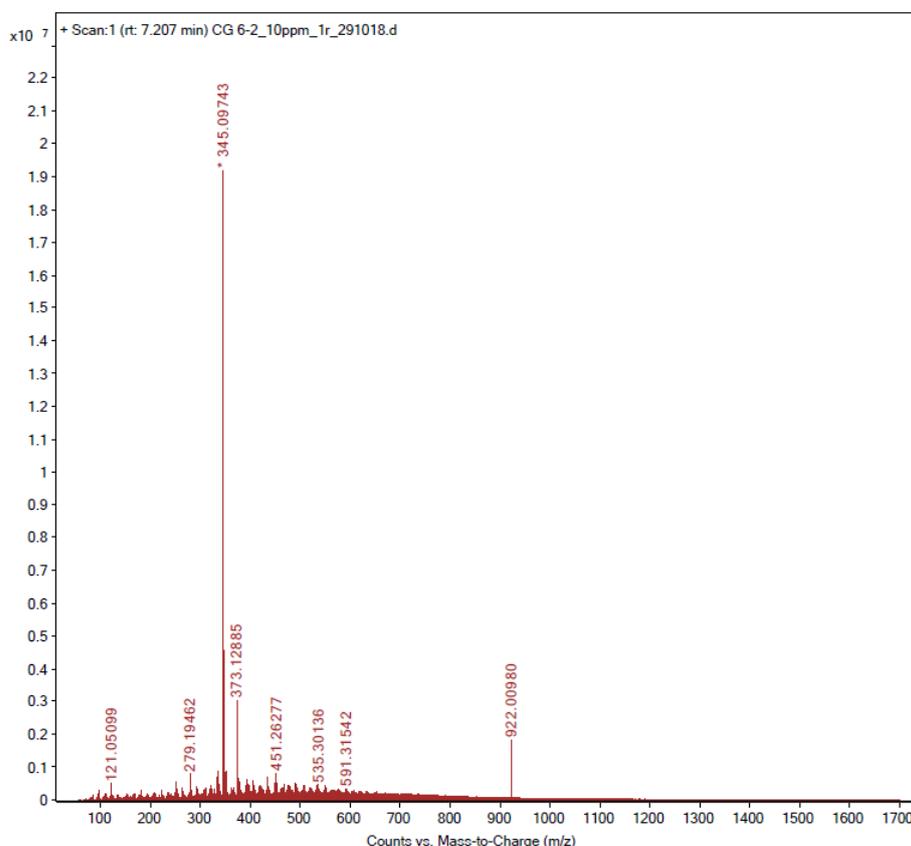


Figure 3.111: The HRESIMS spectrum of compound **115**

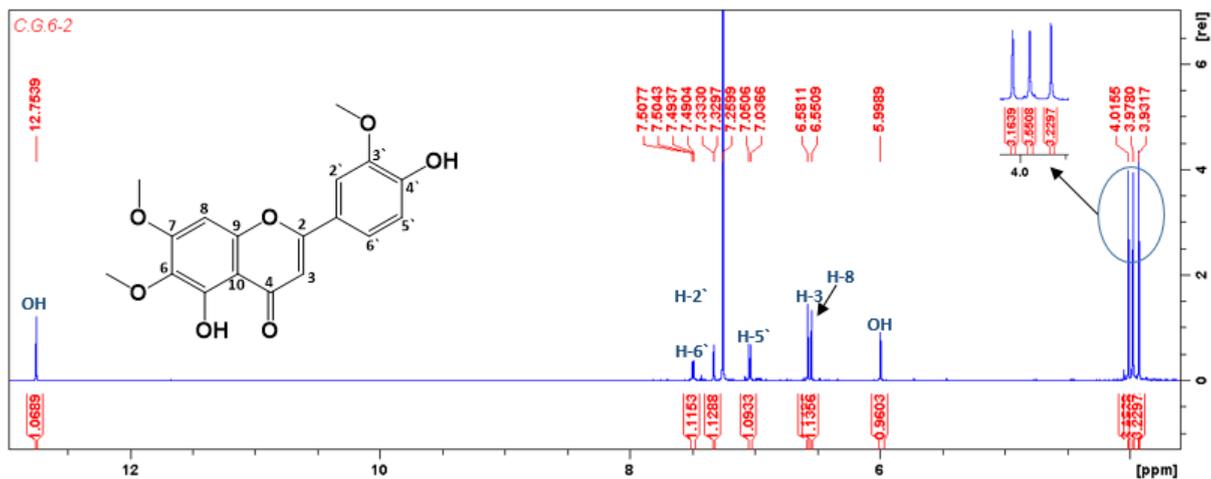


Figure 3.112: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 115

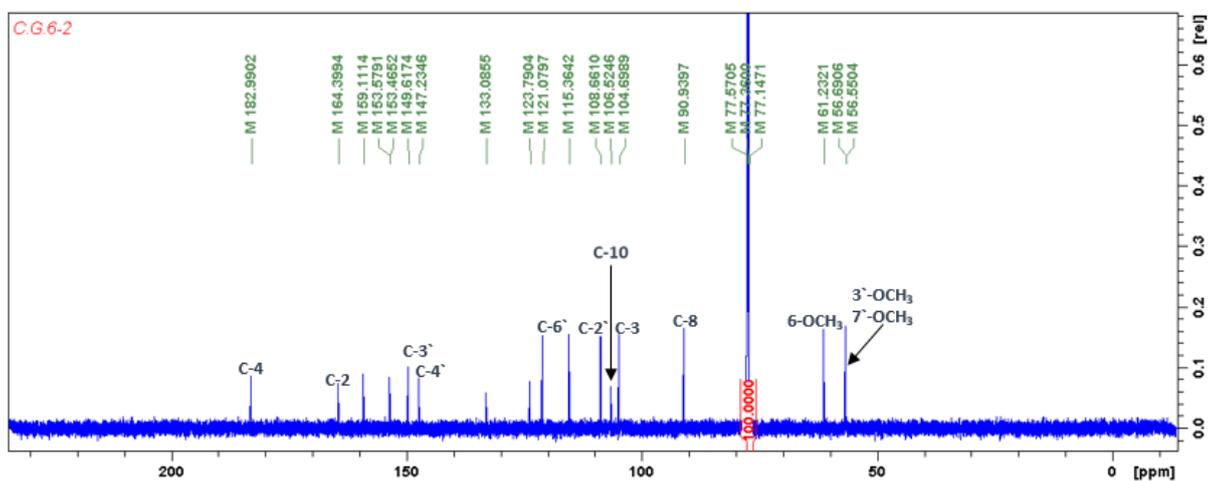


Figure 3.113: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound 115

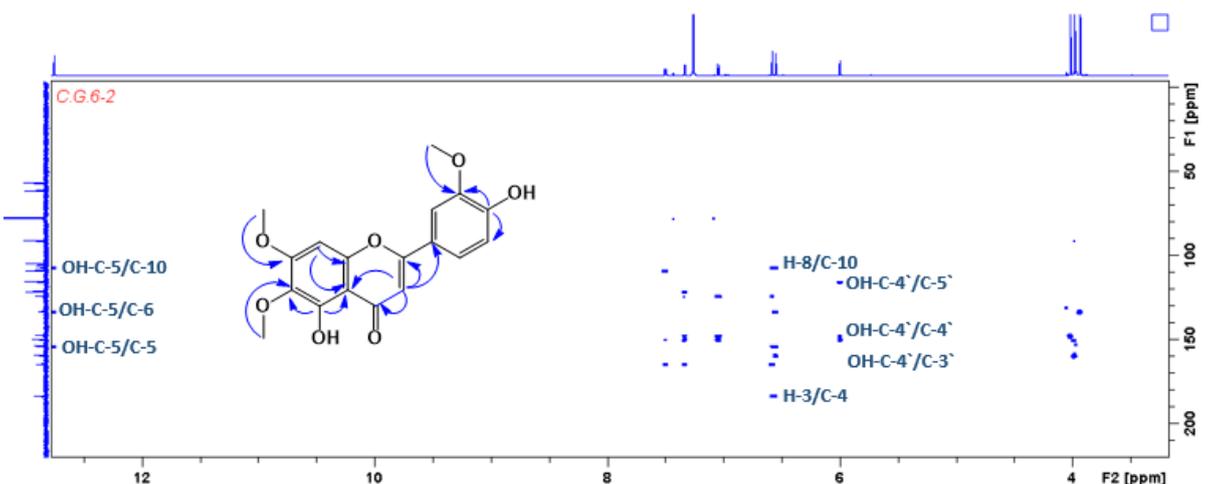


Figure 3.114: The HMBC correlation compound 115

Table 3.24: ^1H NMR (600 MHz) and ^{13}C NMR data of compounds **113-115**

position	Chemical Shift δ_{H} (ppm), J in Hz			Chemical Shift δ_{C} (ppm), J in Hz		
	113	114	115	113	114	115
2	-	-	-	164.7	164.4	164.4
3	6.61 s	6.60 s	6.58 s	104.4	104.9	104.7
4	-	-	-	183.3	183.0	183.0
5	-	-	-	149.9	153.6	153.6
6	-	-	-	137.2	133.6	133.1
7	-	-	-	153.0	159.1	159.1
8	-	6.55 s	6.55 s	131.2	91.0	91.0
9	-	-	-	146.3	153.4	153.5
10	-	-	-	108.2	106.5	106.5
1'	-	-	-	124.1	124.2	123.8
2'	7.43 <i>d</i> (2.1)	7.34 <i>d</i> (2.04)	7.33 <i>d</i> (2.1)	109.14	109.2	108.7
3'	-	-	-	149.75	149.7	147.2
4'	-	-	-	152.08	152.7	149.6
5'	7.00 <i>d</i> (8.46)	6.98 <i>d</i> (8.04)	7.04 <i>d</i> (8.34)	111.65	111.6	115.4
6'	7.59 <i>dd</i> (2.1, 8.46)	7.53 <i>dd</i> (8.04, 2.04)	7.50 <i>d</i> (8.34, 2.1)	120.51	120.4	121.1
6-OCH ₃	4.11 s	-	3.93 s	62.1	-	61.2
7-OCH ₃	3.95 s	3.98 s	3.97 s	61.4	56.7	56.7
8-OCH ₃	3.98 s	3.93 s	-	62.4	61.2	-
3'-OCH ₃	3.97 s	3.99 s	-	56.5	56.48	56.5
4'-OCH ₃	3.98 s	3.97 s	-	56.4	56.5	-
5-OH	-	-	12.77 s	-	-	-
4'-OH	-	-	5.99 s	-	-	-

3.2.9.8 Structure elucidation of tangeritin (116)

The compound **116** was isolated as yellow amorphous powder. The HRESIMS (Figure 3.115) suggested the empirical formula as $\text{C}_{20}\text{H}_{20}\text{O}_7$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 373.1286 (calculated 373.1287). The ^1H NMR spectrum (Figure 3.116, Table 3.25) exhibited three sets of aromatic protons at δ_{H} 7.88 (2H, *d*, $J=8.82$, H-2', 6'), 7.03 (2H, *d*, $J=8.82$, H-3', 5'), singlet at δ_{H} 6.60 (H-3) and methoxy groups at δ_{H} 3.89 (3H), 3.95 (2X 3H), 4.02 (3H) and 4.10 (3H). The ^{13}C NMR spectrum (Figure 3.117, Table 3.25) showed a total of twenty carbons including a carbonyl group at δ_{C} 177.7, five

methoxyls groups, five methines and nine quaternary carbons. The HMBC experiment showed 3J correlations from the methoxyl at δ_H 3.89 (3H), 4.02 (3H) and 4.10 (3H) to the oxygenated quaternary carbons at δ_C 162.6 (C-4'), 138.3 (C-8) and 151.7 (C-5) while the signal at δ_H 3.95 revealed correlations to carbons at δ_C 144.4 (C-6) and 148.8 (C-7). Moreover, a 3J correlations were observed from H-2' and H-6' to carbons at δ_C 161.5 (C-2) and 162.6 (C-4') additionally, long range correlations detected from H-3' and H-5' to carbons at δ_C 124.2 (C-1'). Thus, compound **116** identified as 5,6,7,8,4'-pentamethoxyflavone or tangeritin. The data of compound **116** were in good agreement with respective published data for tangeritin (Hamdan *et al.*, 2011). Antioxidant and anti-inflammatory effect of compound **116** have been reported (Chen *et al.*, 2017). Tangeritin was showed a significant reduction in Parkinson's disease symptoms (Fatima *et al.*, 2017).

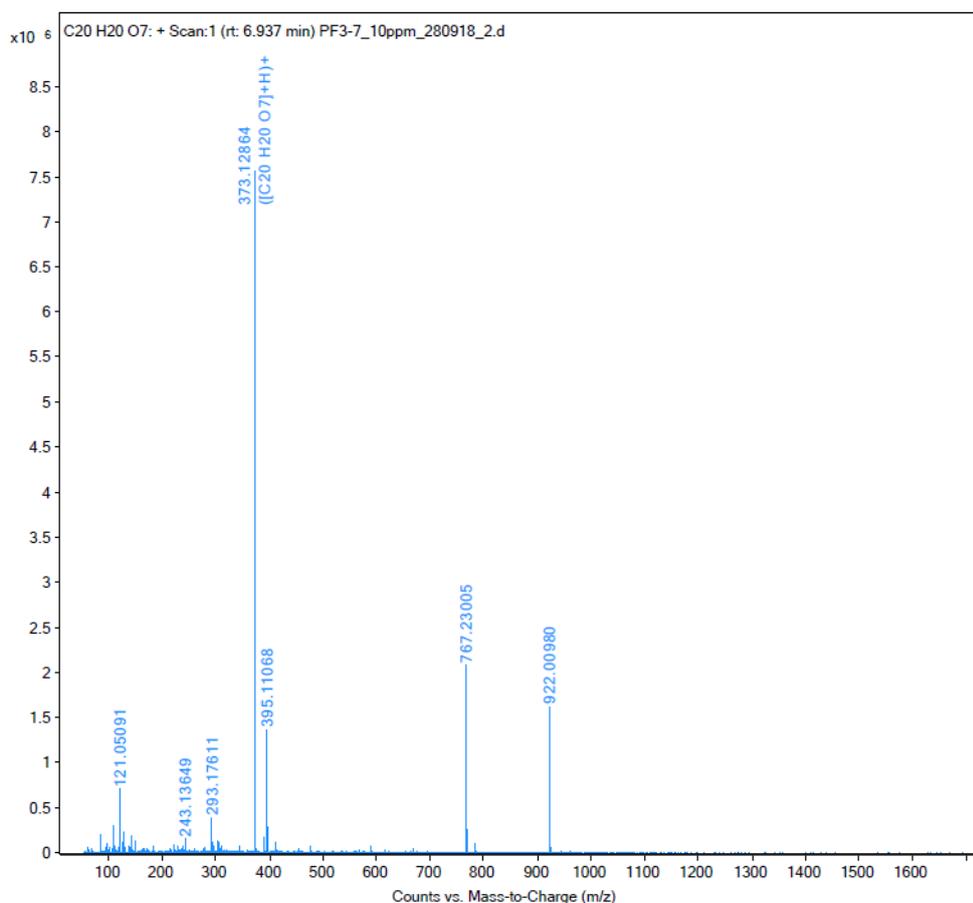


Figure 3.115: The HRESIMS spectrum of compound **116**

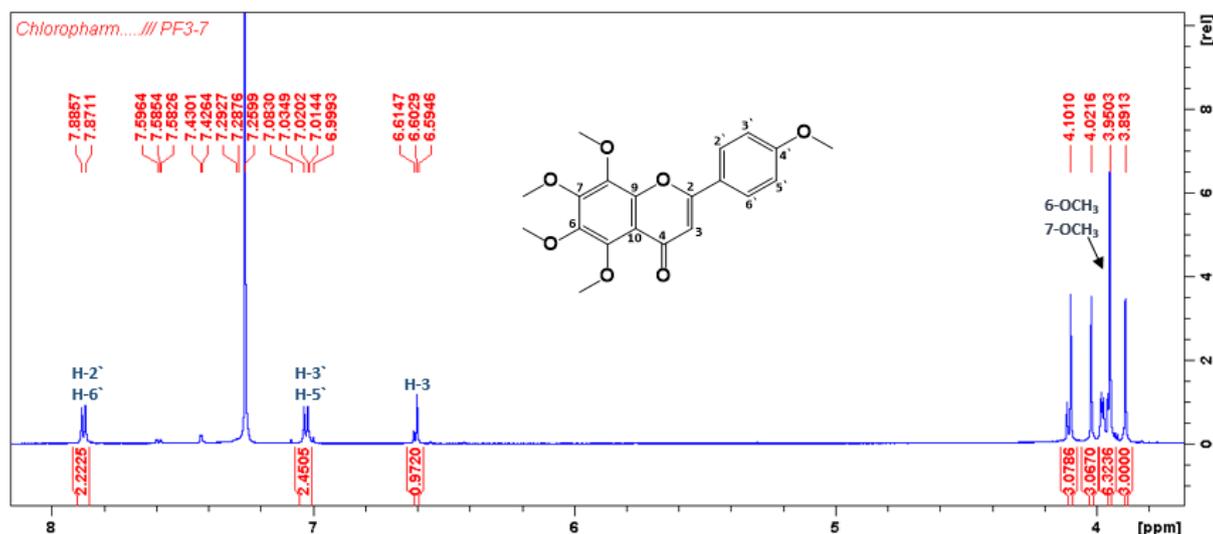


Figure 3.116: ^1H NMR (600 MHz, CDCl_3) spectrum of compound **116**

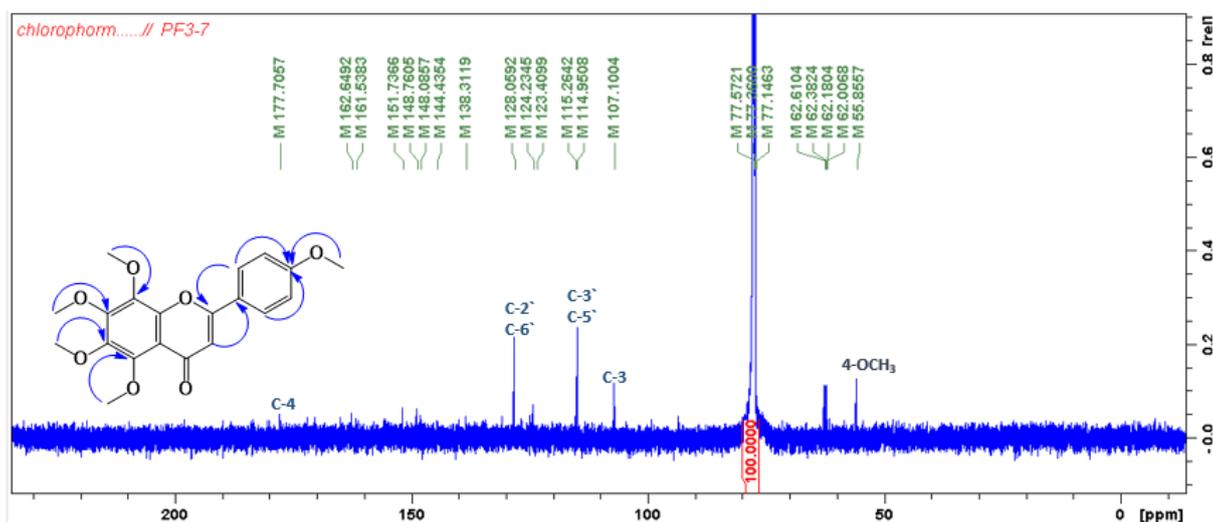


Figure 3.117: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **116**

3.2.9.9 Structure elucidation of tetramethylscutellarein (**117**)

The compound **117** was isolated as a yellow powder. The HRESIMS (Figure 3.118) suggested the empirical formula as $\text{C}_{19}\text{H}_{18}\text{O}_6$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 343.1191 (calculated 343.1181). The 1D and 2D NMR spectra of compound **117** were very similar to compound **116**. The only differences were the presence of an additional olefinic proton at δ_{H} 6.79 (H-8) and four methoxyl signals instead of five in the ^1H NMR spectrum (Figure 3.119, Table 3.25). The ^{13}C NMR spectrum (Figure 3.120, Table 3.25) revealed a total of eighteen carbons including a carbonyl at δ_{C} 177.7, four methoxyl groups,

six methines and the others are quaternary carbons. In the HMBC experiment (Figure 3.121), long-range correlations from the singlet proton at δ_{H} 6.59 (H-8) were revealed to carbons resonating at δ_{C} 161.6 (C-2), 177.7 (C-4), 140.7 (C-6), 154.9 (C-9) and 113.1 (C-10). Moreover, 3J correlations were observed from methoxyl at δ_{H} 3.98 (3H), 3.91 (3H), 3.97 (3H) and 3.88 (3H) to the oxygenated quaternary carbons at δ_{C} 152.9 (C-5), 140.7 (C-6), 158.0 (C-7) and 162.5 (C-4') respectively. Thus, compound **117** was identified as 5,6,7,4'-tetramethoxyflavone. The data of compound **117** were in good agreement with respective published data of tetramethylscutellarein (Li *et al.*, 2018). This compound **117** is reported to have showed antidiabetic activity by inhibiting α -glucosidase enzyme with an IC_{50} 0.75 mM (Damsud *et al.*, 2014).

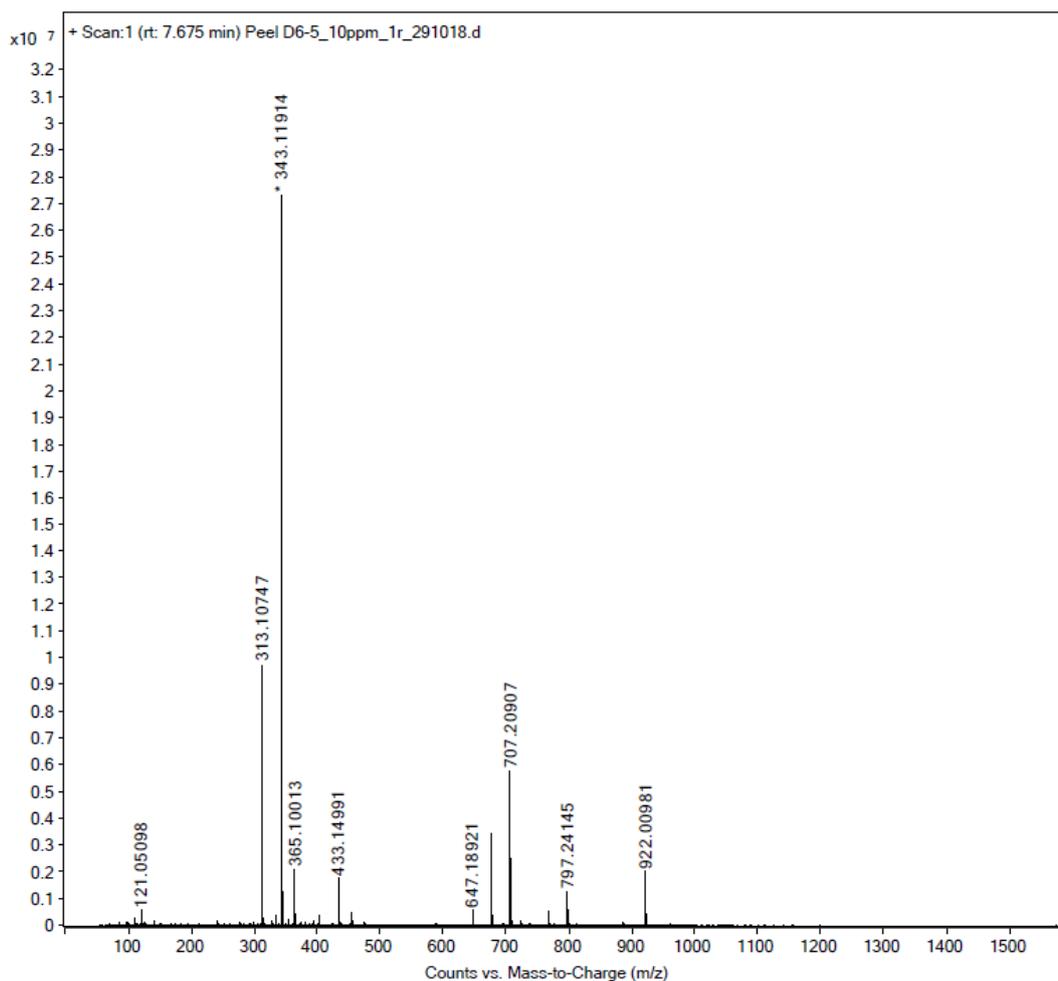


Figure 3.118: The HRESIMS spectrum of compound **117**

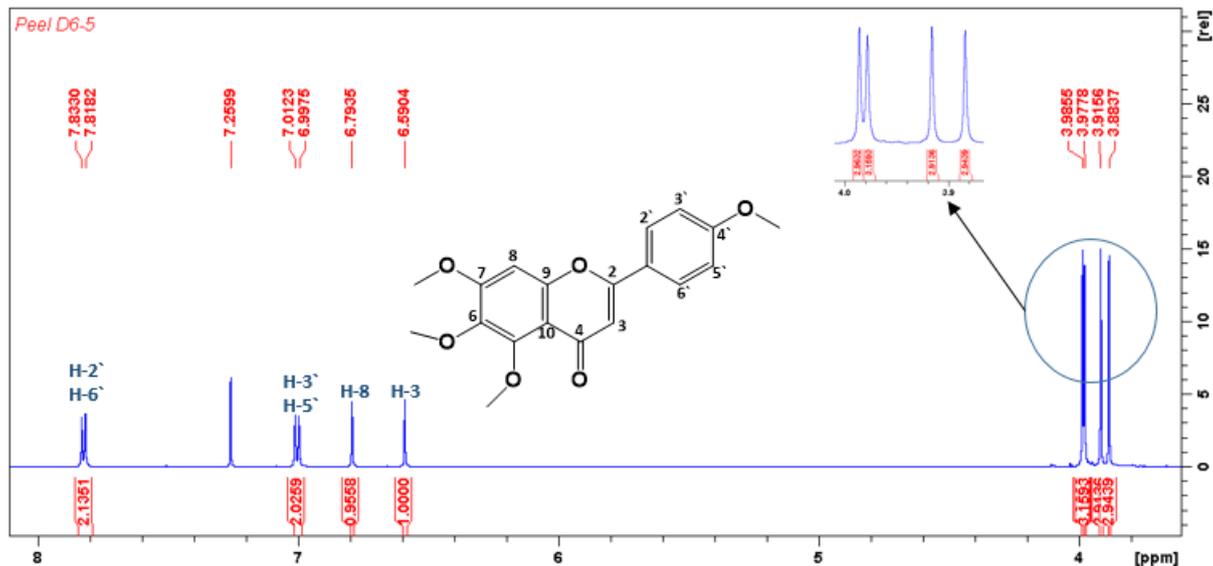


Figure 3.119: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 117

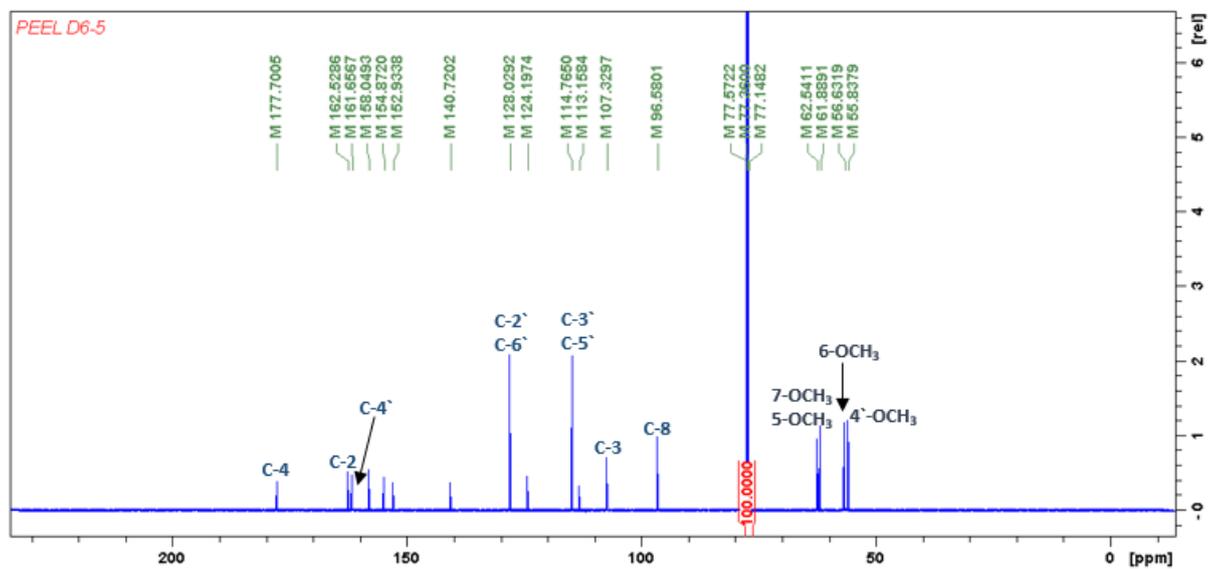


Figure 3.120: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound 117

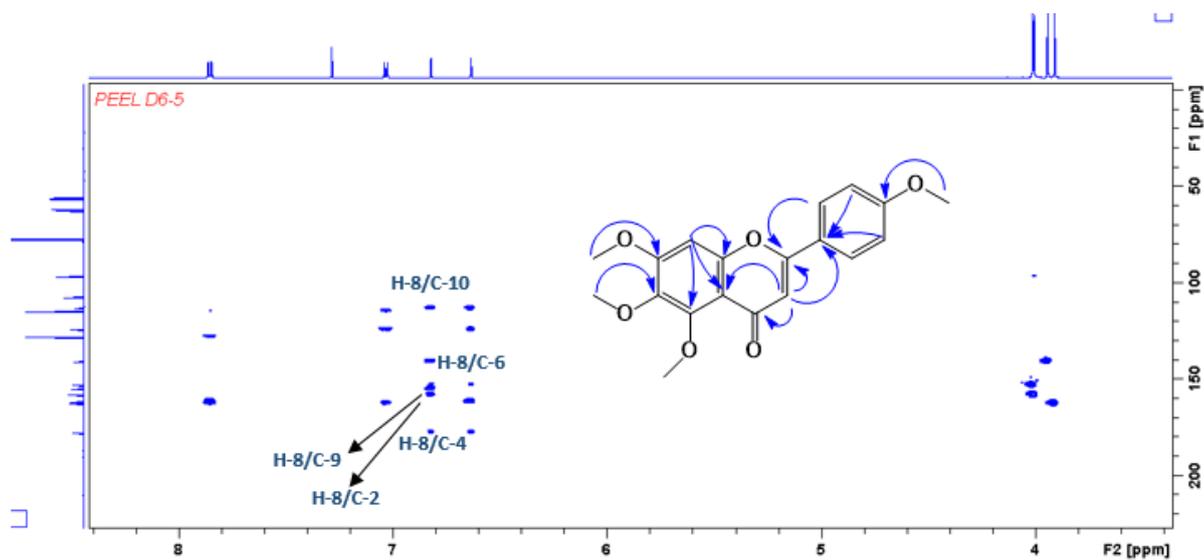


Figure 3. 121: The HMBC correlation compound 117

3.2.9.10 Structure elucidation of salvigenin (118)

The compound **118** was isolated as a yellow powder. The HRESIMS (Figure 3.122) suggested the empirical formula as $C_{18}H_{16}O_6$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 329.1259 (calculated 329.1025). The 1D and 2D NMR spectra of compound **118** were very similar to those of compound **117**. The main differences were the presence of a hydroxyl peak at δ_H 12.75 and the presence of three methoxyl signals instead of four in the 1H NMR spectrum (Figure 3.123, Table 3.25). The ^{13}C NMR spectrum (Figure 3.124, Table 3.25) revealed a total of eighteen carbons including a carbonyl at δ_C 182.8, three methoxy groups, six methines and the remains are quaternary carbons. The HMBC experiment showed long correlation from the hydrogen bonded hydroxyl (δ_H 12.75) to the carbons at δ_C 154.88 (C-5), 131.94 (C-6) and 107.32 (C-10); these correlations confirmed the position of the hydroxyl group through C-5 in the molecule. Moreover, a 3J correlations were observed from methoxyl at δ_H 3.88 (3H), 3.91 (3H) and 3.97 (3H) to the oxygenated quaternary carbons at δ_C 162.1 (C-4'), 131.9 (C-6) and 158.0 (C-7) respectively. Additionally, long range correlations were observed from H-2' and H-6' to carbons at δ_C 124.2 (C-1') and 162.1 (C-4'). Thus, compound **118** identified as 5-hydroxy-6,7,4'-trimethoxyflavone. The spectroscopic data of compound **118** were in good agreement with respective published data of salvigenin (Morocho *et al.*, 2018). Compound **118** possesses anti-inflammation activity (Mansourabadi *et al.*, 2015) and cytotoxic activity agansit the breast cancer cell line (MCF-7) in a mouse model (Noori *et al.*, 2013).

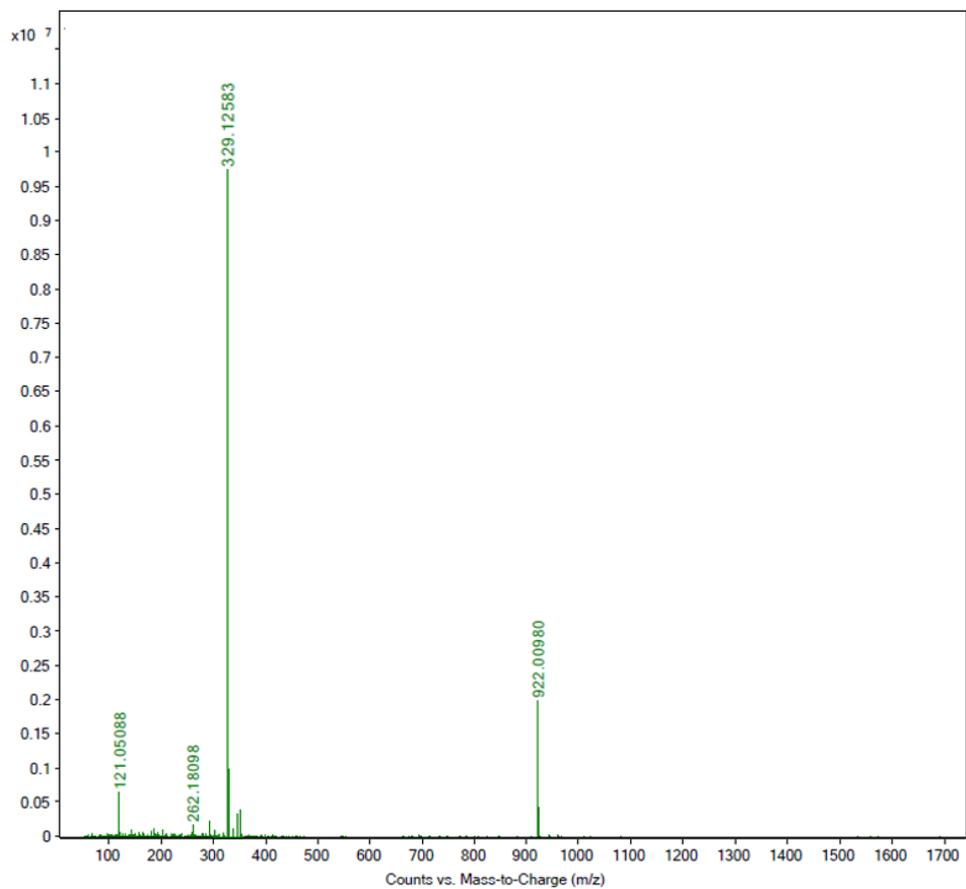


Figure 3.122: The HRESIMS spectrum of compound 118

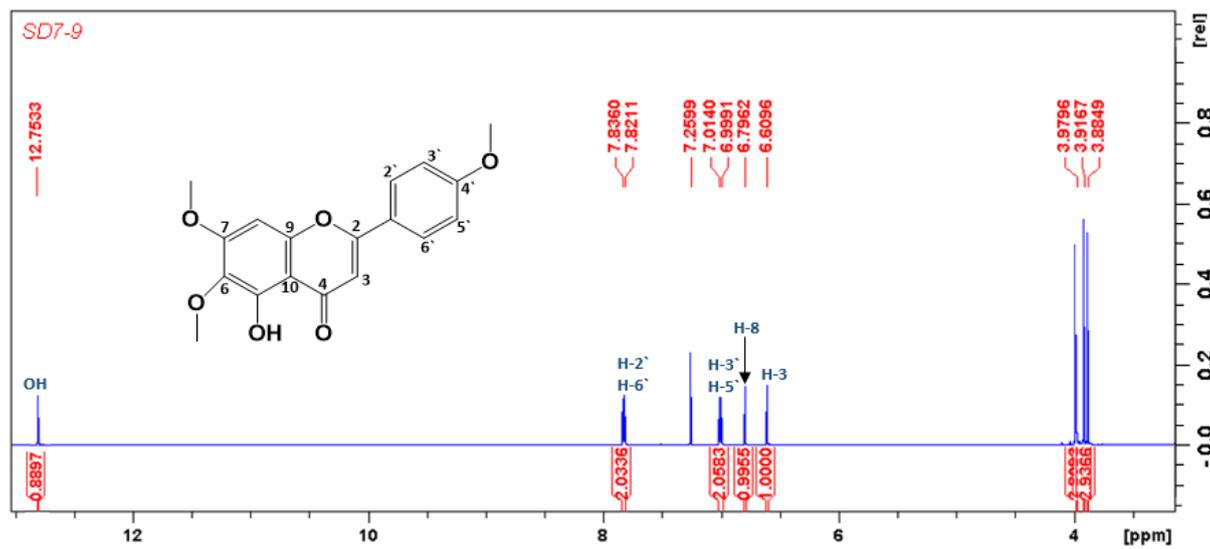


Figure 3.123: ¹H NMR (600 MHz, CDCl₃) spectrum of compound 118

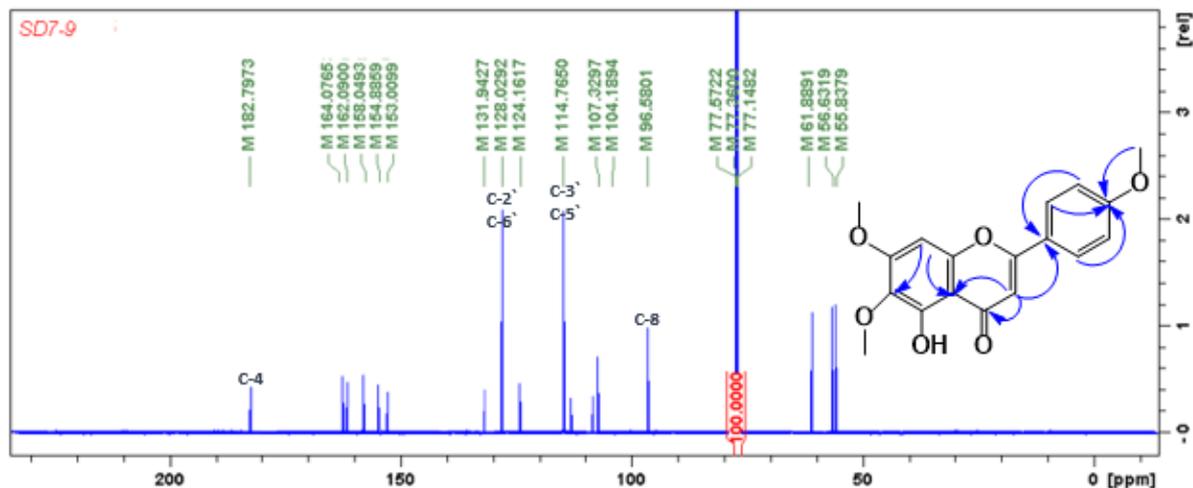


Figure 3.124: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound 118

Table 3.25: ^1H NMR (600 MHz) and ^{13}C NMR data of compounds 116-118

position	Chemical Shift δ_{H} (ppm), J in Hz			Chemical Shift δ_{C} (ppm), J in Hz		
	116	117	118	116	117	118
2	-	-	-	161.5	161.6	164.1
3	6.60 s	6.59 s	6.60 s	107.1	107.1	104.2
4	-	-	-	177.7	177.7	182.8
5	-	-	-	151.7	152.9	154.9
6	-	-	-	144.4	140.7	131.9
7	-	-	-	148.8	158.0	158.0
8	-	6.59 s	6.79 s	138.3	96.6	96.6
9	-	-	-	148.0	154.9	153.0
10	-	-	-	115.3	113.1	107.3
1'	-	-	-	124.2	124.2	124.2
2'	7.88 <i>d</i> (8.82)	7.82 <i>d</i> (8.82)	7.83 <i>d</i> (8.82)	128.0	128.0	128.0
3'	7.03 <i>d</i> (8.82)	7.00 <i>d</i> (8.82)	7.00 <i>d</i> (8.82)	115.0	114.8	114.8
4'	-	-	-	162.7	162.5	162.1
5'	7.03 <i>d</i> (8.82)	7.00 <i>d</i> (8.82)	7.00 <i>d</i> (8.82)	115.0	114.8	114.8
6'	7.88 <i>d</i> (8.82)	7.82 <i>d</i> (8.82)	7.83 <i>d</i> (8.82)	128.0	128.0	128.0
5-OCH ₃	4.10 s	3.98 s	-	62.0	62.5	-
6-OCH ₃	3.95 s	3.91 s	3.91 s	62.4	61.9	61.9
7-OCH ₃	3.95 s	3.97 s	3.97 s	62.6	56.6	56.6
8-OCH ₃	4.02 s	-	-	62.2	-	-
4'-OCH ₃	3.89 s	3.88 s	3.88 s	55.8	55.8	55.8
5-OH	-	-	12.7	-	-	-

3.2.9.11 Structure elucidation of hesperidin (50)

The compound **50** was isolated as a white powder. The HRESIMS (Figure 3.125) suggested the empirical formula as $C_{28}H_{34}O_{15}$ and in the positive ion mode it showed peak $[M+H]^+$ at m/z 611.1953 (calculated 611.1976). The 1H NMR spectrum (Figure 3.126, Table 3.26) exhibited three aromatic protons as ABX pattern resonating at δ_H 6.91 (*dd*, $J=8.34$, 2.1 Hz), 6.93 (*d*, $J=2.1$ Hz), 6.95 (*d*, $J=8.34$ Hz) and two *meta*-coupled ($J=2.5$ Hz) protons at δ_H 6.14 (*d*, $J=2.5$ Hz, H-6) and 6.12 (*d*, $J=2.5$ Hz, H-8). Moreover, two hydroxyl peaks were observed at δ_H 12.96 and 9.08. Moreover, the rutinosyl moiety was revealed by the presence of a glucose anomeric proton at δ_H 4.98 (*d*, $J=8.5$ Hz), while the rhamnose anomeric proton resonated at δ_H 4.52 (*d*, $J=1.92$ Hz) and methyl protons of rhamnose at δ_H 1.09 (3H). The primary elucidation of the 1H NMR spectrum of compound **50** suggested this compound could be a flavanone glycoside. The ^{13}C NMR spectrum (Figure 3.127, Table 3.26) showed a total of twenty eight signals including a carbonyl at δ_C 197.5, a methoxyl, seven quaternary carbons and two aliphatic methine carbons at δ_C 78.8 (C-2) and 42.5 (C-3). Among these, twelve carbons constituted the rutinosyl moiety. The HMBC experiment showed the long range correlation from H-1''' (δ_H 4.52) to carbons at δ_C 66.78 (C-6'') which confirmed the connectivity of glucose and rhamnose. Another 3J correlation from the proton at δ_H 4.98 (H-1'') to the oxygenated quaternary carbon at δ_C 165.60 (C-7) confirmed the linkage of sugar through C-7. The methoxyl δ_H 3.78 revealed a correlation to oxygenated quaternary carbon at δ_C 146.9 (C-3'). Thus, compound **50** was identified as 7-rhamnoglucoside. The spectroscopic data of compound **50** were in good agreement with respective published data of hesperidin (Nieto and Gutierrez, 1986). Compound **50** is reported as a food supplement, decreasing the permeability of blood capillaries, is an antihypercholesterolaemia agent, antihyperlipidemic factor and an antioxidant (Garg *et al.*, 2001; Guardia *et al.*, 2001).

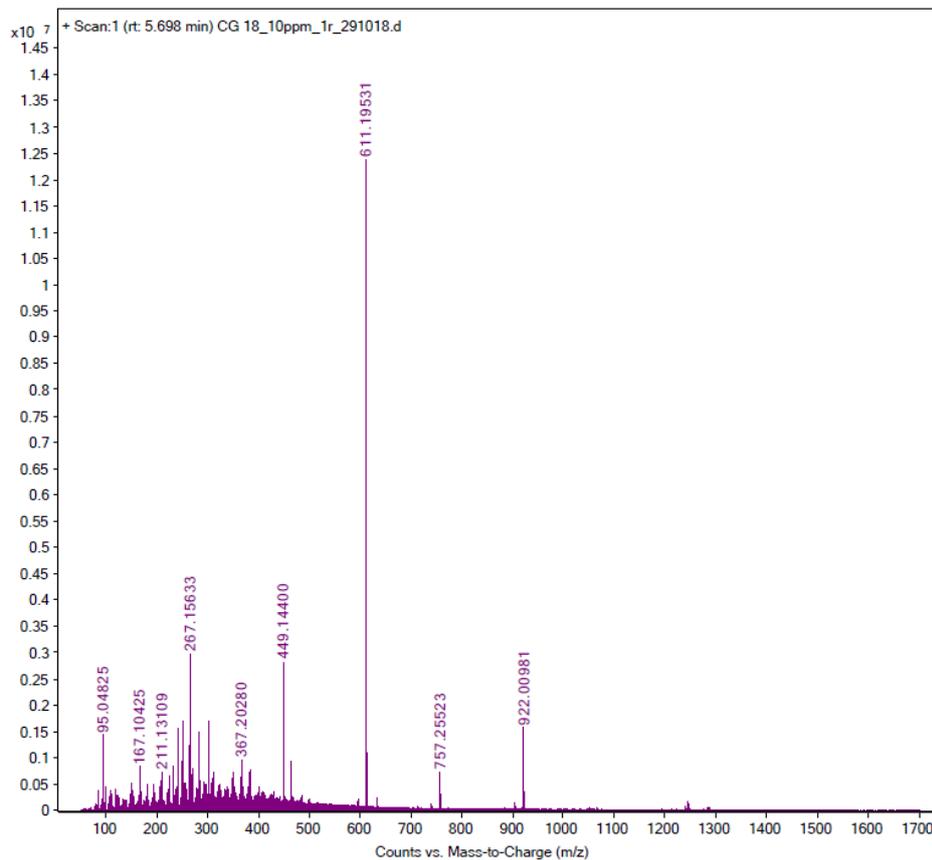


Figure 3.125: The HRESIMS spectrum of compound 50

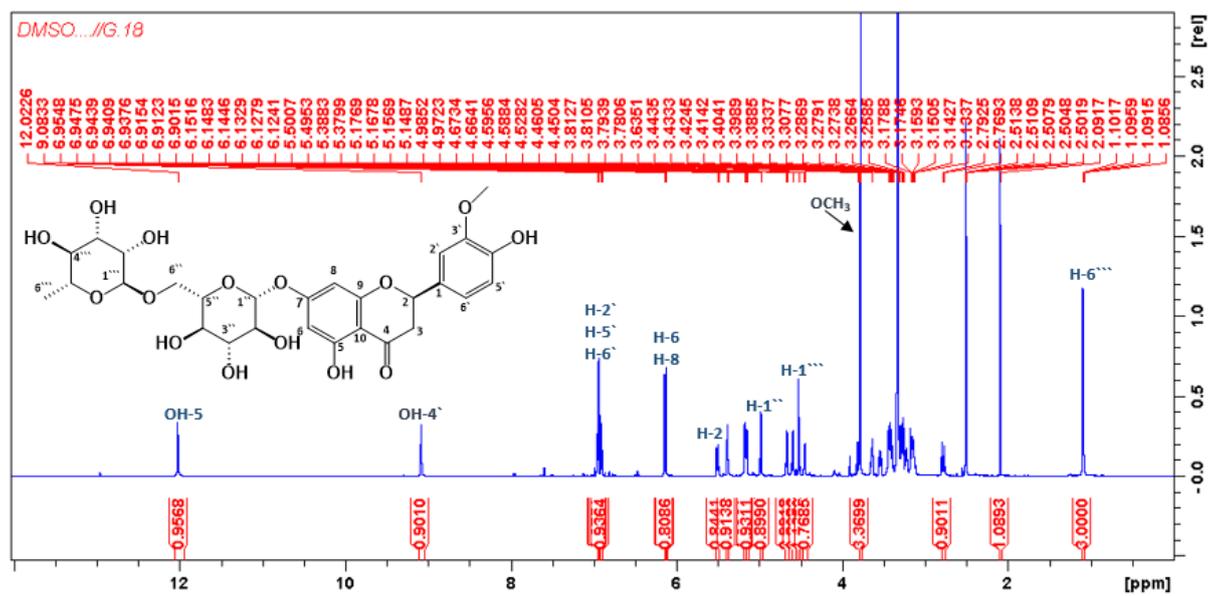


Figure 3.126: ^1H NMR (600 MHz, CDCl_3) spectrum of compound 50

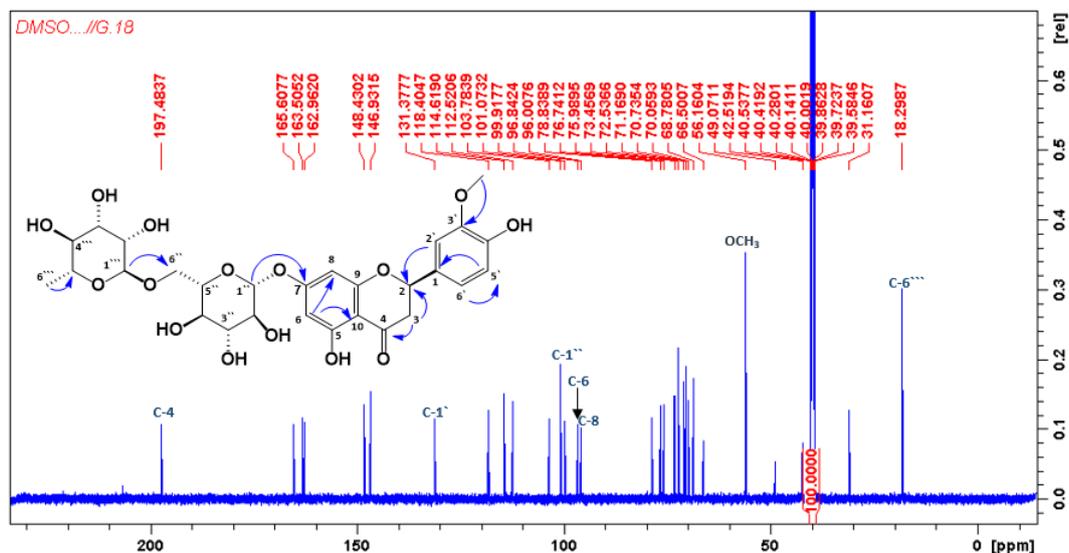


Figure 3.127: ^{13}C NMR (150 MHz, CDCl_3) of compound **50**

3.2.9.12 Structure elucidation of compound narirutin (**51**)

The compound **51** was isolated as a yellow powder. The HRESIMS suggested the empirical formula as $\text{C}_{27}\text{H}_{32}\text{O}_{14}$ and in the positive ion mode it showed $[\text{M}+\text{H}]^+$ peak at m/z 581.1895 (calculated 581.1870). The ^1H NMR spectra (Figure 3.128, Table 3.26) exhibited a methyl signal at δ_{H} 1.21(3H) and two *meta*-coupled olefinic protons at δ_{H} 6.19 (H-6) and 6.22 (H-8) and *ortho* ($J=8.52$) coupled signals resonated at δ_{H} 7.36 (2H, H-2', H-6') and 6.85 (2H, H-3', H-5'). The ^{13}C NMR spectrum (Figure 3.129, Table 3.26) revealed a total of twenty-eight carbons including a carbonyl at δ_{C} 198.7, six quaternary carbons and two aliphatic methine carbons at δ_{C} 78.0 (C-2) and 42.3 (C-3) and twelve carbons forming the rutosyl moiety. The HMBC experiment (Figure 3.130) showed 3J correlation from H-2' and H-6' to carbon at carbons at δ_{C} 149.2 (C-4') and 78.0 (C-2) while H-3' and H-5' revealed long range correlations to carbons δ_{C} 131.0 (C-1', 3J) and 149.2 (C-4', 2J). The cross peak correlation from the anomeric proton of rhamnose at δ_{H} 4.71 and carbon at δ_{C} 69.92 (C-6'') confirmed the linking of rhamnose to glucose while the anomeric proton of glucose at δ_{H} 4.96 revealed a correlation to carbon at δ_{C} 166.95 (C-7) confirming the position of the rutosyl moiety through C-7 in the molecule. Thus, compound **51** was identified as apigenin

7-*O*- β -D-glucopyranosyl (6 \rightarrow 1)- α -L-rhamnopyranoside. The data of compound **51** were in good agreement with respective published data of narirutin (Abu-Gharbieh and Shehab, 2017). Compound **51** was reported to be useful for the treatment of bronchial asthma (Funaguchi *et al.*, 2007) and to reduce liver damage caused by drinking alcohol (Park *et al.*, 2013).

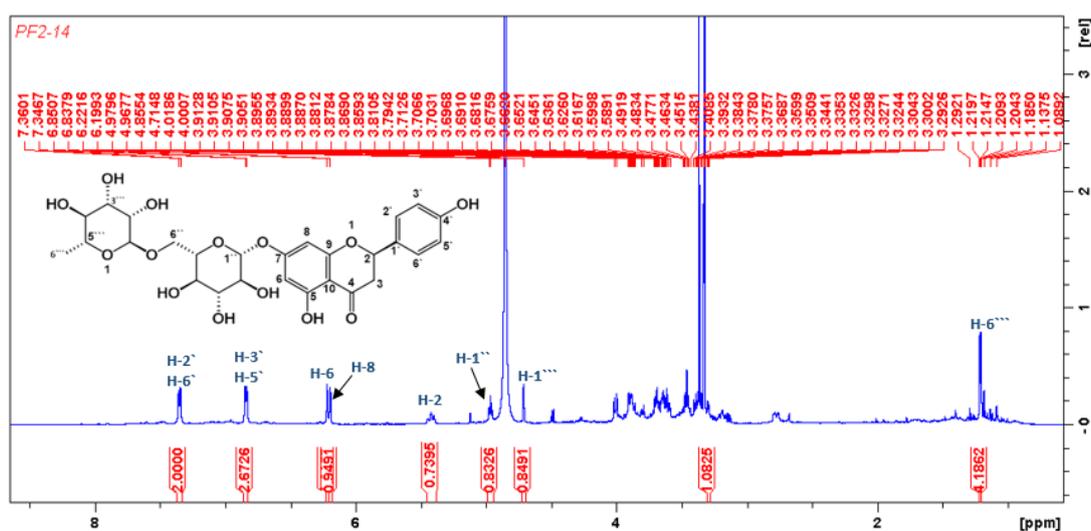


Figure 3.128: ^1H NMR (600 MHz, CDCl_3) spectrum of compound **51**

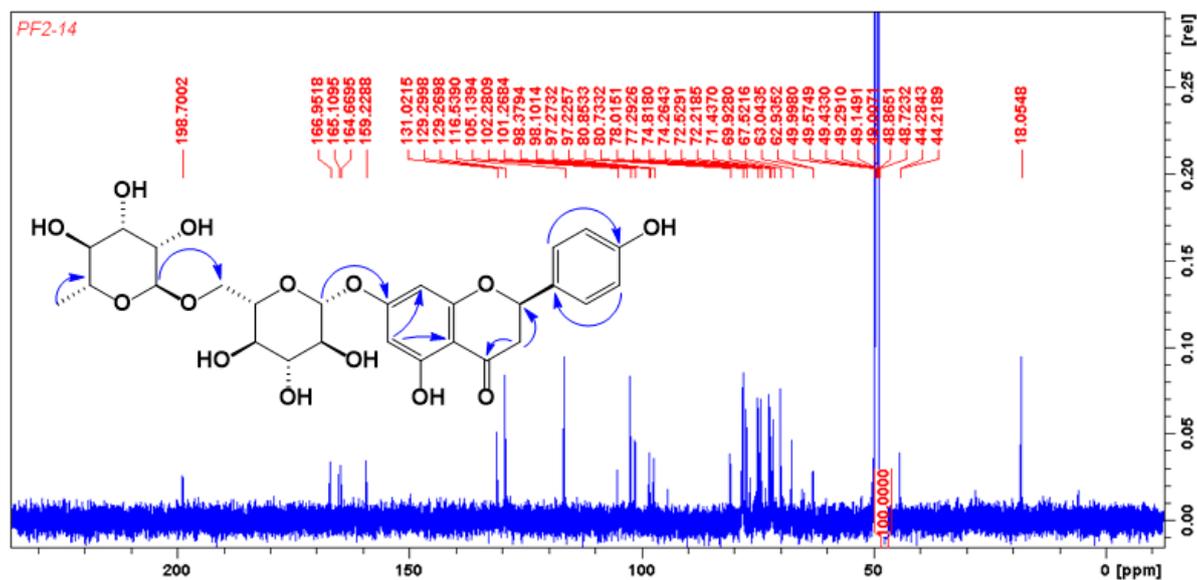


Figure 3.129: ^{13}C NMR (150 MHz, CDCl_3) spectrum of compound **51**

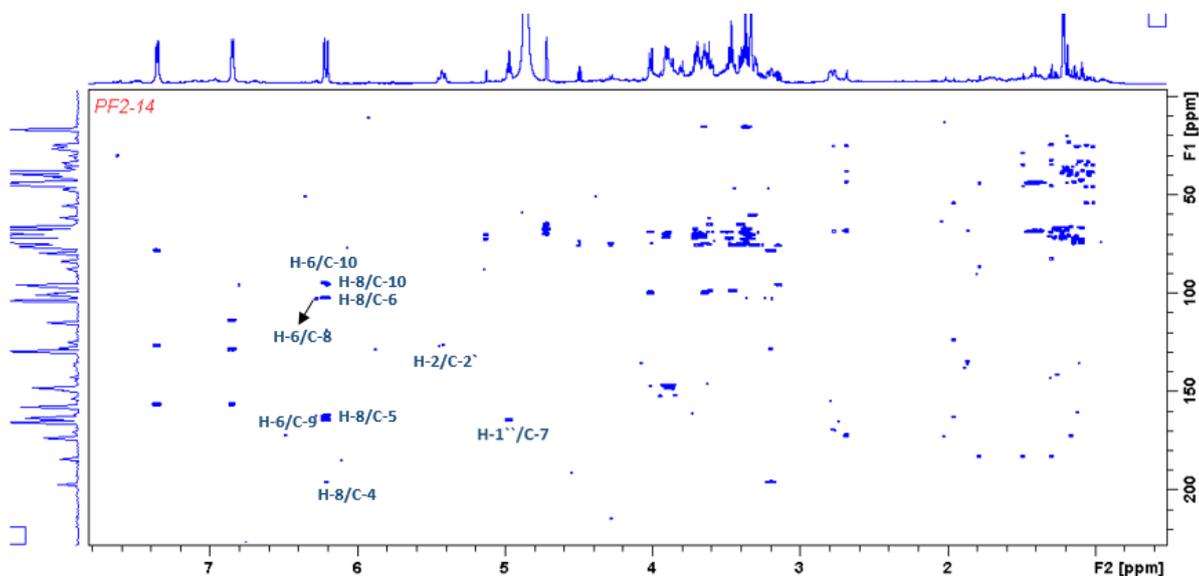


Figure 3.130: The HMBC correlation compound **51**

3.2.9.13 Structure elucidation of compound **marcitrus (119)**

This compound **119** was isolated as a yellow golden oil. The HRESIMS (Figure 3.131) suggested the empirical formula as $C_{28}H_{32}O_{16}$ and in the positive ion mode it showed $[M+H]^+$ peak at m/z 625.1761 (calculated 625.1768). The 1H NMR spectrum (Figure 3.132, Table 3.26) exhibited olefinic protons signals at δ_H 7.47 (2H, d, 8.76 Hz), 7.16 (2H, d, 8.76 Hz) and two anomeric protons at δ_H 4.96 (1H, d, $J=7.00$ Hz), 4.71 (1H, d, $J=1.38$ Hz) and singlet proton at δ_H 6.22 which integrated as 2H and so could be a dioxolane ring. The first suggestion of compound **119** was a glycosylated apigenin connected to a rutosyl moiety with a dioxolane ring. The ^{13}C NMR spectrum (Figure 3.133, Table 3.26) revealed a total of twenty eight carbons including a carbonyl at δ_C 203.6, twelve carbons for rutosyl, seven olefinic carbons and the remains being quaternary carbons while the HSQC (Figure 3.134) showed the direct connections from protons to carbons. The COSY spectrum (Figure 3.135) showed the correlation from H-2' and H-6' to H-3' and H-5' while H-2 revealed a correlation to H-3. In the HMBC experiment (Figure 3.136) 3J correlations were observed from the anomeric proton at δ_H (4.71, H-1'') to carbon at δ_C 69.9 (C-6'') that confirmed the connection between the two sugar moieties. The protons at δ_H 4.96 (H-1'), 7.47 (H-2', H-6') and 7.16

(H-3', H-5') presented a long range HMBC correlation to the carbon at δ_C 160.39 (C-4') confirming the connection of rutinosyl moiety to C-4'. Moreover, the analysis by HMBC showed the correlation from the proton at δ_H 6.22 to the carbon at δ_C 147.7 (C-5) and 135.4 (C-7) that confirmed the position of the dioxolane ring. All these observations confirmed the structure of the compound **119** to be 4,5-dihydroxy-7-(4-(((2*R*,4*S*,5*R*)-3,4,5-trihydroxy-6-(((2*R*,4*R*,5*S*,6*S*)-3,4,5-trihydroxy-6-methyltetrahydro-2*H*-pyran-2-yl)oxy)methyl)tetrahydro-2*H*-pyran-2-yl)oxy)phenyl)-7,8-dihydro-9*H*-[1,3]dioxolo[4,5-*f*]chromen-9-one. This is a new natural product, for which we propose the name marcitrus.

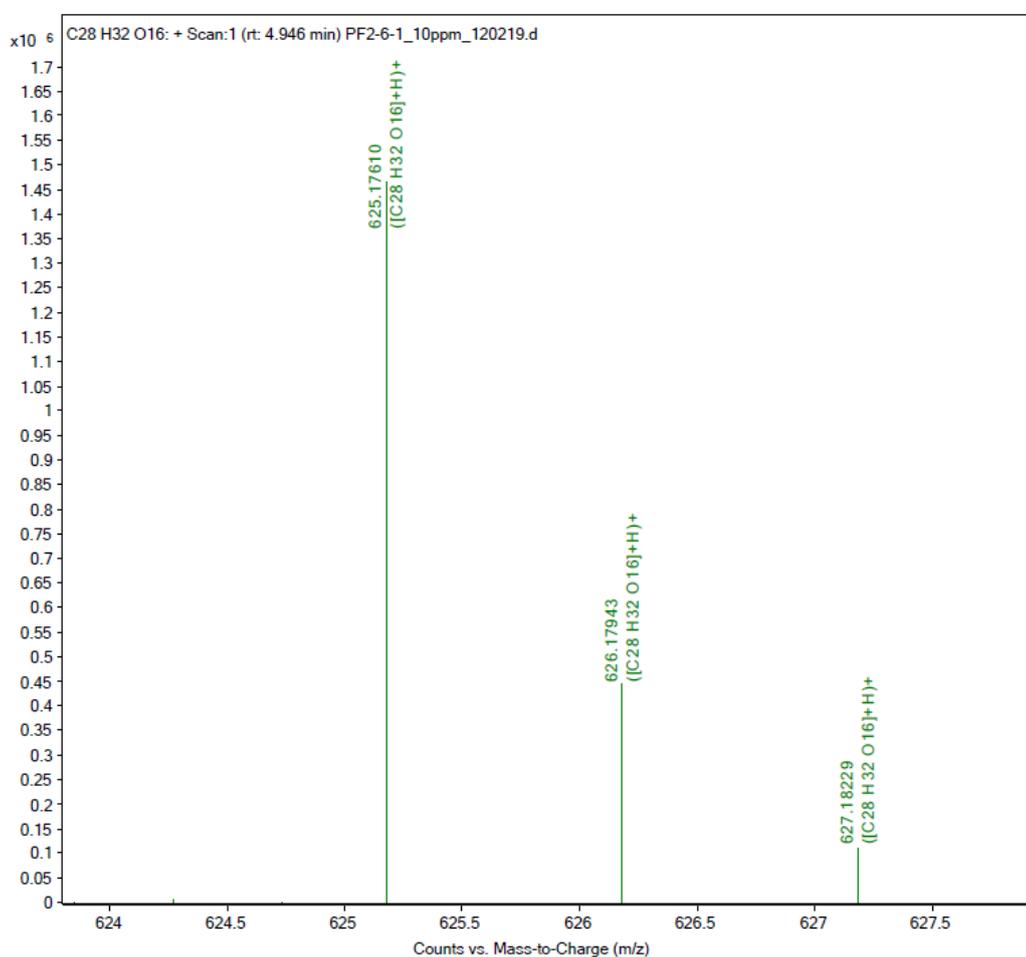


Figure 3.131: The HRESIMS spectrum of compound **119**

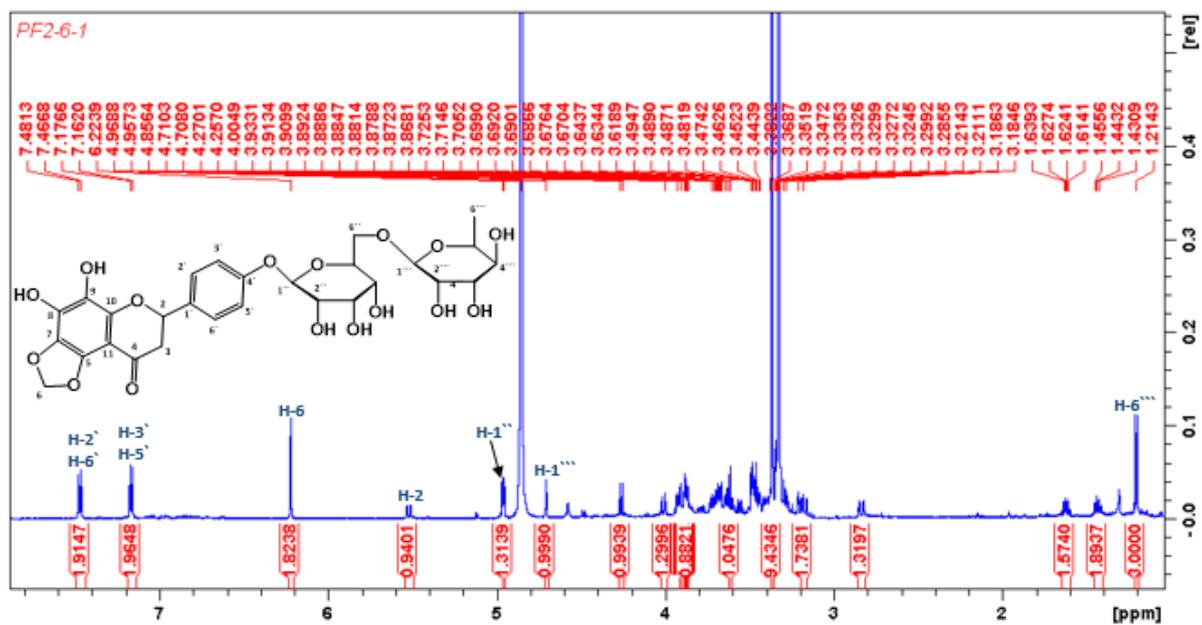


Figure 3.132: ^1H NMR (600 MHz, CD_3OD) spectrum of compound 119

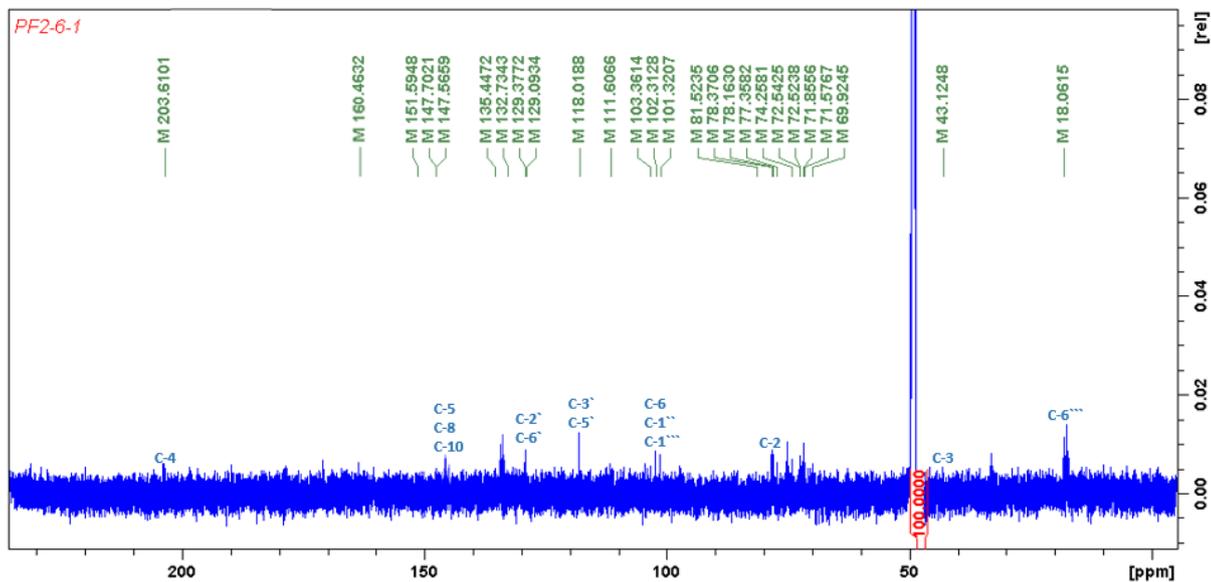


Figure 3.133: ^{13}C NMR (150 MHz, CD_3OD) spectrum of compound 119

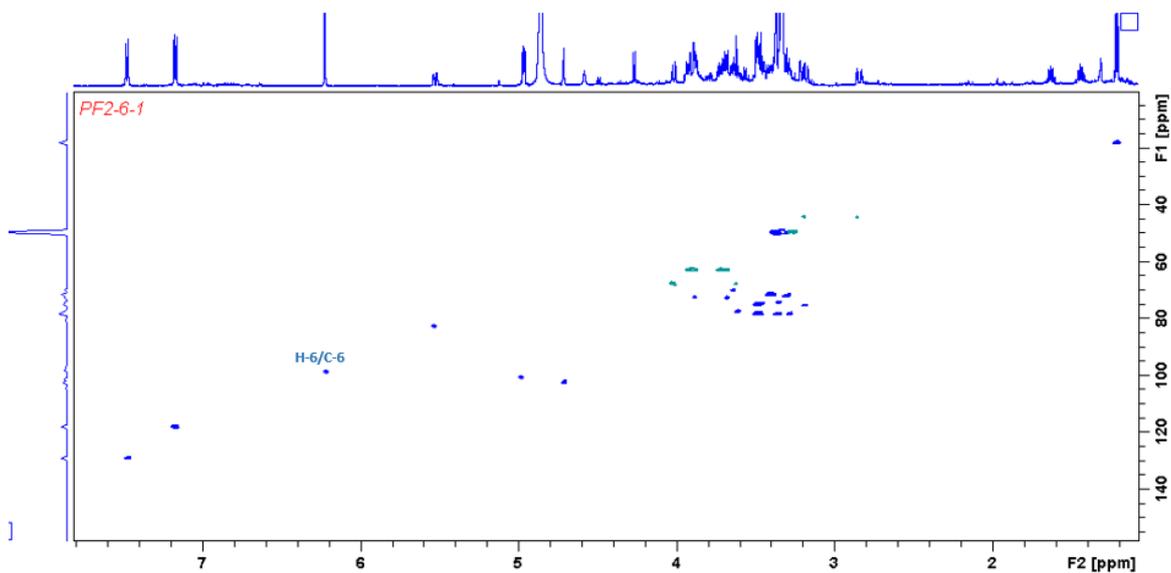


Figure 3.134: HSQC spectrum of compound 119

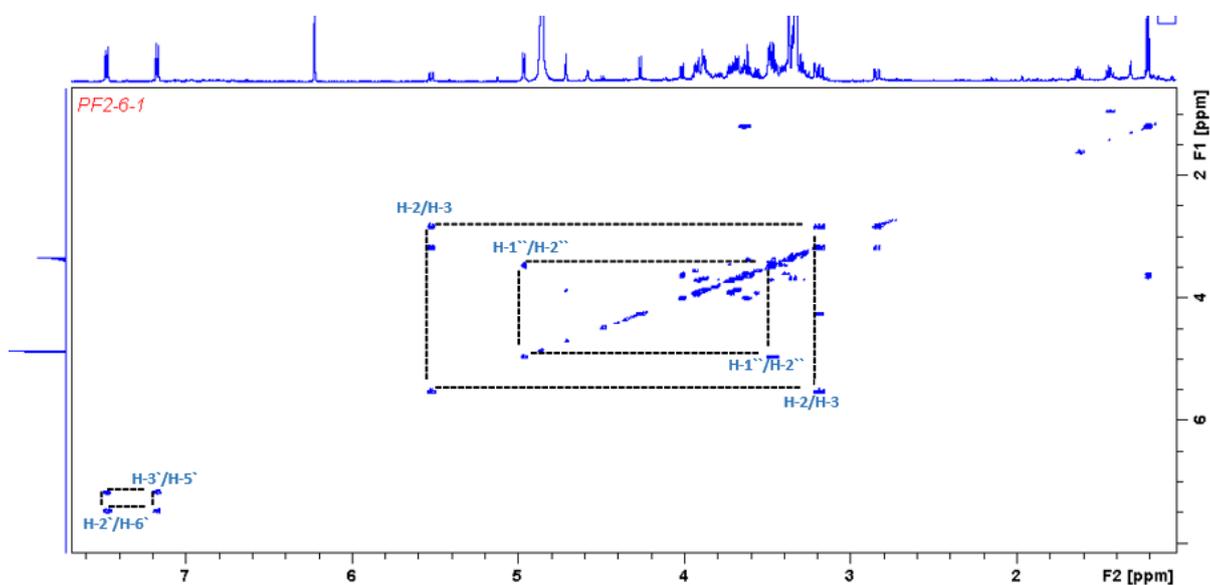


Figure 3.135: COSY spectrum of compound 119

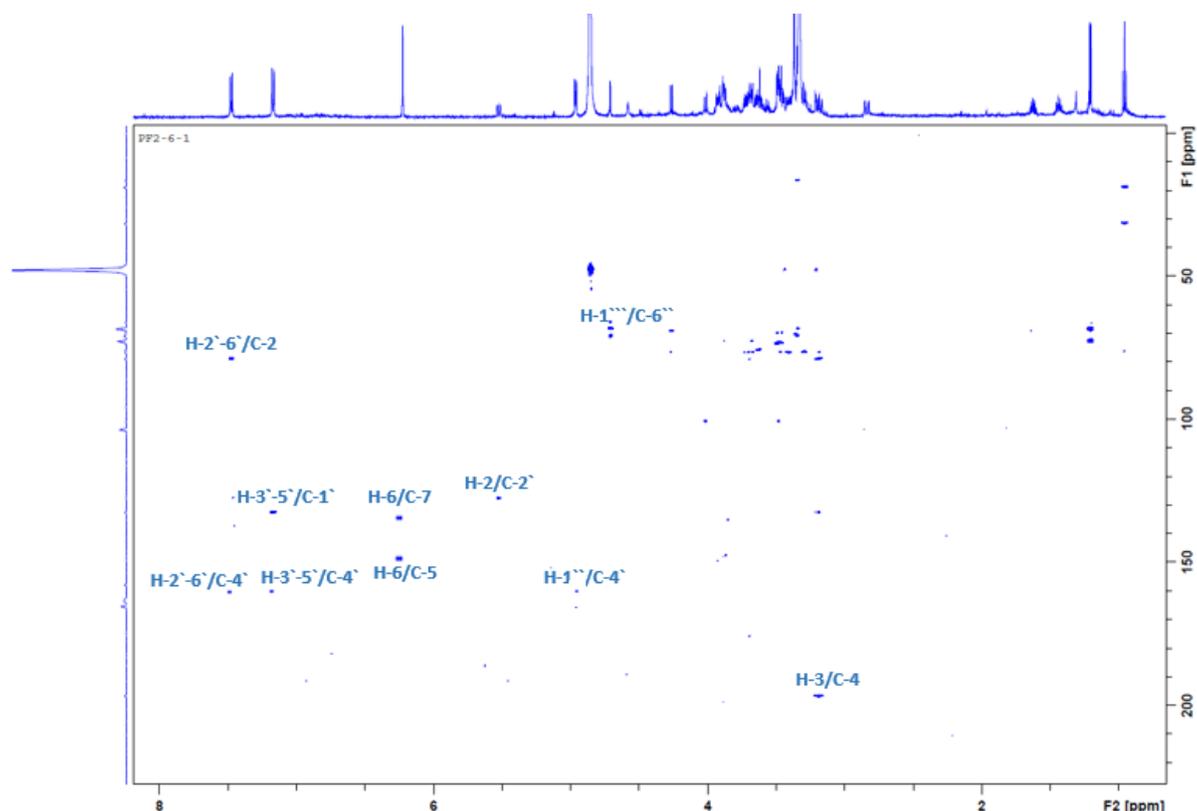


Figure 3.136: HMBC spectrum of compound **119**

3.2.10 Antimicrobial activity of the isolated compounds from *Citrus* species

To evaluate the antimicrobial activity of the compounds isolated from fractions and extracts of *C. grandis* and *C. sinensis*, which showed a notable antimicrobial effect, the isolated compounds were tested against *S. aureus* (NCTC 12981), *E. coli* (NCTC 12241), *P. aeruginosa* (NCTC 12903), *M. luteus* (NCTC 7508) and *C. albicans* (ATCC 90028) and the highest test concentration of compounds was 1 mg/mL using microtitre assay to determine their minimum inhibition concentration (MIC). The observed MIC values are summarized in Table 3.27. The results revealed all the tested compounds except compound **50** inhibited *M. luteus* with highest MIC of 1.25×10^{-1} given by compounds **116** and **119**, while ten compounds efficiently hampering the growth of *C. albicans* and seven compounds exhibited a moderate inhibitory effect against *P. aeruginosa*.

Table 3.26: ¹H NMR (600 MHz) and ¹³C NMR data of compounds **50**, **51**, **119**

Position	Chemical Shift δ_H (ppm), <i>J</i> in Hz			Chemical Shift δ_C (ppm), <i>J</i> in Hz		
	50	51	119	50	51	119
2	5.50 <i>dd</i> (12.00,3.06)	5.42 <i>dd</i> (12.36, 3.06)	5.53 <i>dd</i> (3.12, 12.42)	78.8	78.0	81.5
3	2.78 <i>dd</i> (3.06, 17.16)	2.77 <i>dd</i> (3.06,17.16) 3.29 <i>dd</i> (17.16, 12.36)	2.83 <i>dd</i> (3.12,17.1) 3.29 <i>dd</i> (17.1, 12.42)	42.5	42.3	43.1
4	-	-	-	197.5	198.7	203.6
5	-	-	-	163.5	165.1	147.7
6	6.14 <i>d</i> (2.5)	6.19 <i>d</i> (2.00)	6.22 <i>s</i>	96.8	98.1	101.3
7	-	-	-	165.6	167.0	135.4
8	6.12 <i>d</i> (2.5)	6.22 <i>d</i> (2.00)	-	96.00	97.3	147.6
9	-	-	-	163.0	164.7	129.4
10	-	-	-	103.8	105.1	151.6
11	-	-	-	-	-	111.6
1'	-	-	-	131.37	131.0	132.7
2'	6.93 <i>d</i> (2.1)	7.36 <i>d</i> (8.52)	7.47 <i>d</i> (8.76)	114.6	129.3	129.1
3'	-	6.85 <i>d</i> (8.52)	7.16 <i>d</i> (8.76)	146.9	116.5	118.0
4'	-	-	-	148.4	149.2	160.4
5'	6.95 <i>d</i> (8.34)	6.85 <i>d</i> (8.52)	7.16 <i>d</i> (8.76)	112.5	116.5	118.0
6'	6.91 <i>dd</i> (8.34, 2.1)	7.36 <i>d</i> (8.52)	7.47 <i>d</i> (8.76)	118.4	129.3	129.1
1''	4.98 <i>d</i> (8.5)	4.96 <i>d</i> (7.02)	4.96 <i>d</i> (7.00)	101.1	101.3	102.3
2''	3.63*	3.36*	3.88	76.0	72.5	75.5
3''	3.45*	3.60*	3.48	78.8	78.0	78.5
4''	3.40*	3.90*	3.39	70.7	71.4	71.6
5''	3.54*	3.46*	3.34	76.7	80.8	78.4
6''	3.42 <i>m</i>	4.01*	3.63	66.8	69.9	69.9
1'''	4.52 <i>d</i> (1.92)	4.71 <i>d</i> (2.1)	4.71 <i>d</i> (1.38)	103.8	102.3	103.4
2'''	3.27*	3.65*	3.48	71.7	74.8	74.2
3'''	3.28*	3.33*	3.29	72.5	72.2	71.8
4'''	3.18*	3.29*	3.69	73.4	74.3	72.5
5'''	3.38*	3.65*	3.6	68.8	77.3	77.3
6'''	1.09 <i>d</i> (6.18)	1.21 <i>d</i> (6.24)	1.21 <i>d</i> (6.24)	18.3	18.0	18.1
3`- OCH ₃	3.78 <i>s</i>	-	-	56.2	-	-

* Overlapped peaks – confirmed from COSY, HSQC and HMBC experiments

Compound **111** showed an extraordinary effect on the growth of *S. aureus*, *E. coli*, *M. luteus* and *C. albicans* with highest MIC 6.25×10^{-2} mg/mL. However, compound **50** failed to have any effect against the tested organisms. The modified microtitre assay was the biological test used in this study to examine the antimicrobial activity of the compounds and determine their MIC values. Various antibiotics outlined on Table 3.3 were used as a positive control and the negative controls were water, nutrient broth and normal saline.

Table 3.27: The MIC (mg/mL) of isolated compounds from *Citrus*

Compounds	All concentrations in (mg/mL)				
	<i>S. aureus</i>	<i>E. coli</i>	<i>P. aeruginosa</i>	<i>M. luteus</i>	<i>C. albicans</i>
50	N/A	N/A	N/A	N/A	N/A
57	N/A	N/A	2.5×10^{-1}	5×10^{-1}	2.5×10^{-1}
110	2.5×10^{-1}	N/A	5×10^{-1}	5×10^{-1}	2.5×10^{-1}
111	2.5×10^{-1}	5×10^{-1}	2.5×10^{-1}	2.5×10^{-1}	6.25×10^{-2}
112	N/A	N/A	5×10^{-1}	2.5×10^{-1}	2.5×10^{-1}
113	N/A	N/A	5×10^{-1}	2.5×10^{-1}	2.5×10^{-1}
114	N/A	N/A	5×10^{-1}	2.5×10^{-1}	5×10^{-1}
115	N/A	N/A	N/A	5×10^{-1}	N/A
116	N/A	N/A	5×10^{-1}	1.25×10^{-1}	2.5×10^{-1}
117	N/A	N/A	N/A	5×10^{-1}	5×10^{-1}
118	N/A	N/A	N/A	2.5×10^{-1}	5×10^{-1}
119	N/A	N/A	N/A	1.25×10^{-1}	2.5×10^{-1}

*N/A: No Activity

3.2.11 Discussion

Citrus is one of the most interesting pharmaceutical genera of the Rutaceae family. Many studies have documented this genus to contain different groups of secondary metabolites (Benavente-Garcia *et al.*, 1997; Nijveldt *et al.*, 2000; Encyclopedia Britannica, 2019).

One of the largest classes of natural products found in *Citrus* is flavonoids, which are considered as polyphenolic compounds with structural skeletons of flavanone, flavone, and/or flavonol (Široká *et al.*, 2013). Flavonoids have been studied and reported for many

healthf be refiling properties such as, antimicrobial, anti-inflammatory, antioxidant, cardiovascular and anticarcinogenic (Nijveldt *et al.*, 2001).

The present work focuses on two important species of this genus *C. grandis* and *C. sinensis* using different parts for phytochemical study and screening of their antimicrobial properties on two Gram-positive bacterial strains *Micrococcus luteus* (NCTC 7508) and *Staphylococcus aureus* (NCTC 12981), and two Gram-negative strains *Escherichia coli* (NCTC 12241) and *Pseudomonas aeruginosa* (NCTC 12903) and the only fungal strain *Candida albicans* (ATCC 90028), which is considered one of the most common pathogenic fungi.

The result of preliminary antimicrobial screening of *C. grandis* leaves extracts showed the most potent activity with an MIC value of 3.125×10^{-1} mg/mL against *P. aeruginosa* and *M. luteus*. Bioassay-guided chemical investigation led to the isolation of six compounds, **50**, **57**, **111**, **112**, **114** and **115**. Compounds **111** showed significant antimicrobial activity against the microbes tested. To the best of our knowledge, this is the first report on antimicrobial activity of Iraqi *C. grandis* leaves.

The primarily antimicrobial screening for the peel and leaves of *C. sinensis* extracts against the microbes used in this work revealed that the DCM extract of *C. sinensis* peel showed the strongest higher level of activity with MIC 4.88×10^{-3} mg/mL against the bacterium *M. luteus*. Moreover, the most active extract in *C. sinensis* leaves was the DCM extract, which presented a significant MIC 3.12×10^{-1} against *M. luteus* and *C. albicans*. Both chromatographic and spectroscopic techniques led to isolation and identification of compounds **50**, **57**, **112**, **114**, **115** from the DCM extract of *C. sinensis* leaves and **48**, **50**, **51**, **57**, **102**, **110**, **111**, **113**, **116**, **117**, **119** from DCM and methanolic extracts of *C. sinensis* peels.

The analysis of the antimicrobial results for the isolated compounds revealed that the most effective compound was **111**, which is a flavonoid substituted with five oxygenated

methyls. According to Tian-Yang Wang *et al.* (2018), the most active antimicrobial flavonoids have double bond on C2=C3 and are substituted on position 5 and 7 by OH. Seemingly, occupation of positions 5, 6, 7 by OCH₃ could offer notable inhibitory activity. Compound **50** was not active with the maximum concentration was used in this assay. However, this compound was reported to possess major antiviral effect, anti-inflammatory and analgesic activity (Kumar and Pandey, 2013).

The number of the hydroxyl group results is another influencing factor. Increasing hydroxyl group numbers led to reducing polarity of the compound, which explained inactivity on biological membranes.

The primary testing to determine the mechanism of action for the tested compounds revealed all compounds tested were bacteriostatic.

Chapter 4

The Anti-MRSA Activity of Isolated Compounds

4.1 Methicillin-resistant *Staphylococcus aureus* (MRSA)

Methicillin-resistant *Staphylococcus aureus* (MRSA) is a human pathogen, which is resistant to several types of antibiotics. The organism can be isolated from around a third of the population without significant symptoms (Otto, 2013). The infection levels with MRSA vary from moderate to serious and sometimes leads to death (Lowy, 2003; Moellering Jr, 2011; Otto, 2013).

Staphylococci have been found on earth for more than a billion years depending on many historical indicators (Moellering Jr, 2011). In the 19th century, it was recorded as pathogenic, because it was identified as the main reason for wound infections. Most civilisations successfully treated infections caused by this bacterium using different agents such as honey, copper salts and myrrh. The discovery of penicillin from *Penicillium notatum* in 1928 provided a good new way to inhibit *S. aureus* especially in the 1940s, when penicillin was introduced presented as medication from *S. aureus*. Unfortunately, in 1942s *S. aureus* developed penicillin resistance.

The basic molecular structure of penicillin is a β -lactam ring, which inhibits the synthesis of the bacterial peptidoglycan cell wall. On the other hand, *S. aureus* produces the enzyme β -lactamase, which is considered as the main reason of inactivation of penicillin by attacking the β -lactam ring and changing it to open chain (Figure 4.1) (Palavecino, 2007; Moellering Jr, 2011; Kuriyama *et al.*, 2014; Elhassan *et al.*, 2015).

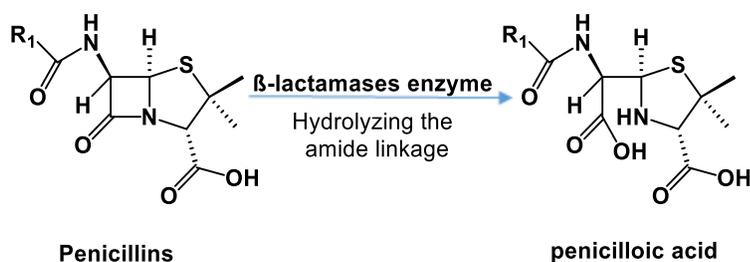


Figure 4.1: Inactivation of penicillin by β -lactamase enzymes

In 1959, methicillin (Figure 4.2) became available as an alternative to penicillin. This antibiotic initially showed significant inhibitory activity against penicillin-resistant *Staphylococcus aureus*. After two years of clinical use, however, the first MRSA strain was detected in the United Kingdom (Jevons, 1961). A short time later, the resistant strain was isolated from other European countries and subsequently from Japan, Australia, and the United States of America (Enright *et al.*, 2002; Moellering Jr, 2011). Methicillin resistance is due to the presence of the *mecA* gene in conjunction with other genetic factors. The *mecA* gene encodes a unique penicillin-binding protein, designated 2' (PB2') or 2a (PBP2a), that has reduced affinity to β -lactam antibiotics (Tsubakishita *et al.*, 2010).

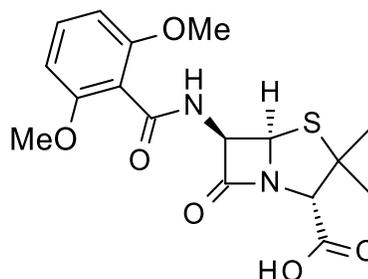


Figure 4.2: Structure of methicillin

In the 1970s, North America, some European countries and two hospitals (East Anglia and Kettering) in England suffered from outbreaks of MRSA. The Kettering outbreak affected most hospitals in the UK by an epidemic of MRSA (Cox *et al.*, 1995; Newsom, 2004). In the 1980s, the MRSA had spread globally. Between 1999 to 2000, in the USA, around 125,969 incidences of hospitalisations were identified with MRSA infections. Furthermore, between January 1998 and June 2003, the annual average percentage of *S. aureus* isolates that were MRSA increased further to 51.6% of ICU and 42% of inpatient non-ICU *S. aureus* isolates (System, 2004; David and Daum, 2010).

It has been reported that the percentage of MRSA infections in children under 15 years of age in England was 0.9% in 1990 and 13% in 2000 (Khairulddin *et al.*, 2004). In Sweden, the general infections were 325 in 2000 and raised in 2003 to 544 (Stenheim *et al.*, 2006). The Data of National Statistics in England and Wales recorded that MRSA infections caused the death of 364 individuals in 2011 compared to 485 in 2010 (Kyte, 2011). The number of infections decreased from 2013 to 2016 (CDC, 2019). A total of 12,784 *Staphylococcus aureus* infections were reported through both the MRSA bacteraemia and methicillin-susceptible *S. aureus* (MSSA) bacteraemia surveillance schemes (Public Health England, 2018). The figure represents a 3.7% increase in the numbers of bacteraemias caused by *S. aureus* from 2016-17 (total number 12,324) and a staggering 29.4% increase from 2011-12 (total number 9,883) when MSSA reporting was made mandatory. However, of the *S. aureus* infections reported during 2017-18, only about 6.6% were MRSA cases, which was a 41.4% decrease from 2011/12, in which 11.3% of all reported cases of *S. aureus* infections were caused by MRSA, and a 1.1% decrease from 2016/17 in which the figure was 6.7%. Whilst the actual reasons behind this continuous decline in MRSA cases are not absolutely clear, it can be assumed that there may be many factors that are behind this decline in MRSA infection ratio such as developed hospital practices including preferred cleaning and expanded regard for the significance of sterile medications for open injuries. Nevertheless, it is necessary to develop new antibiotics to tackle against the issues of antibiotic resistance including MRSA.

4.2 Types of MRSA

Two main types of MRSA have been identified. These are community-associated MRSA (CA-MRSA) and health care-associated MRSA (HA-MRSA).

4.2.1 Community-associated MRSA (CA-MRSA)

Healthy people who have never attended hospitals can become infected with MRSA. It has been reported that the cause of such infections was by coming in contact with people suffering from MRSA especially when sharing household facilities. CA-MRSA is the main cause of skin infections and septicemia.

4.2.2 Healthcare-associated MRSA (HA-MRSA)

Healthcare-associated infections, also known as nosocomial infections, are infections that are acquired in a hospital or other healthcare facility. The HA-MRSA is caused mainly because of lack of nosocomial factors in hospital facilities or as a result for medical or surgical treatments. This type of MRSA infection poses a serious risk to patients, staff and visitors. Moreover, the treatment requires significant costs and long time to heal (Miller *et al.*, 2011; NHS, 2017).

4.3 Treatment of MRSA infections

It is a big challenge to treat MRSA infections due to resistance factors. Many antibiotics have been used for the treatment of MRSA infections. These include trimethoprim-sulfamethoxazole, clindamycin, minocycline, fluoroquinolones, linezolid, daptomycin, tigecycline, ceftaroline and doxycycline. Recovery time, drug dosage, location of the infection and course of treatment depend on a patient's condition and the severity of the infection (Gould *et al.*, 2012).

4.4 Natural products with potent anti-MRSA activity

Natural products are considered potential sources for novel medications, including remedies for MRSA infections. Many studies documented the use of different plants with potential activity against MRSA. One study, for example, demonstrated a significant inhibitory

effect of an essential oil from *Thymus vulgaris* against MRSA ATCC33592 (Tohidpour *et al.*, 2010). Brazilian plants have been shown to have inhibitory activity against MRSA ATCC 33591 (Machado *et al.*, 2003). *Althaea officinalis*, *Melissa officinalis*, *Mentha longifolia* and *Rosa damascene* identified from the Palestinian flora showed remarkable anti-MRSA effects (Abu-Shanab *et al.*, 2007). Traditional Ghana medicinal plants have been used successfully as anti-MRSA agents (Pesewu *et al.*, 2008). Seven Nigerian plants, *Acalypha wilkesiana*, *Ageratum conyzoides*, *Bridella ferruginea*, *Terminalia avicennioides*, *Phyllanthus discoideus* and *Ocimum gratissimum*, proved to have inhibitory effects on *S. aureus* strain (MRSA-NCIB8588) (Akinyemi *et al.*, 2005). Many compounds derived from *Desmodium caudatum* had notable anti-MRSA activity (Sasaki *et al.*, 2012). Xanthones, anthraquinones, quinones, calcarides, aflatoxins and comazaphilones were shown to have significant anti-MRSA effects (Xu *et al.*, 2015). Many compounds isolated from marine bacteria and fungi are also effective anti-MRSA agents (Debbab *et al.*, 2010). Compounds derived from a marine isolate of the fungus, *Pseudallescheria* showed potent anti-MRSA activity (Li *et al.*, 2006).

4.5 Materials and methods

4.5.1 Tested materials

A total of twenty compounds have been tested for anti MRSA activity in this study. These include thirteen compounds from *Ruta chalepensis*, namely, arborinine (**107**), bergapten (**43**), chalepin (**45**), chalepentin (**46**), γ -fagarine (**42**), graveoline (**105**), 6-hydroxy-rutin 3'-7-dimethyl ether (**103**), imperatorin (**104**), isopimpinellin (**44**), kokusaginine (**99**), rutamarin (**47**), rutin (**48**) and rutin 3'-methyl ether (**101**) (Figure 3.5). The compounds isolated from the *Citrus* species, which were tested for anti-MRSA activity, include demethylnobiletin (**113**), hesperidin (**50**), 3-methoxynobiletin (**110**), nobiletin (**57**), salvigenin (**118**), sinensetin (**111**) and tangeritin (**116**) (Figure 3.80). The isolation procedures for these compounds have been described in

Chapter 2 (2.3.3).

4.5.2 Resistant strains of *Staphylococcus aureus*

Five methicillin-resistant *S. aureus* strains (SA1199B, XU212, MRSA340702, EMRSA-15MRSA274819) and standard strain (ATCC25923) were used in this study. All the bacterial strains were obtained from the UCL School of Pharmacy and the experiments were performed in the University of East London.

4.5.3 Materials used for the MRSA assay

A summary of all of the materials, which were used in the anti-MRSA assay is presented in Table 4.1

Table 4.1: Materials used in MRSA assay

Chemical and reagents	Supplier
Ca ²⁺ , Mg ²⁺	Sigma-Aldrich Gillingham, UK
Mueller-Hinton broth	Oxoid
Normal saline	Sigma-Aldrich Company Ltd., UK
3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide	Sigma-Aldrich Company Ltd., UK
DMSO	Sigma-Aldrich Company Ltd., UK
Norfloxacin	Sigma

4.5.4 The anti-MRSA assay

Preparation of culture medium: Mueller-Hinton broth (MHB) was prepared according to the instruction given by the supplier. The MHB was adjusted to contain cations- 20 mg/L Ca²⁺ and 10 mg/L of Mg²⁺.

Preparation of tested compounds: Compounds and antibiotics were dissolved in predetermined amounts of DMSO (less than 1% concentration in the final well), which was further diluted with MHB to obtain the targeted starting concentration (128 µg/mL).

Preparation of MTT: It was achieved by dissolving the required amount of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide in methanol to obtain a concentration of 5 µg/mL.

Suspension of subculture of the bacteria: All bacterial strains were subcultured one day before the experiment. Bacterial strains were subcultured in nutrient agar slopes by streaking the bacteria with a loop followed by incubation at 37° for 12-18 h.

Anti-MRSA assay: This experiment was performed using 96-well plates. The first step on this assay was to add (100 µL) of MHB to all wells except those in column 12. To the first row of the plate, 100 µL of test compounds or antibiotic were added. Using a multi-channel pipette, the materials of the first well were mixed properly, followed by the transfer of 100 µL of the well contents to the wells of the second column and so on until column 10. Finally, 100 µL content from the wells of column 10 were transferred to the wells of column 12.

An inoculum density of 5×10^5 cfu of each of the test organisms was prepared in normal saline (9 g/L) by comparison with a 0.5 MacFarland standard. MHB (125 µL) was dispensed into 10 wells of a 96 well microtitre plate (Nunc, 0.3 mL volume per well). Then microtitre plates were incubated at 37°C for 18 h. To determine the minimum inhibitory concentrations (MICs), 20 µL of MTT was add to the microtitre plate and incubated for 20 min. It is colourimetric method, bacterial growth was indicated by the colour changing from yellow to dark blue and the MIC was recorded as the lowest concentration at which no growth was observed. As a positive control norfloxacin, a well-known antibiotic was used.

The method was used to determine the MIC considered as a broth microdilution method according to National Committee for Clinical Laboratory Standards with modification using nutrient broth as the medium (Rahman *et al.*, 2008; Shiu *et al.*, 2011; Ioannou *et al.*, 2012; Nurunnabi *et al.*, 2018; Rahman *et al.*, 2018; Tareq *et al.*, 2018).

4.6 Results and discussion

Twenty isolated compounds from *R. chalepensis*, *Citrus grandis* and *C. sinensis* were tested for activity against six MRSA strains (Table 4.2). The results revealed that fifteen of them inhibited the MRSA strains with different MIC values. Arborinine (**107**), chalepentin (**46**), 6-hydroxy-rutin 3'-7-dimethyl ether (**103**) and sinensetin (**111**) were the most active against all MRSA strains tested.

In spite of the fact that chalepin (**45**), chalepentin (**46**) and rutamarin (**47**) (Figure 4.3) are all furanocoumarin derivatives, they caused different levels of inhibitions because of structural differences. The order of anti-MRSA potency in these compounds was **46** > **47** > **45** (Table 4.2). Functional groups were the main differences among these three compounds contributing to their differences in lipophilicity. All three compounds are 3-substituted furanocoumarins, among which, except for **46**, the other two compounds are dihydrofuranocoumarins. Rutamarin (**47**), which is simply the acetylated product of chalepin (**45**) was more active than **45**, presumably because of more lipophilicity caused by acetylation.

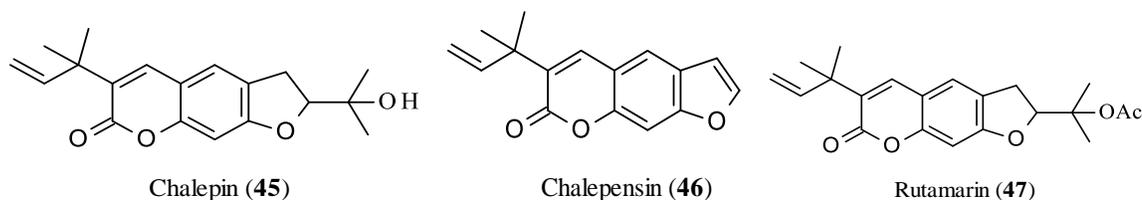


Figure 4.3: Structural similarities and differences in anti-MRSA furanocoumarins **45-47**

6-Hydroxy-rutin 3'-7-dimethyl ether (**103**), rutin (**48**) and rutin 3'-methyl ether (**101**) are flavonoid glycosides, having only differences in the presence/absence and in the number of methyl ether groups in them, giving them varying degrees of lipophilicity (Figure 4.4). Rutin (**48**) does not contain an OMe group, while compound **101** has an OMe group on 3' position, and **103** has two OMe groups in positions 3' and 7. In addition, in **103**, position 6 is occupied

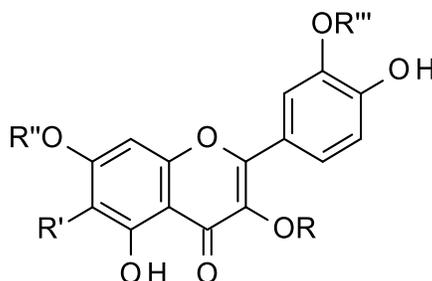
by a hydroxyl group. The highest anti-MRSA potency of compound **103** may be because of the different functional groups and their unique positions that make this compound as the most lipophilic among these three compounds. The order of anti-MRSA selectivity in these compounds was **103** > **101** > **48** (Table 4.2).

Table 4.2: Anti-MRSA activity of selected compounds isolated in this study

Compounds	MIC values in µg/mL					
	XU212	ATCC 25923	SA1199B	EMRSA-15	MRSA340702	MRSA274819
42	-	-	-	-	-	-
44	-	-	-	-	-	-
43	-	-	-	-	-	-
45	256	-	256	-	256	256
46	64	128	-	-	64	64
47	128	-	128	-	128	128
48	-	-	-	-	-	256
50	-	-	-	-	-	-
57	-	256	-	-	-	-
99	-	-	-	-	-	256
101	256	128	-	-	256	256
103	32	64	-	-	128	256
104	-	256	256	-	-	128
105	-	-	-	-	-	-
107	-	256	-	128	64	256
110	-	256	-	256	-	-
111	-	128	-	256	128	256
113	-	256	-	256	-	-
116	-	-	-	-	-	-
118	-	-	-	-	-	-
Norfloxacin	16	2	32	1	64	64

Compound **107** is an acridone alkaloid containing three methyl groups, two of which are oxygenated. This compound has been reported to have many pharmaceutical applications as antimicrobial, antiviral, antiplasmodial, antimalarial and anticancer agents and,

not surprisingly, as anti-MRSA agent (Sohrab *et al.*, 2004; Réthy *et al.*, 2007; Oliveira *et al.*, 2009; Amoa Onguéné *et al.*, 2013; Fouotsa *et al.*, 2013).



Compounds	R	R'	R''	R'''
48	Rutinosyl	H	H	H
101	Rutinosyl	H	H	Me
103	Rutinosyl	OH	Me	Me

Figure 4.4: Structural similarities and differences in anti-MRSA flavonoids **48**, **101** and **103**

Compounds **110**, **111**, **113**, **116** and **118** are flavonoids. Compounds **110** and **111** exhibited three aromatic protons as ABX pattern in positions 2', 5' and 6'. Compound **110** has six oxygenated methyl groups and one free proton in position 6 while, compound **111** contains five oxygenated methyl groups with no substituent at positions 3 and 8. Compounds **116** and **118** are known as a symmetrical ring because it has only one methoxy group in position 4'. Compound **116** consists of five methoxy groups and position 3 is free while, compound **118** contains three methoxy groups, a hydroxyl group in position 5 and two free protons in positions 3 and 8. Among these tested flavonoids, compound **111** showed a distinct effect on the MRSA strains tested. Compounds **42**, **44**, **43**, **50**, **105**, **116** and **118** did not possess any anti-MRSA activity against the strain at the highest concentration used in this assay.

This seems to be the first report on the evaluation of the anti-MRSA effect of the isolated compounds, from *R. chalepensis*, *C. grandis* and *C. sinensis* against a number of MRSA strains.

Chapter 5
**Scanning Electron
Microscopic Analysis**

5.1 What is scanning electron microscope (SEM)

A scanning electron microscope (SEM), a surface imaging tool, is an electron microscope that can generate images of a sample by scanning the surface with a focused beam of accelerated electrons. SEM was introduced to study the topography and composition of solid material, and was first commercially made available in 1965. This microscope consists of an electron optical column, a vacuum system, electronics, and software. The specimen chamber is large and there are lenses above the specimen to focus the beam of electrons onto the specimen surface. At the top of the column an electron gun is located, which produces the electron beam. This beam is scanned in a rectangular area on the specimen. The interaction between the beam of electrons and the specimen generates different signals, which are detected and produce the image. A secondary electron signal is the most commonly used signal and provides topographic information of the specimen surface (Hearle *et al.*, 1972; Lawes, 1987; Goldstein *et al.*, 2017).

5.2 Applications of scanning electron microscope

SEM is used in a number of industrial, commercial, research and forensic applications, some of which are outlined below.

5.2.1 Industrial uses

SEM is a useful instrument used to examine the surface structure of components and products in different industrial sectors such as microelectronics, semiconductors, medical devices and food processing. Moreover, SEM is considered as an important tool in the paper industry to study the paper strength which is generally affected by fibre quality, fibre bonding and network factors (Li, 2002). One of the new trends in applications of SEM is in the cosmetics industry, where it can provide information about the shape and size of tiny particles in cosmetic products before any product reaches the consumer. For instance, the small and round particles

give better impact on the consistency and performance of the product in mixing and flowing, than the overlarge or serrated particles (ATA, 2018)

5.2.2 Commercial

The industrialization or development of products required to study the composition and topography of products. For example, for optimal performance, stainless steel equally coated with specific chemicals. The SEM is considered as a helpful tool to detect any cracks, imperfections or contaminants on the surfaces of such coated products (ATA, 2018).

5.2.3 Research

The SEM has helped scientific research in universities in different fields. For example, biologists use SEM to gain an increased level of information on various biological specimens, and geologists apply this technique to learn more about crystalline structures, soil and rock. Moreover, SEM is considered as one of the fundamental instruments to study the nanoparticles in pharmaceutical research, e.g., in drug delivery, in dental studies, and in biomedical research. Additionally, the SEM has been used in food science, environmental studies, and forensic research (Carr, 1971; ATA, 2018).

5.3 The use of the scanning electron microscopy (SEM) in microbiology

This instrument can provide information about the micromorphology of microorganisms (Kaláb *et al.*, 2008) in spite of the fact that, very limited research has been conducted to study the microorganisms using the SEM (Carr, 1971; Marrie *et al.*, 1983; Franson *et al.*, 1984; Arroyo *et al.*, 2014). One of the earliest SEM research studies on bacteria (Williams and Davies, 1967) was performed at the University of Liverpool where they recommended the use of the SEM to study the morphological features of microorganisms. Greenwood and O'Grady (1969) studied the effect of ampicillin on the surface of bacteria *Staphylococcus aureus* and *Streptococcus*

pyogenes. They found that the antibiotics had an effect on cell division of the bacteria and caused significant damage to the cell wall.

In the present study, the effect of many pure compounds, which had shown significant antimicrobial activity, was investigated by scanning electron microscopy.

5.4 Materials and methods

The isolation and antimicrobial activity of the purified compounds are described in *Chapters 2 and 3*. Six compounds, chalepentin (**46**), nobiletin (**57**), 6-hydroxy-rutin-3'-7-dimethyl ether (**103**), imperatorin (**104**), sinensetin (**111**) and arborinine (**107**) were tested. The effect of the pure compounds on the morphology of two Gram-positive bacteria, *Micrococcus luteus* and *Staphylococcus aureus*, two Gram-negative bacteria, *Escherichia coli* and *Pseudomonas aeruginosa*, and the fungus, *Candida albicans* was investigated.

The test samples were selected from microtitre plates at the MIC concentrations. 100 μ L of 5% v/v unbuffered glutaraldehyde (Agar Scientific, UK) was added to the target well and fixed at 4°C. After 24 hours fixation in glutaraldehyde, the samples were removed and passed through isopore membrane filters (0.4 micron filters from Merk), washed three times with distilled water and then air dried at room temperature. The filters were attached to aluminium specimen stubs with sticky carbon tabs (Agar Scientific, UK). All samples were sputter-coated with gold. They were then examined with an FEI InspectS scanning electron microscope using a range of operating voltages and working distances.

5.5 Results and discussion

In this study, the SEM was chosen because it is one of the useful tools, which has been used to study the morphological features of microorganisms for viewing the coated samples and detecting any surface damage or change. This study was designed to capture images of untreated

(control) and treated strains of *Escherichia coli*, *Candida albicans*, *Micrococcus luteus*, *Staphylococcus aureus* and *Pseudomonas aeruginosa*.

Candida albicans is one of the important opportunist pathogenic fungi that can cause serious systemic infections. The treatment of this microbe is considered as a big challenge because it is eukaryotic like the host cell (Endo *et al.*, 2010). *C. albicans* is a dimorphic yeast *i.e.* it can grow either as typical budding yeast cells or as hyphae depending on the environmental conditions (Sudbery *et al.*, 2004; Mayer *et al.*, 2013). In reality, it can produce a variety of cell forms and therefore it really should be described as polymorphic. Untreated *C. albicans* under the SEM showed budding cells and hyphae (Figure 5.1). The compounds **46**, **57**, **103**, **111** and **107** caused apparent damage to the surface of *C. albicans* (Figure 5.1), which was clear to detect under the SEM. Compounds **46**, **103** and **111** showed the same action against the surface of *C. albicans*. Some cells, following treatment with compound **57**, possessed particles or small swellings on the surface of the cell wall. Moreover, compound **107** generated a mucous layer around the cells and changed their shape.

The Gram-positive bacterial species, *Micrococcus luteus* has a primary coccoid shape. This organism can cause ‘simple’ infections particularly in individuals with suppressed immune systems (Bonjar, 2004). In the present study, a sample of the untreated strain of *M. luteus* was examined with the SEM (Figure 5.2). The growth of *M. luteus* was significantly inhibited by compounds **46**, **103**, **104**, **107** and **111**. The treated samples were all examined with the SEM (Figure 5.2). The compounds caused changes to the shape of the cells, increased the agglutinability of the cell walls and affected the progress of cell division (Figure 5.2). These effects could explain the possible mechanism of action of the compounds. There have been very few studies on the effects of antibacterial compounds on the cell surface of *M. luteus*

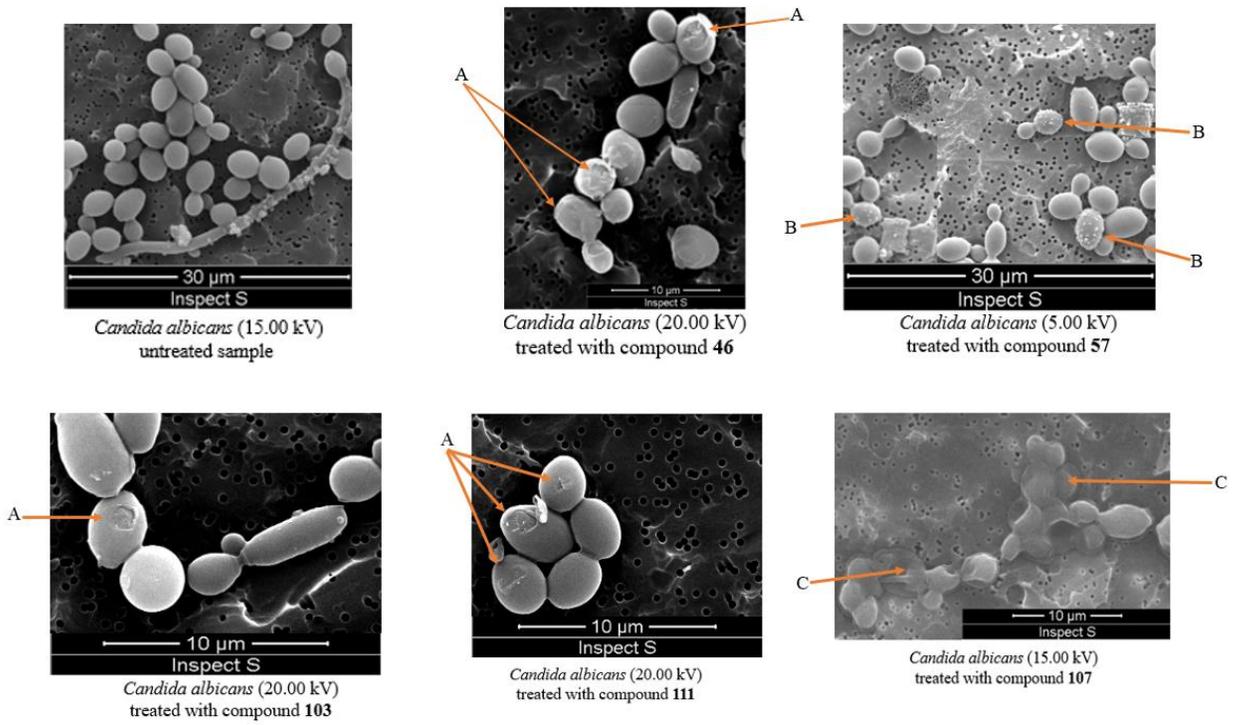


Figure 5.1: Untreated and treated samples of *Candida albicans* strain

A: surface damage; B: surface particles/small swellings; C: mucous layer

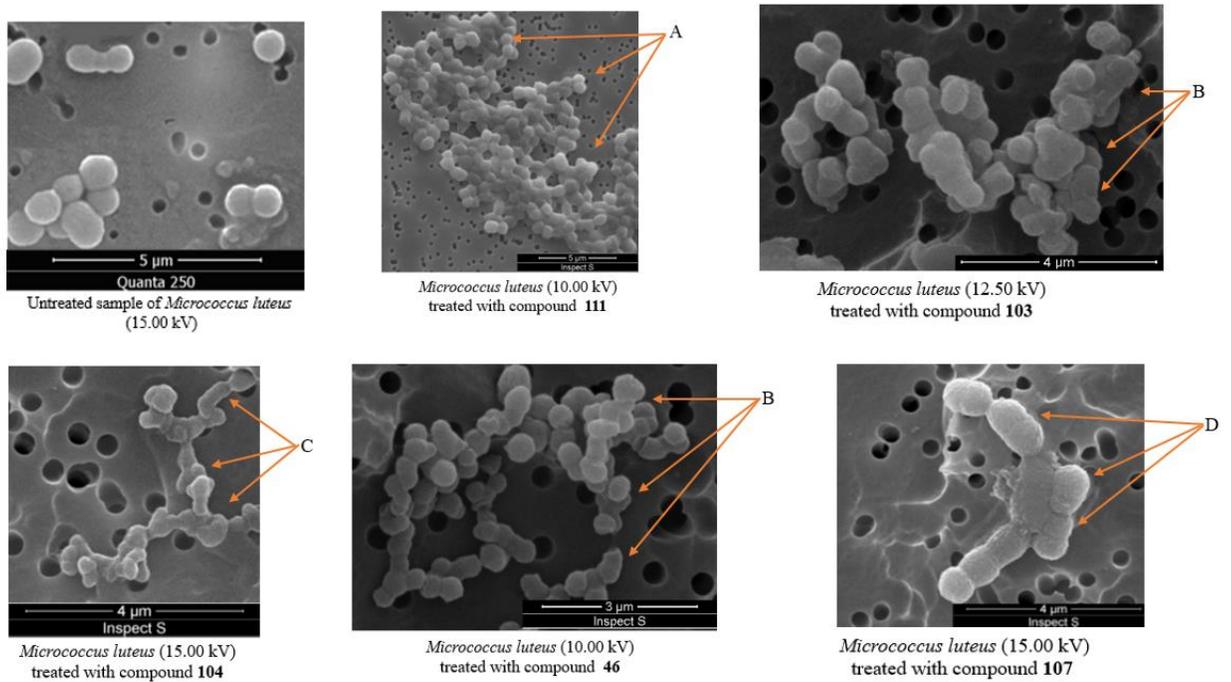


Figure 5.2: Untreated and treated samples of *Micrococcus luteus*

A: shrinking cells; B: changing shape; C: elongated cells; D: swelling cells

(Greenwood and O'Grady, 1969a; Greenwood and O'Grady, 1972; Greenwood and O'Grady, 1973a; Greenwood and O'Grady, 1973b; Monodane *et al.*, 1989; De Lillo *et al.*, 1997).

Staphylococcus aureus is widely found on the skin or in the nose. In general, these bacteria cause simple skin infections. This bacterial infection can be fatal, however, if the infection is transferred to blood, bones, lungs or heart (Wertheim *et al.*, 2005; Taylor and Unakal, 2017). *S. aureus* cells are cocci and, following cell division, produce clusters of cells (Lowy, 1998; Taylor and Unakal, 2017). The untreated cells of *S. aureus* (Figure 5.3) and the treated samples (Figure 5.4) were examined with the SEM. Compound **46** inhibited the growth of *S. aureus* by producing cells of a variable and reduced size. Compound **111** affected cell division and cell shape. Compound **107** caused damage to the cell wall of some cells (Figure 5.3). The *S. aureus* cell wall consists of 80-90% of peptidoglycan, teichoic acids, and proteins (Umeda, 1988; Dmitriev *et al.*, 2004).

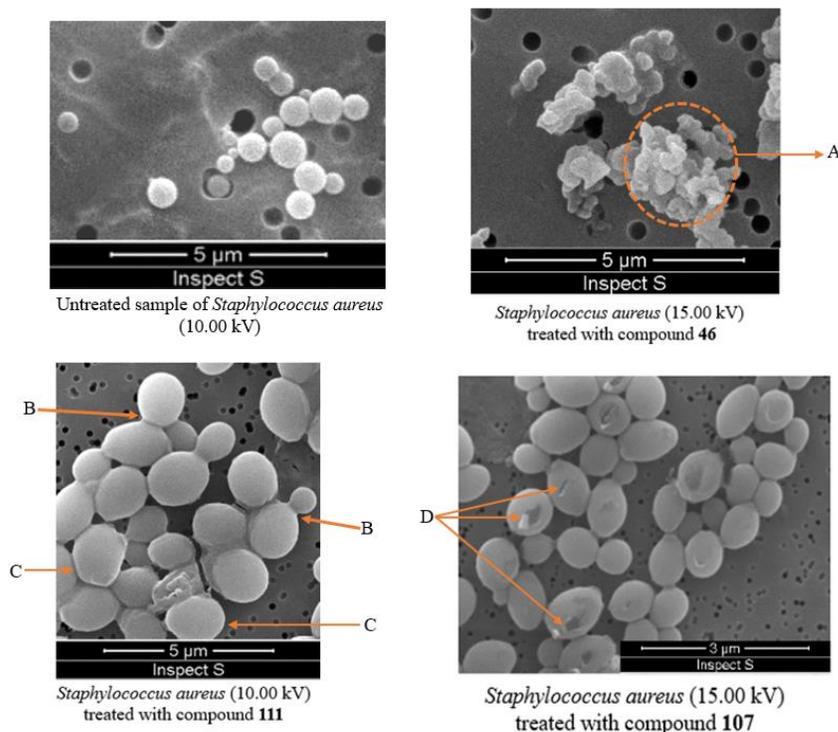
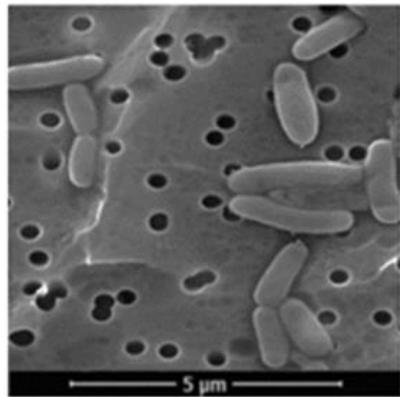


Figure 5.3: Untreated and treated samples of *Staphylococcus aureus*
 A: reduced size cells; B: cell division; C: variable shape; D: surface damage

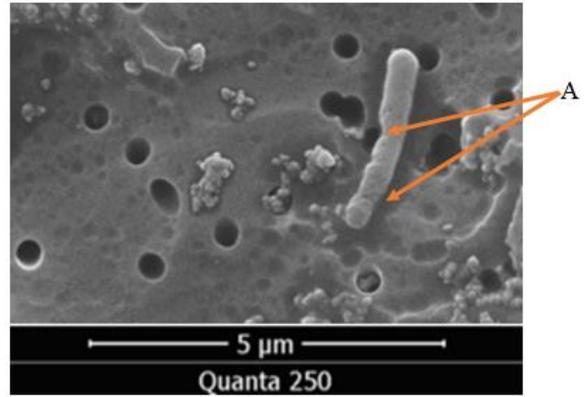
Pseudomonas aeruginosa is a common environmental bacterial species and is classified as Gram-negative with a rod shape ($0.5\text{-}0.8 \times 1.5\text{-}8 \mu\text{m}$) (Bennik, 1999). These bacteria can be isolated from soil, water, plants and animals (Stover *et al.*, 2000). It was first isolated from infections by Gessard in 1882 and identified as a pathogenic organism in 1890 by Charrin (Bodey *et al.*, 1983; Lyczak *et al.*, 2000) and in the meantime, research proved it to be responsible for community-acquired and hospital-acquired infections (Driscoll *et al.*, 2007). Both untreated and treated samples *P. aeruginosa* (Figure 5.4) were investigated using the SEM to observe the morphological effects of the selected compounds. Compounds **107** and **111** caused pronounced morphological damage, while the effect of compound **103** was simply on the bacterial surface or no fundamental difference could be detected between the untreated and treated samples.

Gram-negative bacterium *Escherichia coli* resides in the normal microflora in the human intestine, while some modified strains are pathogenic and cause health hazards. This rod-shaped bacterium is able to cause many infections such as in the central nervous system, urinary and gastrointestinal system (Nataro and Kaper, 1998; Dho-Moulin and Fairbrother, 1999). The normal strain of *E. coli* and the treated specimen (Figure 5.5) were analysed and compared under SEM to detect any surface damage. Compound (**107**) caused significant modifications on the surface of bacteria while compound **111** caused simple damage (Figure 5.5).

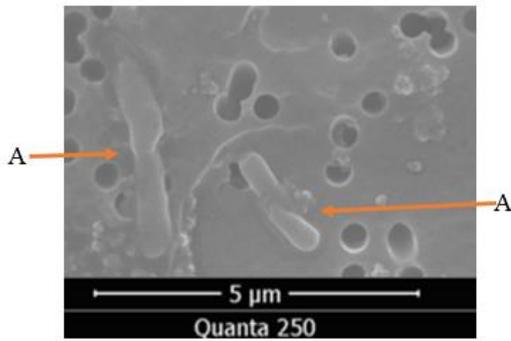
There are only a handful of preliminary studies available where SEM was used to study the effect of phytochemicals on the morphological aspects of microorganisms. To the best of our knowledge, this assay and the possible mechanism of action of selected compounds are reported here for the first time.



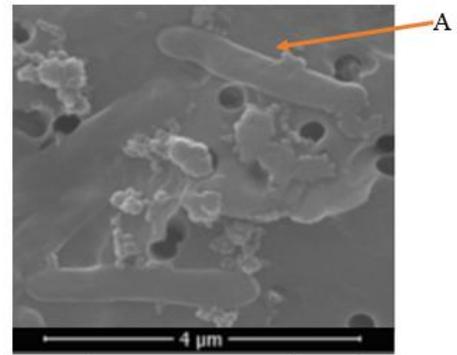
Untreated sample of *Pseudomonas aeruginosa* (20.00 kV)



Pseudomonas aeruginosa (25.00 kV)
treated with compound **103**



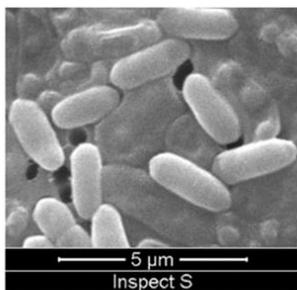
Pseudomonas aeruginosa (20.00 kV)
treated with compound **107**



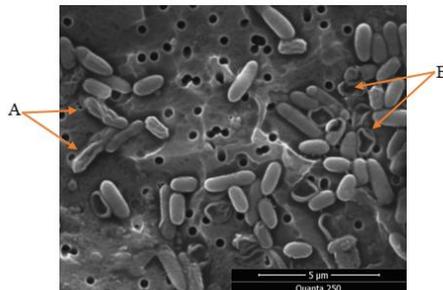
Pseudomonas aeruginosa (25.00 kV)
treated with compound **111**

Figure 5.4: Untreated and treated samples of *Pseudomonas aeruginosa*

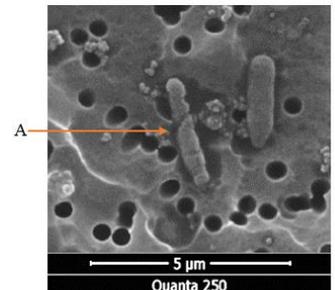
A: A: degradation of the cell wall



Untreated *Escherichia coli* (10.00 kV)



Escherichia coli (25.00 kV)
treated with compound **111**



Escherichia coli (25.00 kV)
treated with compound **107**

Figure 5.5: Untreated and treated samples of *Escherichia coli*

A: A: degradation of the cell wall; B: Cell lysis

5.6 Conclusion

This new SEM assay was carried out in conjunction with the resazurin assay and in the presence of standard samples to detect any morphological changes. There was evidence of the cell wall degradation in the treated samples when compared with untreated samples, for example the effect of compound **107** on *P. aeruginosa* (Figure 5.4). This indicates that this compound could be damaging the cytoplasmic membrane proteins or it could be binding with the proteins. Moreover, the treated sample of *C. albicans* with compound **107** (Figure 5.1) created a mucilage layer around the cells, which may be a protective response. Some compounds caused shrinkage of cells or example compound **46** against *S. aureus* (Figure 5.3). On the other hand, some compounds caused swelling of the cells, for example compound **107** against *M. luteus* (Figure 5.2). This suggests that these particular compounds may have the ability to affect the osmotic concentration of the cells. . Moreover, compound **111** affected cell division of *S. aureus*, which could be a possible effect of the compounds on the cell-cycle phases.

One of the main benefits of this assay could potentially be a better understanding of structure-activity relationships among the tested phytochemicals based on their effects on the morphology of microbial cells.

Chapter 6
**Conclusion and
Recommendations**

In the present study, three Iraqi medicinal plants from the Rutaceae family viz: *Citrus grandis* (leaves), *Citrus sinensis* (leaves and peel) and *Ruta chalepensis* (fruits, stems, leaves and roots) have been investigated for their phytochemical and antimicrobial properties following a bioassay-guided approach.

The antimicrobial activity of the selected plants was evaluated using the microtitre resazurin assay, and performed using two Gram-positive bacterial strains, *Micrococcus luteus* and *Staphylococcus aureus*, two Gram-negative bacterial strains *Escherichia coli* and *Pseudomonas aeruginosa* and one fungal strain *Candida albicans*.

The DCM extracts of *C. grandis* and *C. sinensis* leaves gave strong inhibitory activity against all of the bacterial strains used in this study. While, both of the DCM and the MeOH extracts of *C. sinensis* peel exhibited significant activity, the DCM extract was particularly potent against *M. luteus* (MIC 4.88×10^{-3} mg/mL). The phytochemical studies on the active extracts/fractions of *C. grandis* and *C. sinensis* revealed fourteen flavonoids (Figure 3.80) including hesperidin (**50**), rutin (**48**), narirutin (**51**), rutin 3'-methyl ether (**101**), 3-methoxynobiletin (**110**), nobiletin (**57**), sinensetin (**111**), 6,7,8,3',4'-pentamethoxyl-flavone (**112**), demethylnobiletin (**113**), 5-desmethylnobiletin (**114**), cirsilineol (**115**), tangeritin (**116**), tetramethylscutellarein (**117**), salvigenin (**118**) and Marcitrus (**119**), which is reported for the first time from the Iraqi species of *C. grandis*. Compound **119** is also reported for the first time as a natural product.

Ruta chalepensis, which is well known as a good source of a variety of secondary metabolites, e.g., alkaloids, coumarins, flavonoids and terpenes, has been investigated in the current project for the antimicrobial activity of its fruits, leaves, stems and roots. The *n*-hexane, DCM and MeOH extracts of the fruits, leaves, and stems showed notable antimicrobial activity, particularly, the leaf methanolic extracts against *C. albicans* (MIC 1.95×10^{-2} mg/mL). While,

the root MeOH extract exhibited a low level of antimicrobial activity, both the *n*-hexane and the DCM extracts did not display any effect with the maximum concentration 10 mg/mL. The phytochemical studies carried out on *R. chalepensis* active parts/extracts/fractions led to the isolation of nineteen compounds (Figure 3.5) including bergapten (**43**), kokusaginine (**99**), isokokusaginine (**100**), skimmianine (**41**), rutin (**48**), rutin 3'-methyl ether (**101**), rutin 7,4'-dimethyl ether (**102**), 6-hydroxy-rutin-3'-7-dimethyl ether (**103**), chalepin (**45**), chalepentin (**46**), rutamarin (**47**), isopimpinellin (**44**), γ -fagarine (**42**), imperatorin (**104**), graveoline (**105**), ribalinium (**106**), arborinine (**107**), 3',6'-disinapoylsucrose (**108**) and hexadecane (**109**). To the best of our knowledge, this is the first report on the phytochemical studies on the fruits of the Iraqi species of *R. chalepensis*. Compounds **100-102** and **108** are reported here for the first time from *R. chalepensis*. Moreover, compound 6-hydroxy-rutin-3'-7-dimethyl ether (**103**) is a new natural product.

The compounds isolated from *C. grandis*, *C. sinensis* and *R. chalepensis* extracts exhibited potent antimicrobial activity. Compounds **42, 43, 44, 45, 46, 47, 48, 50, 57, 99, 101, 103, 104, 105, 107, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118** and **119** were screened to evaluate their antimicrobial properties against the microbial strains mentioned earlier, and 1 mg/mL concentration was selected as the maximum dosage (stock concentration).

Compounds **42, 43, 44, 45, 46, 47, 48, 50, 57, 99, 101, 103, 104, 105, 107, 110, 111, 113, 116** and **118** were assessed for their anti-MRSA activity against five methicillin-resistant *Staphylococcus aureus* strains (SA1199B, XU212, MRSA340702, EMRSA-15MRSA274819) and standard strain (ATCC25923). Compounds **45, 46, 47, 57, 101, 103, 104, 107, 110, 111** and **113** showed anti-MRSA efficacy.

It is important to understand the mechanism of action of any medication or bioactive compound including antimicrobials. This may help to determine how the tested compounds

inhibit the growth of microorganisms. Compounds **46**, **57**, **103**, **104**, **111** and **107** were investigated using the scanning electron microscope (SEM) to understand the mechanism of action of these compounds. The results revealed how some of these compounds affected the morphological features of the microorganisms by damaging the cell wall, shrinking cells, dilating cells or affecting the cell division progress.

This seems to be the first report of a study on antimicrobial the mechanism of action of compounds directly from resazurin microtitre assay plates using the SEM and looking for morphological changes connected with the compound

Future studies:

- Investigation of the antimalarial activity and antioxidant properties of the isolated compounds. The previous studies on both of *R. chalepensis* and *Citrus* extracts were indicated their antimalarial and antioxidant activities but most of the isolated compounds need to be studied.
- Assessment of the performance of isolated compounds from *Citrus* in the cosmetics industry. As the *Citrus* extracts are an important components in cosmetics and they contain different secondary metabolites, there is a need to identify the responsible compounds responsible for the activity.
- Synthesis of structural analogues of active antimicrobial compounds identified in this study. Plants extracts provided unique compounds, which is difficult to obtain and maybe with a simple synthesis these compounds can be produced in larger quantities to carry out more extensive antimicrobial studies toward their mechanisms of action and structure activity relationships.

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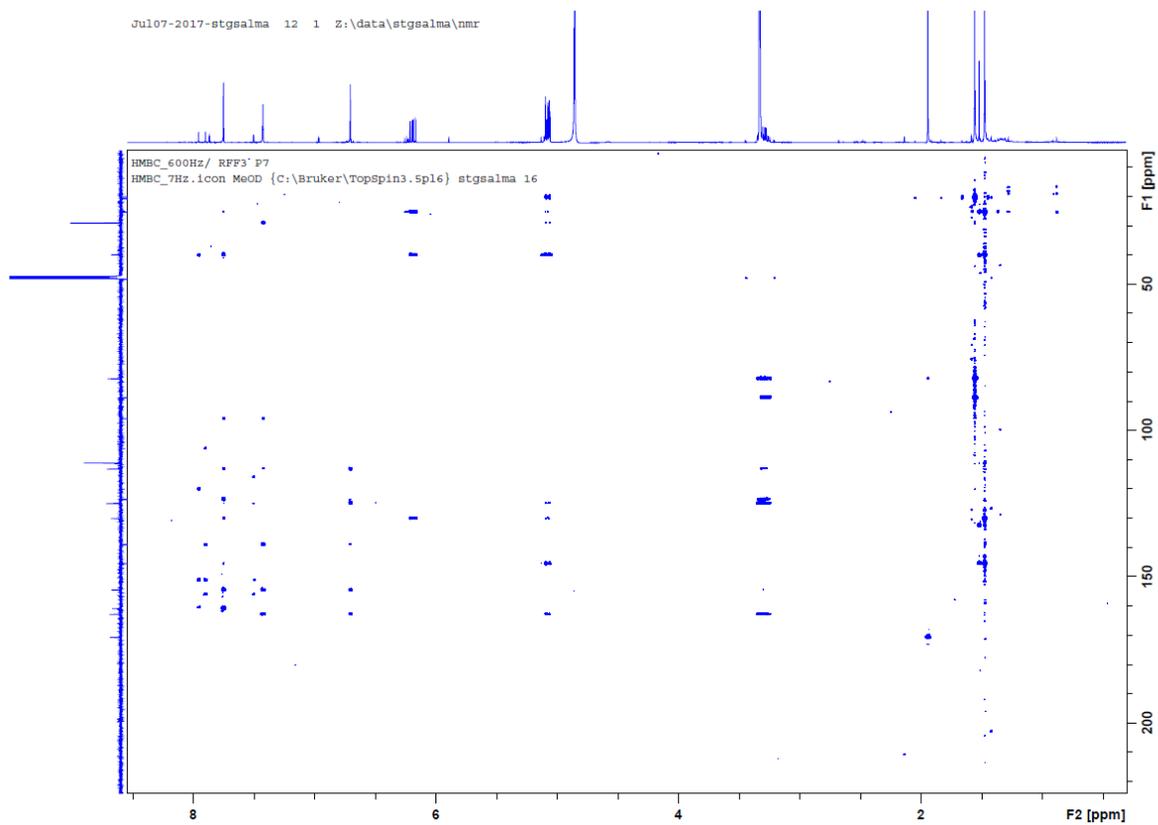
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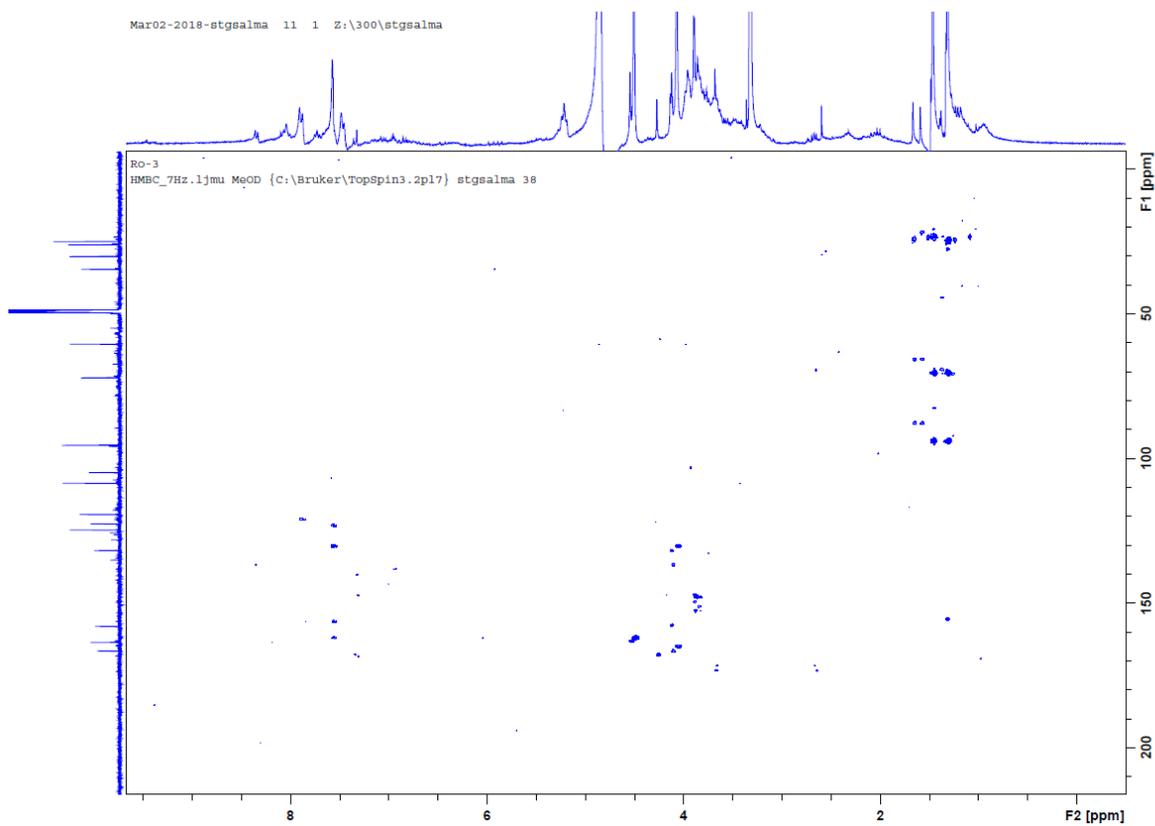
Appendix 1: The numbers and names of the isolated compounds from *Ruta chalepensis* and *Citrus species*

Compound number	Compound name	Compound number	Compound name
41	Skimmianine	110	3-methoxynobiletin
42	γ -fagarine	111	Sinensetin
43	Bergapten	112	6,7,8,3',4'-pentamethoxyl-flavone
44	Isopimpinellin	113	Demethylnobiletin
45	Chalepin	114	5-desmethylinensetin
46	Chalepensin	115	Cirsilineol
47	Rutamarin	116	Tangeritin
48	Rutin	117	Tetramethylscutellarein
50	Hesperidin	118	Salvigenin
51	Narirutin	119	Marcitrus
57	Nobiletin		
99	Kokusaginine		
100	Isokokusaginine		
101	Rutin 3'-methyl ether		
102	Rutin 7,4'-dimethyl ether		
103	6-hydroxy-rutin-3'-7'-dimethyl ether		
104	Imperatorin		
105	Graveoline		
106	Ribalinium		
107	Arborinine		
108	3',6'-disinapoylsucrose		
109	Hexadecane		

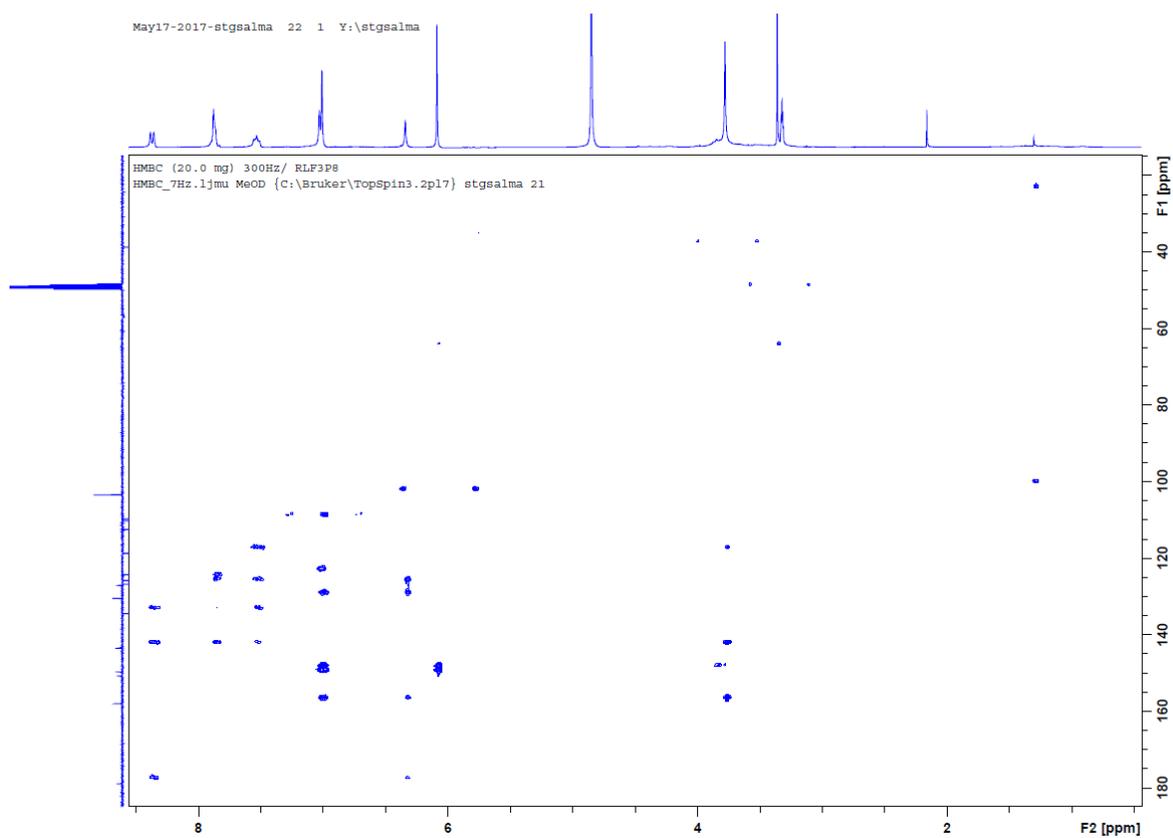
Appendix 2: HMBC for compound 47



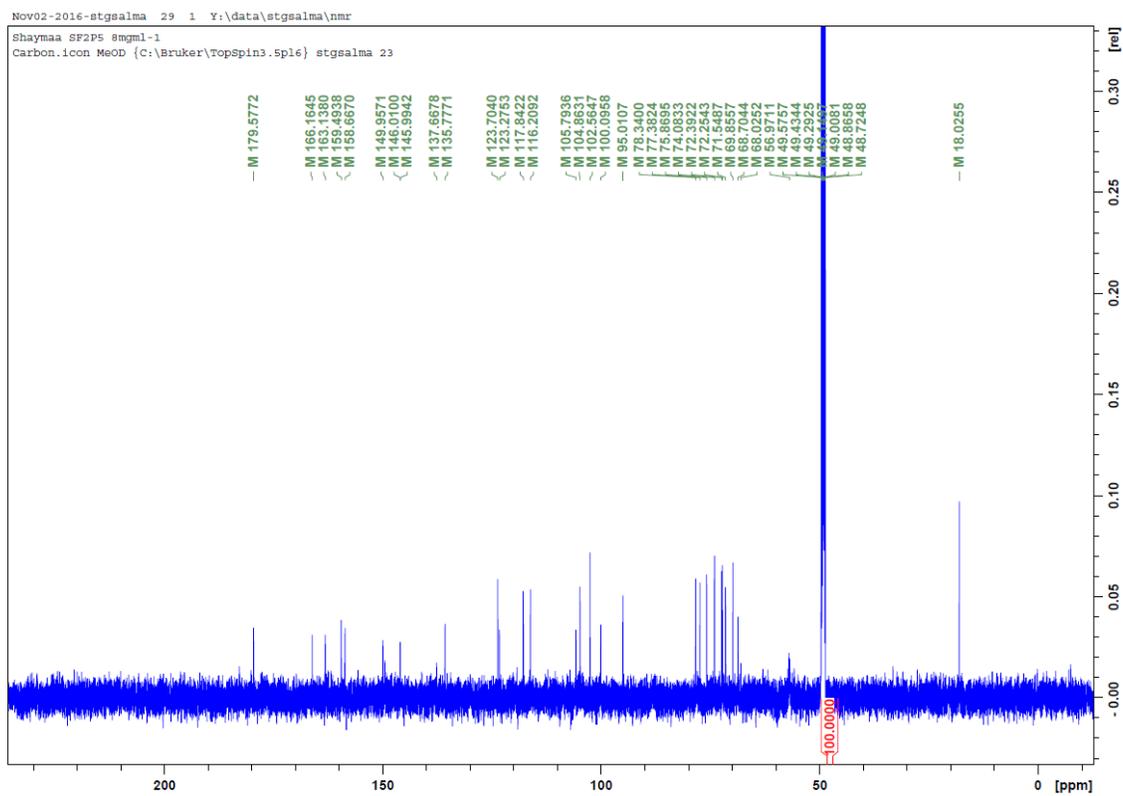
Appendix 3: HMBC for compound 106



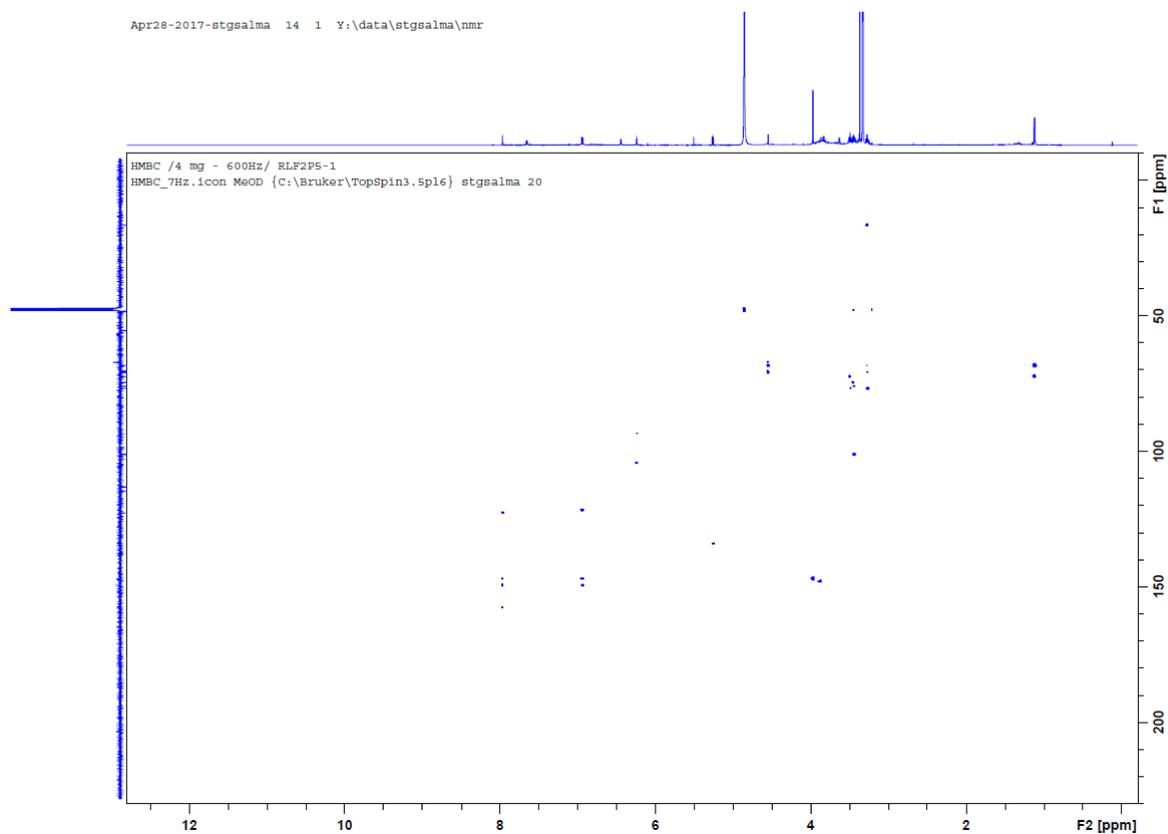
Appendix 4: HMBC for compound 105



Appendix 5: ^{13}C for compound 48



Appendix 6: HMBC for compound 101



Appendix 7: List of presentations

- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2016). Phytochemical and Antimicrobial Studies on Selected Medicinal Plants from the Iraqi Flora. Poster presentation- UK, Liverpool John Moores University, June 2016.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2017). Phytochemical and Antimicrobial Studies on Selected Medicinal Plants from the Iraqi Flora. Poster presentation- UK, The 9th MMU Postgraduate Research Conference.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2018). Screening for Phytochemical and Antimicrobial Activity on Some Iraqi Traditional Herbal Medicine. Poster presentation- PHYTOPHARM 2018, Horgen, Switzerland, 25-27 June 2018.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2018). Screening for Phytochemical and Antimicrobial Activity for *Citrus sinensis* and *Citrus grandis* on Iraqi Flora. Poster presentation- PSE-YSM 2018, Liverpool, UK, 02-05 July 2018.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2017). Phytochemical and Antimicrobial Studies on *Ruta chalepensis* L. (Rutaceae). Oral presentation- UK, Liverpool John Moores University, Monday 19th June 2017.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2018). HPLC-PDA Based Chemical Profiling and Antimicrobial Activity Studies on *Citrus grandis* and *C. sinensis* from the Iraqi Flora. Oral presentation- ISCNP30 and ICOB10, Athens, Greece, 25-29 November 2018.
- Al-Majmaie, S., Nahar, L., Sharples, G. P. and Sarker, S. D. (2018). *In vitro* antimicrobial activity of *Ruta chalepensis* L. growing in Iraq. Oral presentation- PSE-YSM 2018, Liverpool, UK, 02-05 July 2018.
- Al-Majmaie, S., Sharples, G. P., Nahar, L., Wadi, K., Nath, S., Dempster, N., Jasim, H., Al-Groshi, A. and Sarker, S. D. (2019). Antimicrobial activity and mechanisms of action of selected flavonoids from the Rutaceae. Oral presentation- Trends in Natural Product Research – PSE Young Scientists’ Meeting, 19-22 June, Budapest, Hungary.

Appendix 8: List of papers during the PhD project

- Al-Majmaie**, S., Nahar, L., Sharples, G. P., Wadi, K. and Sarker, S. D. (2019) Isolation and antimicrobial activity of rutin and its derivatives from *Ruta chalepensis* (Rutaceae) growing in Iraq, *Records of Natural Products* **13**, 64-70.
- Nurunnabi, T. R., Nahar, L., **Al-Majmaie**, S., Rahman, S. M. M., Sohrab, M. H., Billah, M. M., Ismail, F. M. D., Rahman, M. M., Sharples, G. P. and Sarker, S. D. (2018) Anti-MRSA activity of oxysporone and xylitol from the endophytic fungus *Pestalotia* sp. growing on the Sundarbans mangrove plant *Heritiera fomes*, *Phytotherapy Research* **32**, 348-354.
- Nurunnabi, T. R., **Al-Majmaie**, S., Nakouti I., Nahar, L., Rahman, S. M. M., Sohrab, M. H., Billah, M. M., Ismail, F. M. D., Sharples, G. P. and Sarker, S. D. (2018) Antimicrobial activity of the ethyl acetate extract and its major metabolite, kojic acid, from endophytic fungus *Colletotrichum gloeosporioides* isolated from *Sonneratia apetala*, a mangrove plant of the Sundarbans, *Asian Pacific Journal of Tropical Medicine* **11**, 350-354.
- Uddin, S., Alnsour, L., Segun, P., Servi, H., Celik, S., Gokturk, R. S., Al Groshi, A., **Al-Majmaie**, S., Guetchueng, S. T., Nahar, L., Dempster, N. M., Ismail, F. M. D., Ritchie, K. J. and Sarker, S. D. (2017). Flavonoids from two Turkish *Centaurea* species and their chemotaxonomic implications, *Trends in Phytochemical Research* **1**, 243-248.
- Nurunnabi, T. R., **Al-Majmaie**, S., Nahar, L., Nakouti, I., Rahman, S. M. M., Sohrab, M. H., Billah, M. M., Ismail, F. M. D., Sharples, G. P. and Sarker, S. D. (2019) Sonneratinone: A new antimicrobial benzofuranone derivative from the endophytic fungus *Aspergillus niger* isolated from the Mangrove plant *Sonneratia apetala* Buch.-Ham., *Trends in Phytochemical Analysis* (submitted).