

**THE PHYTOCHEMICAL INVESTIGATION ON THE SURFACE EXUDATE OF
Microglossa pyrifolia: CASE STUDY AS ANTI-PLASMODIAL, ANTI-OXIDANT, ANTI-
LEISHMANIAL AND ANTI-MICROBIAL**

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**A thesis Submitted to the Graduate School in Partial Fulfillment for the Requirements of
the Award of the Degree of Master of Science in Chemistry of Egerton University**

**EGERTON UNIVERSITY
OCTOBER, 2014**

DECLARATION AND RECOMMENDATION

DECLARATION

This thesis is my original work and has not been submitted wholly or in part for award in any institution.

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This thesis has been prepared under our supervision and has our recommendation to be presented for examination as per the Egerton University regulations.

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ACKNOWLEDGEMENTS

I would like to sincerely thank my lead supervisor, Prof Jacob O. Midiwo for taking the role of a promoter and mentor. Without his constant words of encouragement and support this work would have been much harder to accomplish. I am particularly grateful to him for his financial support, endless patience and vast source of knowledge in attempting to make a chemist of me.

I am grateful to Prof. J. Matasyoh, Dr F. Okanga, Dr L. Kerubo for guiding and encouraging me throughout my studies. I also express my appreciation to the academic, technical and support staff of the Department of Chemistry, University of Nairobi for their assistance in various aspects of my study.

Many thanks to Dr Muhammad Ilias of the National Center for Natural Products Research School of Pharmacy University of Mississippi for retrieving the spectra for structural analysis, and his group for helping with the antiplasmodial, antimicrobial and antileishmanial tests as well. I also thank Mr. Patrick Mutiso of Department of Botany, University of Nairobi for his assistance in collection and identification of the plant material. I am also grateful to my laboratory mates and graduate students at the Department of Chemistry, University of Nairobi and all research students whom I met in the course of my studies for the friendship they accorded me, their kind advice, assistance and collaboration.

ABSTRACT

Malaria is a global disease prevalent in the tropics caused by Plasmodium species. It affects 300-500 million people and kills 1.5-2.7 million people annually. To date, a large number of herbal remedies are used to treat malaria and manage related fever; nevertheless, in most of these cases of plant therapies, efficacies have not been proven nor have their active components been identified. Most malaria episodes are accompanied by clinical complications attributable to radical oxygen species. In an effort to address the problem of malaria and the associated complications, phytochemical analysis of *Microglassa pyrifolia* (Asteraceae), used to treat malaria ethnomedically, was investigated for compounds with anti-plasmodial, antimicrobial, antileishmanial and anti-oxidant activities. In the preliminary investigation, the crude extracts of aerial parts of the plant [dichloromethane: methanol (1:1) extract] had an anti-plasmodial activity against D6 and W2 strains of *Plasmodium falciparum* of IC₅₀ value of 8.0 and 13.0 µg/ml respectively. In this study crude leaf surface extract was obtained by dipping a handful of fresh leaves into ethyl acetate for a short period of 15 seconds to avoid the extraction of internal tissue components. Separation of compounds was carried out using chromatographic methods and their identification done by spectroscopic methods. A total of seven compounds were isolated, namely; 5,7,4'-trihydroxy-3,8,3'-trimethoxyflavone (**81**), 5,7,4'-trihydroxy-3,8,3',5'-tetramethoxyflavone (**82**), 5,3'4'-trihydroxy-7-methoxyflavanone (**83**), 5,7,3'-trihydroxy-3,8,2',5'-tetramethoxyflavone (**84**), 5,3'4'-trihydroxy-3,7,8-trimethoxyflavone (**85**), 8-acetoxyisochiliolide lactone (**86**) and 7,8-epoxyisochiliolide lactone (**87**) out of which compounds **82**, **84** and **86** were new. The *in vitro* anti-plasmodial tests were done by a colorimetric assay that determined the parasitic lactate dehydrogenase (pLDH) activity in 96-well microplate. Anti-oxidant activity was performed by UV-VIS spectrometry method based on the stable radical compound DPPH. The compounds showed no anti-plasmodial, anti-bacterial, anti-fungal and anti-leishmanial activities except 5,3,4'-trihydroxy-7-methoxyflavone (**83**) which exhibited mild anti-leishmanial activity (IC₅₀ 13.13 µg/ml). Anti-oxidant activities were exhibited, with IC₅₀ of 6.02 ± 0.2 for 5,7,4'-trihydroxy-3,8,3'-trimethoxyflavone (**81**), 6.45 ± 0.3 for 5,7,4'-trihydroxy-3,8,3',5'-tetramethoxyflavone (**82**), 8.5 ± 0.3 for 5,3'4'-trihydroxy-7-methoxyflavanone (**83**), 8.79 ± 0.3 for 5,7,3'-trihydroxy-8,4'5'-trimethoxyflavone (**84**) and 6.02 ± 0.2 for 5,3',4'-trihydroxy-3,7,8-trimethoxyflavone (**85**)

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LIST OF ABBREVIATIONS AND ACRONYMS

ACT	Artemisinin Combination Therapies
DCM	Dichloromethane
DDT	Dichlorodiphenyltrichloroethane
DEPT	Distortionless Enhancement by Polarization Transfer
DMSO	Dimethylsulfoxide
DPPH	2, 2-Diphenyl-1-picrylhydrazyl
EIMS	Electron Impact Mass Spectroscopy
HMBC	Heteronuclear Multiple Bond Correlation
HMQC	Heteronuclear Multiple Quantum Correlation
HR-EIMS	High Resolution Electron Impact Mass Spectrometry
HSQC	Heteronuclear Single Quantum Correlation
IC ₅₀	Inhibition Concentration-50: Concentration of substance that produce 50% inhibition of certain process
LC ₅₀	Lethal Concentration-50: Concentration that kills 50% of test animal
MIC	Minimum Inhibition Concentration
NMR	Nuclear Magnetic Resonance
pLDH	Plasmodium lactate dehydrogenase
PTLC	Preparative Thin Layer Chromatography
RBM	Roll Back Malaria
ROS	Reactive Oxygen Species
TLC	Thin Layer Chromatography
UNICEF	United Nation Children Fund
UV-VIS	Ultra Violet-Visible
WHO	World Health Organization

CHAPTER ONE

INTRODUCTION

1.1 Background information

Plants through an integrated network of enzymes and carefully regulated chemical reactions, synthesize crucially important molecules including carbohydrates, proteins, fats, and nucleic acids which are called primary metabolites. Despite the extremely varied characteristics of living organisms, the pathways for modifying and synthesizing these metabolites are found to be essentially the same in all organisms, apart from minor variations. These processes demonstrate the fundamental unity of all living matter, and are collectively described as primary metabolism, with the compounds involved in the pathways being termed primary metabolites (Dewick, 2002). In contrast to these primary metabolic pathways, there also exists an area of metabolism concerned with compounds which have a much more limited distribution in nature. Such compounds, called secondary metabolites, are found in only specific organisms, or groups of organisms, and are an expression of the individuality of species or groups of species called taxons. Secondary metabolites are not necessarily produced under all conditions, and in the vast majority of cases the function of these compounds and their benefit to the organism is not clear. These compounds sometimes have important ecological functions, providing protection against attack by herbivores and microbes and serving as attractants for pollinators and seed-dispersing agents. They may also contribute to competition and invasiveness by suppressing the growth of neighboring plant species, a phenomenon known as allelopathy (Osbourn and Lanzotti, 2009). In addition to their function in plants, the natural products have great impact on both human and animal life. Plants have been an integral part of life in many indigenous communities, and Africa is no exception (Sidigia *et al.*, 1995). Mankind has throughout the ages used natural products for various purposes including flavors, fragrances, dyes and most important, for medicine.

The use of plants for medical purposes dates back to antiquity (Sofowara, 1982). Most of Africa's biodiversity play major specific roles in the cultural evolution of human societies (Mugabe and Clark, 1998). The use of natural products for treatment of parasitic diseases is well known and documented since ancient times, and stems from the fact that some of these natural products are biosynthesized as defense agents against plant pathogens (Kaur *et al.*, 2009). Initially these medicines took a form of crude extracts contained in herbal formulations like teas, tinctures, poultices and powders (Balunas and Kinghorn, 2005).

But with the enormous advancements in separation and isolation chemistry and pharmacological testing, natural products have continued to be significant sources of drugs and leads. Their dominant role is evident in the approximately 60% of anticancer compounds and 75% of drugs for infectious diseases that are either natural products or natural product derivatives (Newman *et al.*, 2003; Cragg *et al.*, 2005).

With developments in synthetic and computational chemistry, some drugs are designed through molecular modeling (Meyer *et al.*, 2000). However, despite all these developments, natural products and their derivatives represent more than 50% of all the drugs in clinical use in the world, including the treatment of malaria (Gurib-Fakim, 2006). Historically, majority of antimalarial drugs have been derived from medicinal plants or from structures modeled on plant lead compounds (Klayman, 1985). Quinine, artemisinin, derivatives and analogues, the drugs of choice for treatment of malaria, were either obtained directly from plants or developed using chemical structures of plant derived compounds as templates (Basco *et al.*, 1994; Kayser *et al.*, 2003). Many more compounds with significant anti-plasmodial activities have been isolated from plants and efforts are being made to develop some of these into future drugs (Kaur *et al.*, 2009). In underdeveloped communities, crude natural products made in the form of herbal formulations are still used for malaria therapy. Several plants with traditional use in curing malaria by different communities have been documented (Gessler *et al.*, 1995; Kvist *et al.*, 2006; Muthaura *et al.*, 2007a).

The recent emergence of multi-drug resistant malaria parasites and insecticide resistant mosquitoes, coupled with the limited number of chemotherapeutic agents available to treat and control the disease has triggered an alarming increase in the incidence and spread of the disease therefore highlighting the need for new chemically diverse effective drugs (Pamisha *et al.*, 2007). Another challenge encountered in the use of natural product-derived drugs and drugs from other sources in combating malaria has been the complications arising from oxidative stress during malaria infections. Oxidative stress has been linked to several malaria complications such as cerebral and pulmonary oedema, anemia and poor eyesight (Taoufiq *et al.*, 2006). The use of natural product-derived drugs from other sources in combating malaria has however been faced with several challenges, including the emergence of drug resistant parasites to drugs world-wide as well as complications arising from oxidative stress during malaria infections.

The increase in resistance to the anti-malarial drugs has necessitated continued search for alternative drugs to chloroquine and other commonly used anti-malaria drugs (Muthaura *et al.*, 2007a). In this regard, several plants are being screened for anti-plasmodial activity.

In a collaboration programme between University of Nairobi and University of Mississippi, 140 plants were screened out of which six plants namely *Clerodendrum eriophyllum* (Verbenaceae), *Sphaeranthus bullatus* (Asteraceae), *Microglossa pyrifolia* (Asteraceae), *Vernonia galamensis* (Asteraceae), *Albizia gummifera* (Fabaceae) and *Alibizia schimperiana* (Fabaceae) showed activities as indicated in Table 1:1. These activities were interpreted as good to moderate activities, based on WHO guidelines on activity, which stipulates that extracts with IC₅₀ of less than 10 µg/ml have promising activity, whereas those with IC₅₀ of 11-50 µg/ml have moderate activity. The ethanol extract of *M.pyrifolia* showed antimicrobial effects which were poor to moderate with MIC values no less than 500 µg/mL as indicated in table 1.2. The same ethanol extract exhibited antioxidant properties with IC₅₀ value of 47.31µg/mL for free radical scavenging activity (Dickson *et al.*, 2006). The active ingredients in the extracts of these plants were therefore to be sought through chromatographic fractionation and further bioassay.

Table 0.1: Anti-plasmodial activities of the selected plant extracts towards D6 (Chloroquin sensitive) and W2 (Chloroquin resistant) strains of *Plasmodium falciparum*

Plant	<i>In vitro</i> activity (IC ₅₀ ,µg/ml)	
	Pf D6	Pf W2
<i>Microglossa pyrifolia</i> (whole plant)	8.0	13
<i>Albizia gummifera</i> (root bark)	1.1	1.0
<i>Sphaeranthus bullatus</i> (whole plant)	9.7	15
<i>Alibizia schimperiana</i> (stem bark)	0.97	1.5
<i>Vernonia galamensis</i> (aerial parts)	9.0	9.2
<i>Clerodendrum eriopyllum</i> (aerial parts)	8.8	8.8

Table 0.2: Mean minimum inhibitory concentration (MIC; $\mu\text{g/mL}$) of *M. pyrifolia* on various microorganisms

Test organism	<i>M.pyrifolia</i>	Tetracycline	Miconazole
Bacteria			
<i>Staphylococcus aureus</i>	>1000	2.5	NT
<i>Bacillus subtilis</i>	500	5	NT
<i>Micrococcus flavus</i>	>1000	1	NT
<i>Streptococcus faecali</i>	1000	5	NT
<i>Escherichia coli</i>	0	10	NT
<i>Pseudomonas aeruginosa</i>	0	20	NT
<i>Salmonella abony</i>	0	2.5	NT
<i>Klebsiella aerogenes</i>	0	5	NT
Fungi			
<i>Candida albicans</i>		NT	10
<i>Saccharomyces cerevisiae</i>		NT	10
<i>Trichophyton interdigitale</i>		NT	20
<i>Microsporium floccosum</i>		NT	20

0, no inhibition observed; NT, not tested. All tests were done in triplicates. Tetracycline and miconazole served as positive control for antibacterial and antifungal assays respectively.

1.2 Statement of the problem

Malaria is the world's deadliest parasitic disease. It is transmitted by female *Anopheles* mosquitoes, and is a serious significant impediment to socioeconomic development in poor countries. It is one of the most important diseases of the African and Asian countries, killing 1-3 million people and causing disease in 300-500 million people in 107 countries annually.

Of these deaths, over 90% occur in tropical Africa. Some of the fatalities are caused by elevated production of radicals in host cells. Efforts to combat the disease are hampered by growing resistance of malaria parasites to the readily available drugs. There is, therefore, a need for continued efforts in the search for antioxidants and compounds that are active against malaria parasites. These can be developed to new and more effective drugs.

1.3 Objectives

1.3.1 General objective

To identify phytochemicals with anti-plasmodial antimicrobial antileishmanial and antioxidant activities from the plant *Microglossa pyrifolia*

1.3.2 Specific objectives

1. To prepare a crude extract of *M. pyrifolia*
2. To isolate secondary metabolites from *M. pyrifolia* crude extract and elucidate the structures of the isolated pure compounds using spectroscopic methods.
3. To determine the *in vitro* anti-plasmodial, antimicrobial, antileishmanial and antioxidant activities of pure compounds of *M. pyrifolia*

1.4 Justification

In more recent times, natural products have continued to be significant sources of drugs and leads for drug development through structural modification. Some of the most important antimalarial compounds that have been isolated from traditional medicinal plants include quinine (**1**) from *Cinchona succiriba* (Rubiaceae) and artemisinin (**4**) from the *Artemisia annua* (Asteraceae). The ethno pharmacological approach for the search of new antimalarial agents from plant sources has proved to be predictive and very significant due to the success of the antimalarial drug quinine and the recent discovery of artemisinin. In Kenya, several plants have been in use traditionally for the treatment of malaria. Around Lake Victoria, Western Kenya, and in some West African countries, *M. pyrifolia* is used for the treatment of malaria. Although a lot of research has been carried out on this plant to establish its antimicrobial, antiplasmodial and antioxidant activities, phytochemical analysis to investigate the compound(s) behind these biological activities have not been undertaken. It is thus important to undertake further investigations on this plant for metabolites and to establish the compounds that are antiplasmodial (*in vitro*) and anti-oxidant.

CHAPTER TWO LITERATURE REVIEW

2.1 Malaria problem

Malaria is a life-threatening parasitic disease transmitted by several species of *Anopheles* mosquitoes and is predominant in the tropics particularly in Africa and Asian countries (Korenromp, 2004; Geissbühler *et al* 2007). It is the female sex which is the vector. Malaria is caused by protozoa of the genus *Plasmodium*. The four human-pathogenic *Plasmodium* species *P. falciparum*, *P. vivax*, *P. ovale* and *P. malariae* differ in pathogenesis and global distribution (Odugbemi *et al.*, 2007). *P. falciparum* causes most of the severe and fatal cases of malaria and is the predominant species throughout sub-Saharan Africa. Since *P. vivax* can develop in mosquitoes at a lower temperature than *P. falciparum*, its geographical distribution is wider. *P. vivax* and *P. ovale* can form resting stages in the liver (hypnozoites) which may cause clinical relapses many months after the first attack (Hetzl *et al.*, 2007).

In humans, malaria parasites are found in the liver and in the blood where they invade, grow and replicate in erythrocytes. Once sexual forms (gametocytes) of the parasites have developed and circulated in the blood stream, they may be taken up by a female *Anopheles*, and the parasite's sexual reproduction can take place in the gut of the mosquito. Once in the mosquito gut, they undergo extensive replication and when the mosquito bites another human for blood meal, the parasites can pass via the salivary glands into the blood stream of the victim usually 10-14 days after the initial blood meal (Warrel and Giles, 2002; Greenwood *et al.*, 2005). Clinical malaria is manifested by a range of symptoms. In semi-immune populations, malaria often results in a symptomatic parasitaemia; which is manifested by symptoms such as headaches, cough, body pains, chills, vomiting and diarrhea (Warrel and Gilles, 2002; Nkuo-Akenji and Menang, 2005). Severe malaria is a complex multi-system disorder, characterized by one or more of the following clinical features; Severe anaemia, prostration, impaired consciousness, respiratory disease, multiple convulsions, circulatory collapse, pulmonary oedema, abnormal bleeding, jaundice and haemoglobinuria (WHO, 2000b; Mackintosh *et al.*, 2004).

Cerebral malaria and severe anaemia are responsible for long term developmental impairments and is the cause of most deaths in children (Carter *et al.*, 2005). Severe anaemia results from the destruction of red blood cells, either directly by the malaria parasite or indirectly by immune mechanisms.

In children it may lead to long-term neurological, cognitive and developmental impairments while in pregnant women it may lead to low birth-weight, increase in infant and child mortality (Bremner, 2001; Schellenberg *et al.*, 2003b; Marchant, 2004.). Where malaria prospers most, human societies have prospered least. Malaria imposes economic costs on households and states. In some African countries, it is estimated that malaria is responsible for an economical decline of up to 13% per year (Sachs and Malaney, 2002). There is a strong link between malaria and poverty. Due to the heavy financial burden it imposes on the family and the loss of workdays and income, it leads to poverty and consequently, poverty makes the family more vulnerable to it.

In terms of economy, malaria is known to be both a disease and a cause of poverty. It is estimated that malaria is responsible for a GDP economic growth penalty of up to 1.3% per year in some African countries (Sachs and Malaney, 2002). When compounded over the years, this penalty leads to substantial differences, totaling US\$ 12 billion annually in GDP between countries without malaria and countries with malaria, and severely restrains the economic growth of entire Africa region [Kamau, 2006].

The problem of controlling malaria has been aggravated by inadequate health structures and poor socio-economic conditions. The recent emergence of multi-drug resistant malaria parasites and insecticide resistant mosquitoes has compounded the problem further and triggered an alarming increase in the incidence and spread of the disease. (Olliaro and Yuthavong, 1999; Berry, 2000). Efforts to overcome the problem are hindered by two obstacles. One is the growing resistance of malaria parasite *Plasmodium falciparum* to chloroquine and other commonly used cheap synthetic drugs, as well as the growing resistance of the vector *Anopheles* to DDT and other insecticides (Bilia, 2006). The second obstacle is limited availability, coupled with higher cost and greater toxicity of alternative drugs (Saidu *et al.*, 2000).

In recognition of the gravity of the malaria problem, WHO and UNICEF launched the Roll Back Malaria (RBM) partnership in 1998, with the aim of reducing malaria burden by at least 50% by the year 2010 through application of evidence based interventions and strengthening health delivery services (Kamau, 2006). This was supported by the Abuja declaration in 2000 where African heads of state agreed on a concerted effort to reduce malaria burden on the continent and endorsed the goals of RBM partnership of halving the number of malaria deaths by the year 2010 (Hetzl *et al.*, 2007). Currently, efforts to promote three core malaria control interventions are being undertaken concurrently: use of insecticide treated nets, intermittent preventive treatment during pregnancy and infancy, and effective treatment of clinical cases with artemisinin based combination therapies (ACTs) (Kamau, 2006).

In Kenya, like other tropical countries, malaria is endemic and resistance of malaria parasites to readily available and cheap drugs like chloroquine is growing fast. According to the Ministry of Health, twenty four million Kenyans are at risk of malaria infections and majority cannot afford the costs of the adopted artemisinin based combination therapy. The government therefore found it necessary to provide free ACT therapy in health facilities alongside expansion of healthcare provision (Muthaura *et al.*, 2007b; World Report, 2007). Despite this effort however, many regions remain underserved and subsequently the communities use herbal remedies and other cheaply available alternatives (Kirira *et al.*, 2006; Muthaura *et al.*, 2007b). Amid the malaria deaths and suffering, progress is being made. Though for too long the global community has been reluctant to invest sufficient resources in fighting malaria, leaving it near the bottom of the world's health agenda, there is gradual increase in international awareness of the problem and malaria is now on the agenda of the health community, political arena and international financial institutions. In 1997, the multi-lateral initiative of malaria (MIM), an alliance of agencies, institutions and governments, was formed to maximize the impact of scientific research on malaria through capacity building in Africa and global collaboration.

2.2 The use of plants and natural products in chemotherapy of malaria

Natural products have been investigated and utilized to alleviate disease since early human history. In early 1900s, before the “synthetic Era”, 80% of all medicines were obtained from roots, barks and leaves (James *et al.*, 2006). Thus plants and natural products have had a great contribution to the fight against malaria since time immemorial.

In Kenya and other developing countries, the use of herbal remedies in treatment of malaria is well known as evidenced by overwhelming literature on plants used traditionally to cure malaria (Kokwaro, 1976; Muregi *et al.*, 2003; Koch *et al.*, 2005; Kirira *et al.*, 2006; Muthaura *et al.*, 2007a; Muthaura *et al.*, 2007b). Apart from their use in traditional medicine, plants and natural products have also contributed to modern treatment of malaria, being the source of some commonly used anti-malarial drugs like quinine (1) and artemisinin (4) from *Cinchona* and *Artemisia* respectively and their synthetic modifications and analogues (Figure 2.1).

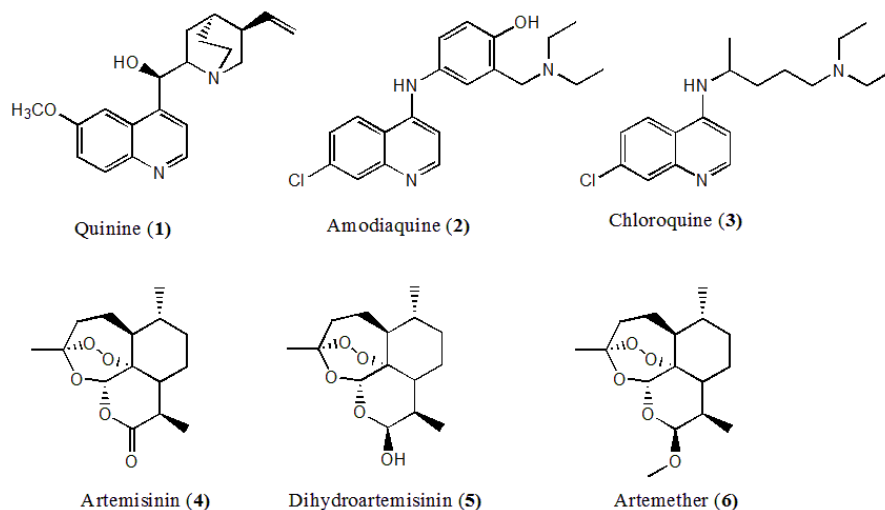


Figure 2.1: Some Natural Products and their synthetic analogues used against malaria.

2.3 Malaria and oxidative stress

Normal physiological processes in the body result in the production of Reactive Oxygen Species (ROS), which are also produced by the influence of exogenous chemicals. In the absence of parasitic attack, numerous enzymatic and non enzymatic antioxidant systems in cells and biological fluids are able to counter these ROS and their subsequent adverse effects.

However, in malaria patients, there is an increase in the production of ROS and the antioxidant defense system is weakened. This therefore results in oxidative tissue damage leading to oxidative stress associated with several pathological phenomena such as cerebral and pulmonary oedema, poor eyesight, atherosclerosis, cardiac ischemies, rheumatic diseases and cancer (Bahorun *et al.*, 1996). Antioxidants are thought to act by suppressing the enzymes responsible for superoxide production, chelating metal ions that induce free radicals and reduction of radicals by donation of hydrogen atoms (Pieta, 2000), as is illustrated in figure 2.2

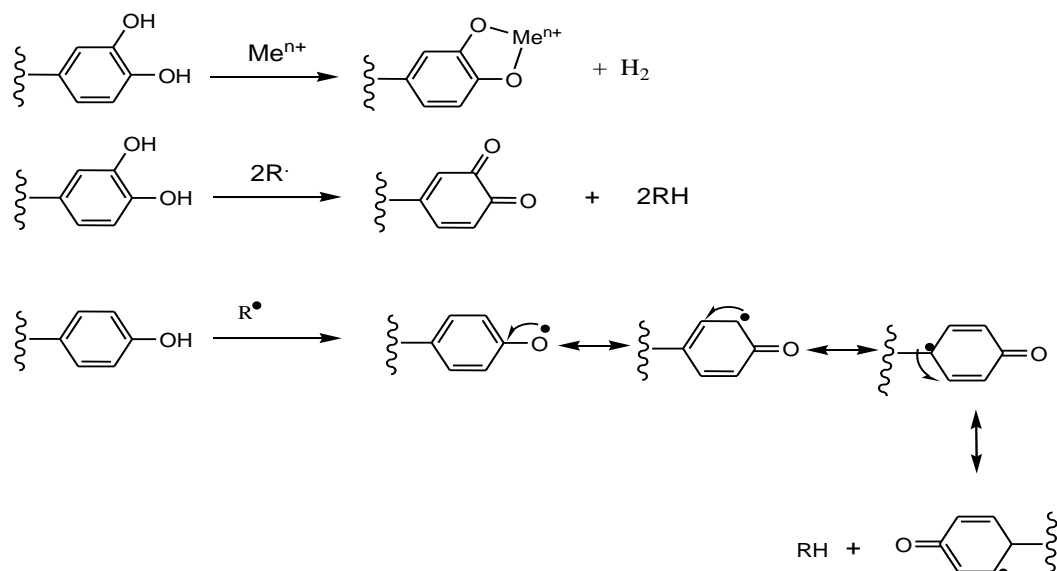


Figure 2.2: Mechanisms of antioxidant activity through chelation of metal ions and reduction by hydrogen donation.

As malaria parasites proliferate in malaria patients, large quantities of ROS are generated which damage macromolecules leading to several malaria complications. The major source of ROS is heme, a by-product of hemoglobin digestion degradation by *Plasmodium*, whereby the iron II in heme is oxidized to iron III releasing an electron that is used to ionize oxygen molecules and subsequently form peroxides which are easily broken to radicals (Figure 2.3).

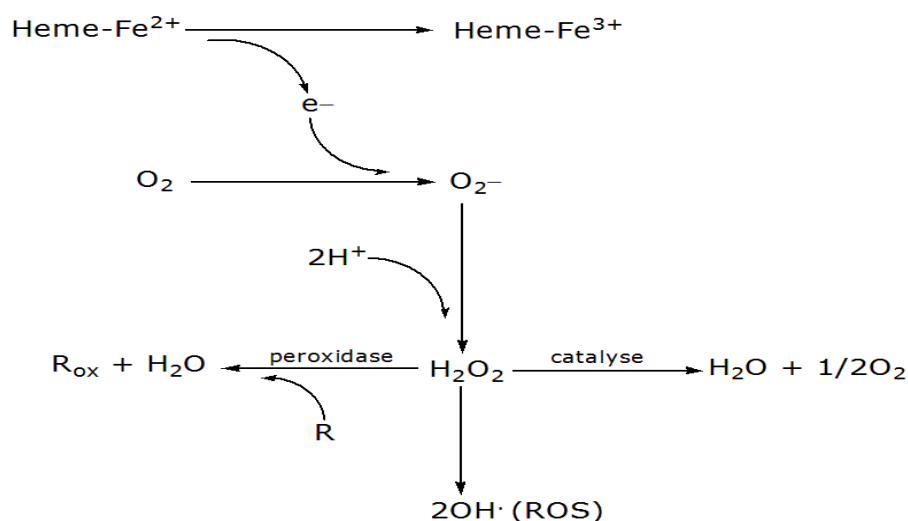


Figure 2.3: Generation of ROS from heme.

ROS are also produced by host immune system for the purpose of eradicating the parasite, as the parasite is highly susceptible to oxidative burden (Kawazu *et al*, 2008). Malaria parasites are therefore equipped with anti-oxidant defenses that are meant to establish redox equilibrium for their survival. Such defenses are the ones targeted in malaria control strategies, some of the anti-malarials like chloroquine and artemether are known to act by increasing the production of ROS (Taoufiq *et al.*, 2006).

The effect of ROS to malaria patient therefore, though beneficial as it suffocates the parasite, is also detrimental since it leads to the damage of some cells and hence bringing about more complications to the patient. It is therefore important to investigate if some of anti-plasmodial compounds can also have anti-oxidant effect.

2.4 Leishmaniasis

Leishmaniasis is a protozoan disease that affects many people in the world; 12 million are currently infected and around 350 million others are threatened (Croft *et al.*, 2003). It is transmitted by a bite of female phlebotomine sand flies which infect the blood with parasites of the genus *Leishmania* (Tonui, 2006). Two common forms of leishmaniasis are known; cutaneous leishmaniasis (CL) which causes sore (ulcers) at the bite site (Figure 2.4) and visceral leishmaniasis (VL) which affects vital organs. Sandflies *Phlebotomus martini* and *Phlebotomus orientalis*, are known vectors for *Leishmania donovani*, the causative agent for visceral leishmaniasis (VL), also known as Kalaazar. *Phlebotomus duboscqi* and *Phlebotomus guggisbergi* are vectors for *Leishmania major* and *Leishmania tropica* respectively, the aetiological agents for cutaneous leishmaniasis (CL).



Figure 2.4: Cutaneous leishmaniasis ulcer on the hand of an infected adult human being.

Leishmaniasis is found in 88 countries hosting 350 million people, being widespread in tropical and sub-tropical regions of the world where it is a serious health problem (Kigundu *et al.*, 2009). More than 90% of the global cases of visceral leishmaniasis (VL) occur in India, Bangladesh, Nepal, Brazil and Africa (particularly East and North Africa including Sudan). Globally, the number of new cases of cutaneous leishmaniasis (CL) and visceral leishmaniasis (VL) is estimated at 1.5 million and 500,000 annually, respectively. People of all ages are at risk of infection by leishmaniasis if they live or travel where the disease is endemic.

In Kenya, the disease has been reported in Baringo, Kitui, Machakos, Meru, West Pokot, Elgeyo Marakwet and Turkana districts, which are known to be endemic foci for kala-azar (VL) (Kigundu *et al.*, 2009). Currently, the recommended drugs for both VL and CL are pentavalent antimonials: sodium stibogluconate (pentosam®) and meglumine antimoniate (glucantime®). Both drugs require long courses of parenteral administration and have toxic side effects (Kigundu *et al.*, 2009).

2.5 Ethno-botanical, ethnopharmacological and phytochemical information on *Microglossa pyrifolia*

Microglossa pyrifolia (Figure 2.5) which belongs to the family Asteraceae is a climbing shrub indigenous in Africa and tropical Asia (Schmidt *et al.*, 2003). The plant is also called *Conyza pyrifolia* or *Microglossa volubilis* and is in the genus *Microglossa*, with about ten species (Kuiate *et al.*, 1999). It is traditionally used against fever and malaria in Kenya and Ghana (Kokwaro, 1976; Kohler *et al.*, 2002). It is also used as an abortifacient, as an analgesic to relieve headaches and stomach pains as well as for treatment of colds, wounds and eyes (Akah and Ekekwe, 1995; Johns *et al.*, 1990; Kuiate *et al.*, 1999; Schmidt *et al.*, 2003). Its organic solvent extract showed anti-plasmodial activity (Kohler *et al.*, 2002).



Figure 2.5: Photo of *Microglossa pyrifolia*

2.6 Phytochemistry of the genus *Microglossa*

Microglossa pyrifolia is the only species reported to have medicinal uses and biological activities amongst *Microglossa* species. Phytochemical information however is available on this plant together with *M. angolensis*, *M. mespilifolia*, *M. pyrhopappa* and *M. zeylanica*. Clerodane and *seco*-clerodane diterpenoids have been reported from *M. pyrifolia* (7 – 16, Figure 2.6) (Zdero *et al.*, 1990a; Kohler *et al.*, 2002), *M. pyrhopappa* and *M. angolensis* (17 – 43, Figure 2.7) (Zdero *et al.*, 1990b; Tene *et al.*, 2005).

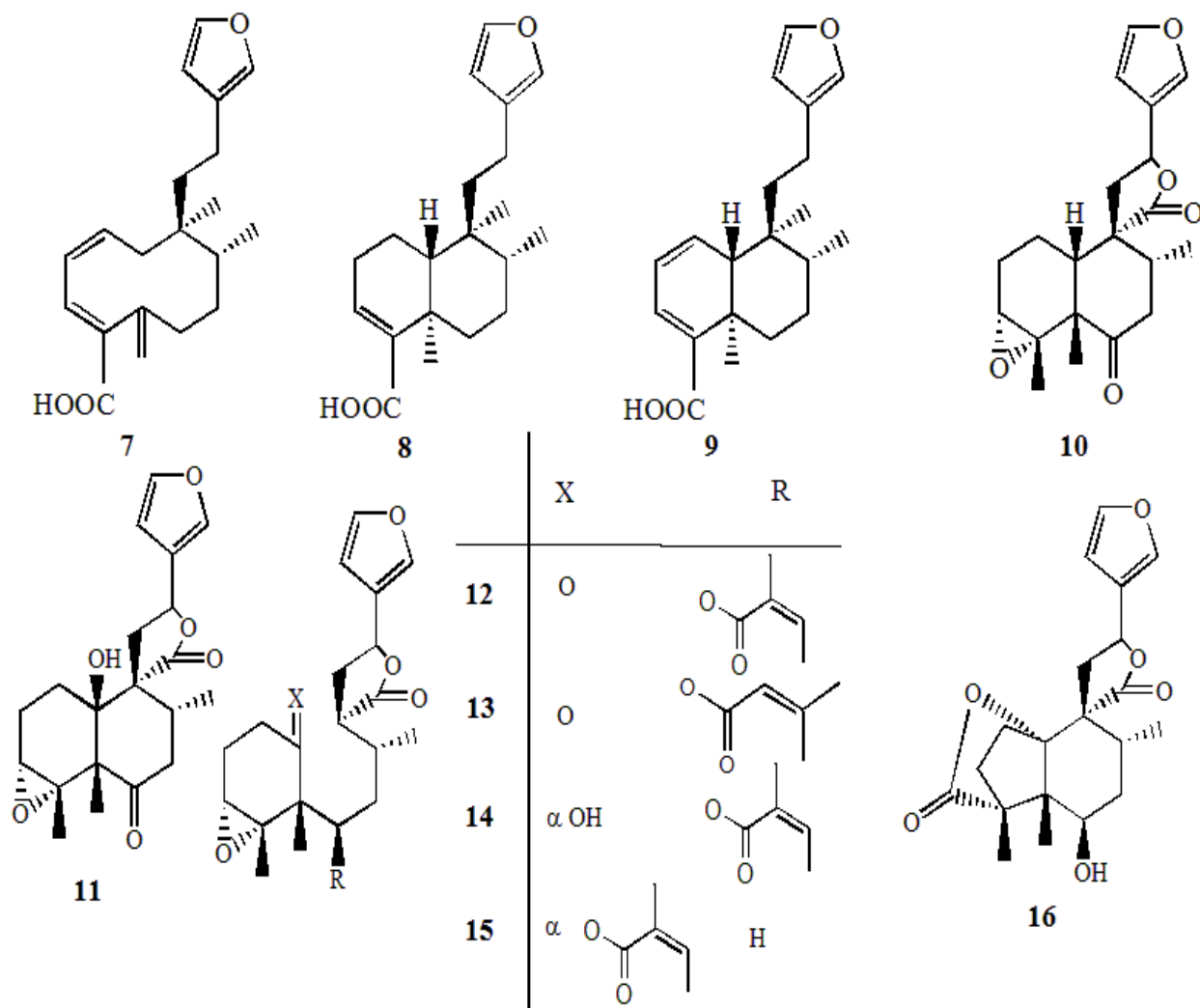


Figure 2.6: Clerodane and *seco*-clerodane diterpenoids from *M. pyrifolia*

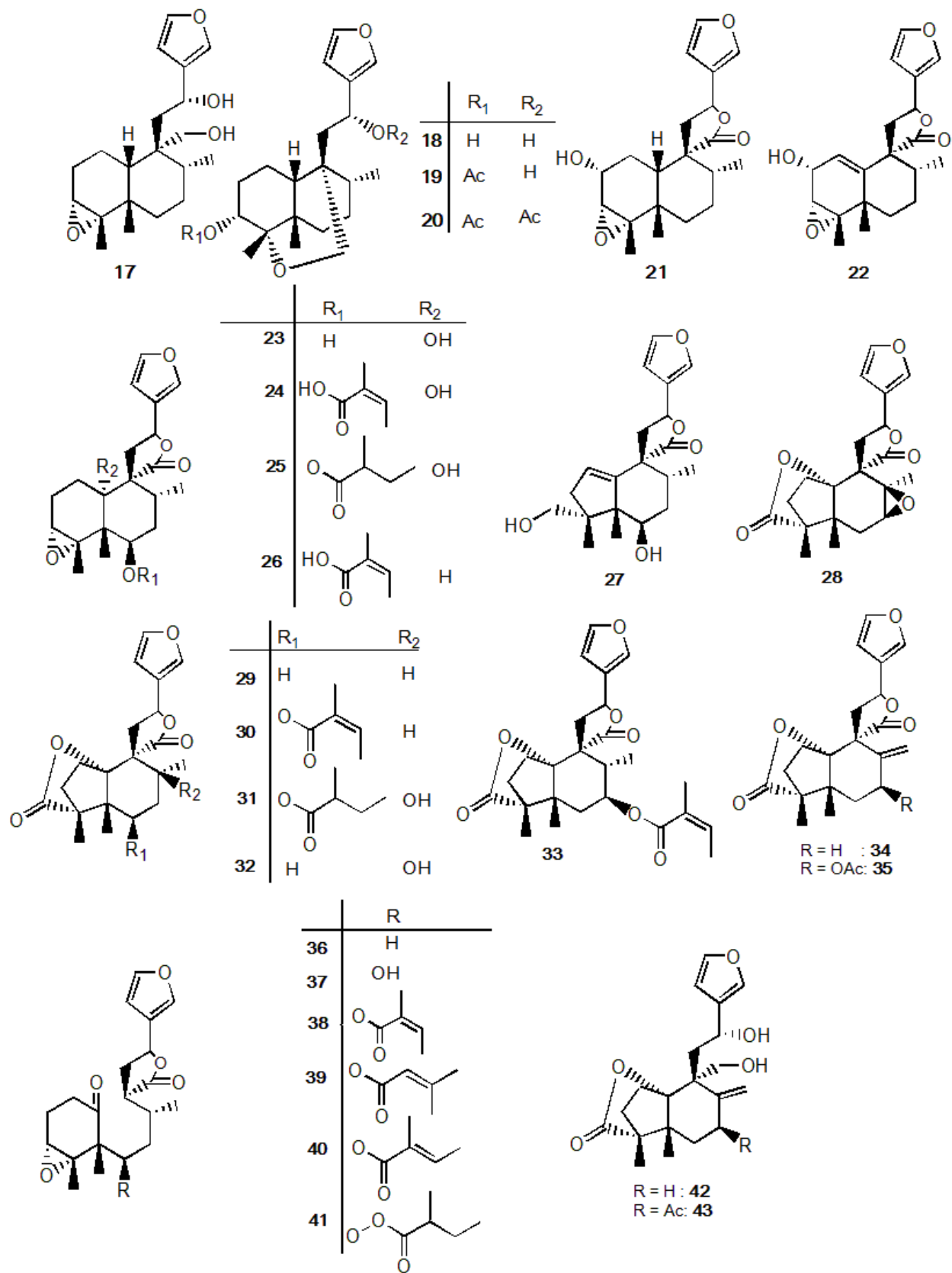


Figure 2.7: Clerodane and seco-clerodane diterpenoids from *M. pyrhopappa* and *M. angolensis*

Another class of compounds isolated from the genus *Microglossa* is phytane diterpenoids. These have been isolated from *M. pyrifolia* (**44** –**49**, Figure 2.8) (Kohler *et al.*, 2002; Zdero *et al.*, 1990a), *M. pyrhopappa* (**48** –**50**, Figure 2.8) (Zdero *et al.*, 1990b) and *M. zeylanica* (**51** – **52**, Figure 2.8) (Gunatilaka *et al.*, 1987). Out of these, phytol (**44**) and 6*E*-geranylgeraniol-19-oic acid (**46**) exhibited antiplasmodial activities of 8.5 and 12.9 μ M respectively against chloroquine sensitive strains of *Plasmodium falciparum* (PoW). Compounds **44** and **46** also exhibited antiplasmodial activities of 11.5 and 15.6 μ M respectively against chloroquine resistant strains of *Plasmodium falciparum* (Dd2) (Kohler *et al.*, 2002).

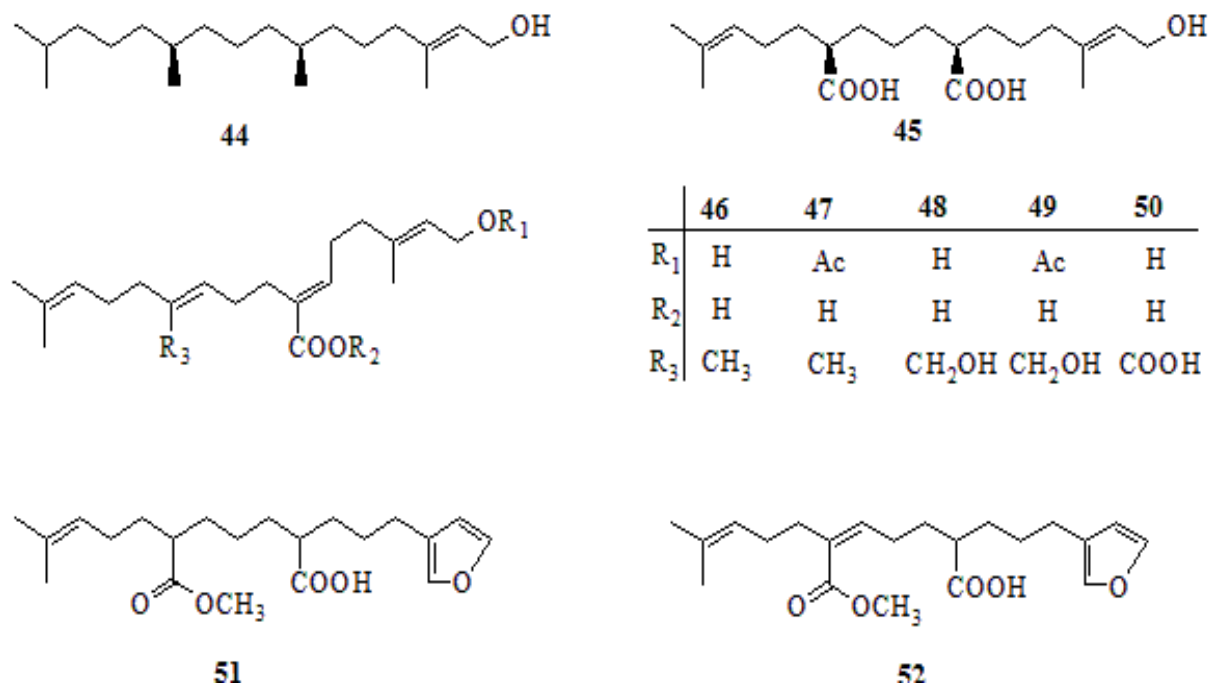


Figure 2.8: Phytane diterpenoids from *M. pyrifolia*, *M. Pyrhopappa* and *M. zeylanica*

Other compounds reported from *Microglossa pyrifolia* are dihydrobenzofurans and aurone flavonoids (**53** – **60**, Figure 2.9), (Schmidt *et al.*, 2003; Rucker *et al.*, 1992), triterpenoids (**61** – **67**, Figure 2.10), (Schmidt *et al.*, 2003), sesquiterpenes (**68**-**72**, figure 2.11), (Zdero *et al.*, 1990a; Kohler *et al.*, 2002) as well as acetylenes and acetylenic glucosides (**73** –**80**, Figure 2.12) (Rucker *et al.*, 1992). Compound **74**, namely 2- β -D-glucopyranosyloxy-1-hydroxytrideca-3,5,7,9,11-pentayne exhibited antibacterial activities against the bacteria *Escherichia coli*, *Pseudomonas aeruginosa*, *Enterococcus faecalis* and *Staphylococcus aureus* (Rucker *et al.*, 1992).

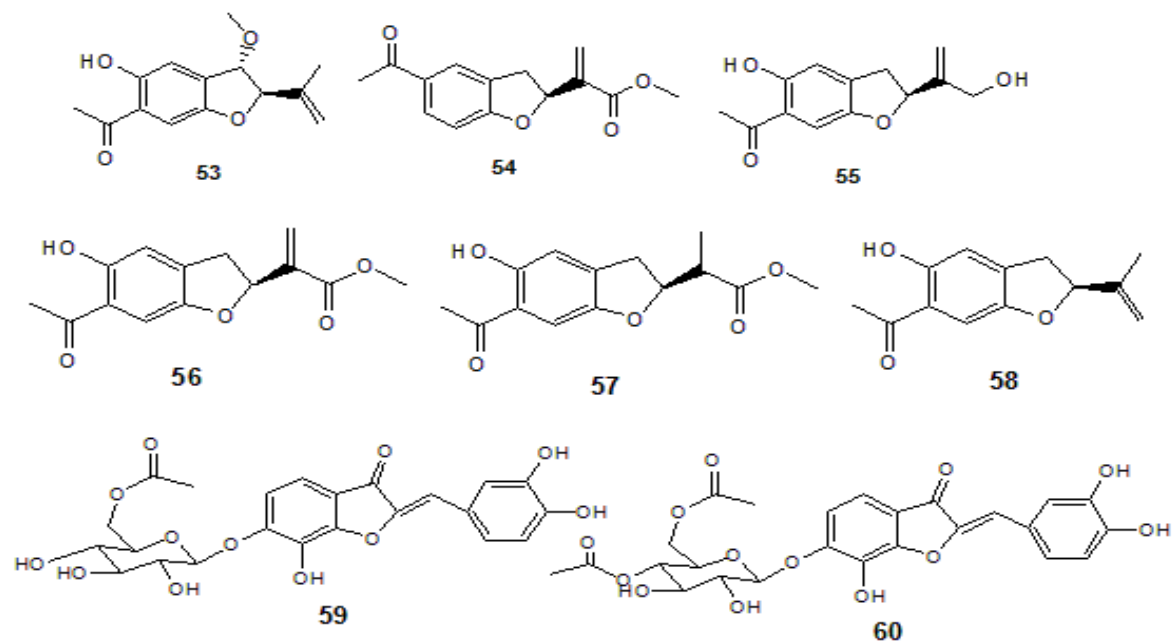


Figure 2.9: Dihydrobenzofurans and aurone flavonoids from *M. pyrifolia*

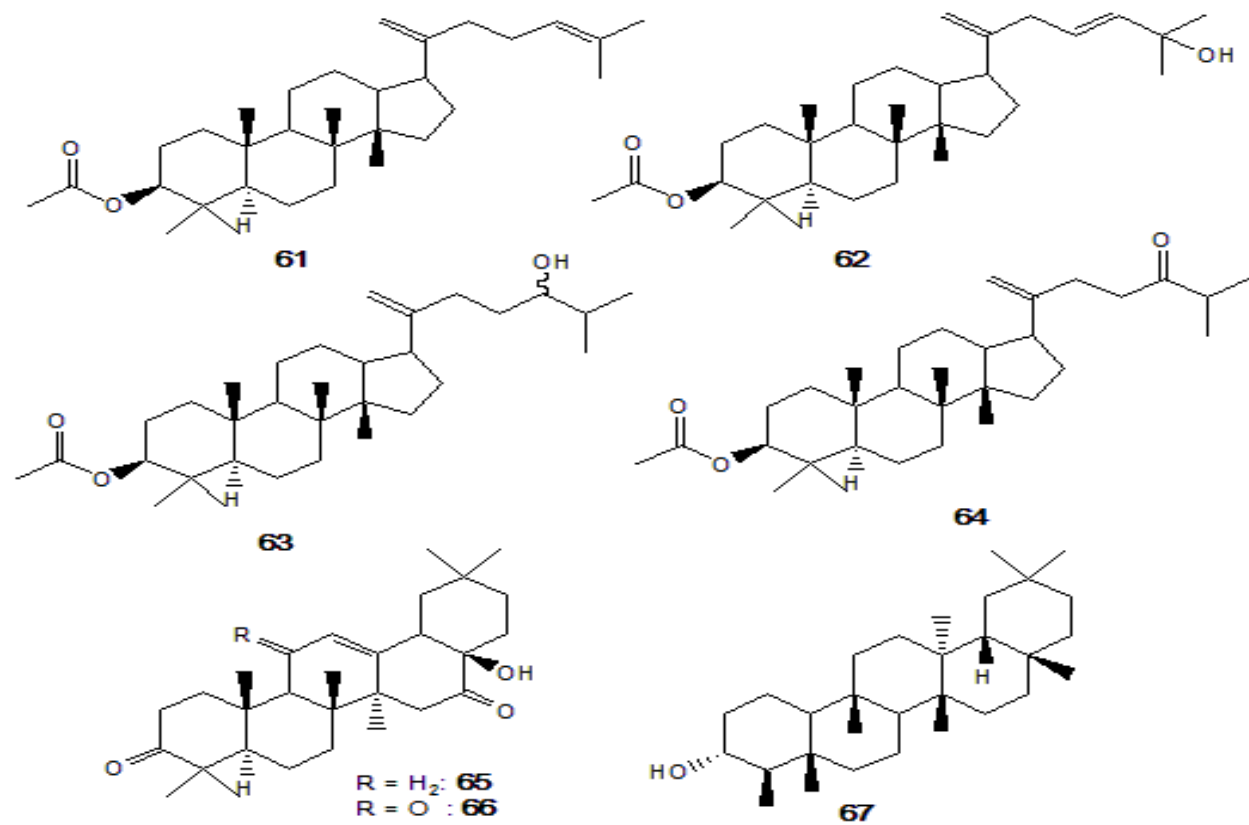


Figure 2.10: Triterpenoids from *Microglossa pyrifolia*

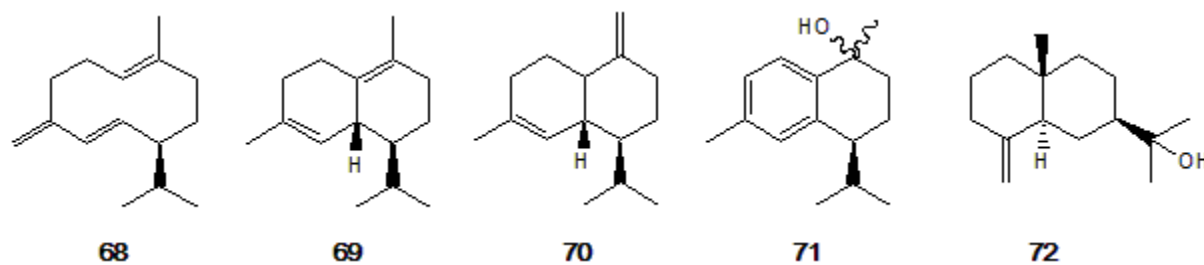


Figure 2.11: Sesquiterpenes from *Microglossa pyrifolia*

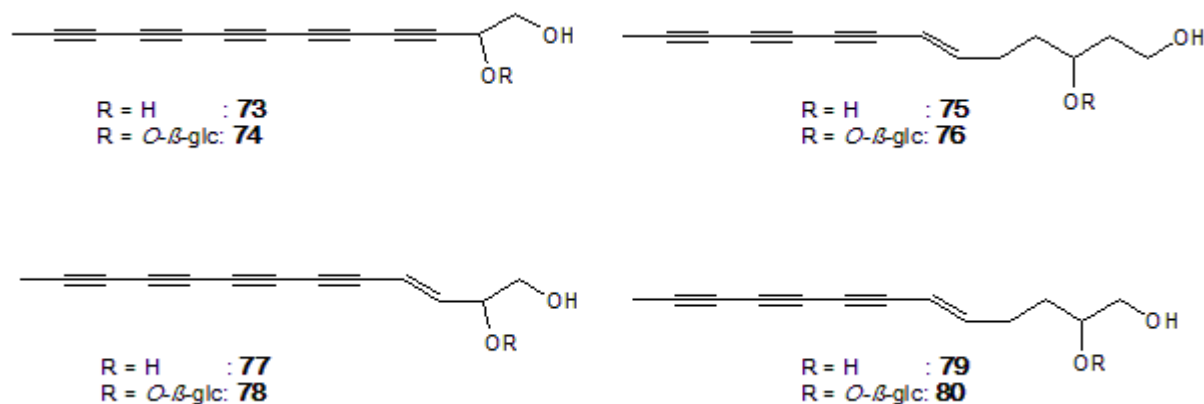


Figure 2.12: Acetylenes and acetylenic glucosides from *Microglossa pyrifolia*

2.7 Secondary metabolites

Secondary metabolites are metabolic intermediates or products, found as differentiation products in restricted taxonomic groups, not essential to growth and the life of the producing organism. They are biosynthesized from one or more general metabolites by a wider variety of pathways than are available in general metabolism. Secondary metabolites tend to be strain specific and have a wide range of chemical structures and biological activities (Rosenthal and Berenbaum, 1992).

The ability to synthesize secondary metabolites has been selected through the course of evolution in different plant lineage when such compounds address specific needs. Floral scent volatiles and pigments have evolved to attract insect pollinators and thus enhance fertilization. Synthesis of toxic chemicals has evolved to ward off pathogens and herbivores or to suppress the growth of neighboring plants (Bennet and Wallsgrave, 1994; Dixon *et al.*, 1996; Mitchel-Olds *et al.*, 1998; Trossat *et al.*, 1998; Harborne, 1999).

Chemicals found in fruits prevent spoilage and act as signals (in the form of color, aroma, and flavor) of the presence of potential rewards (sugars, vitamins and flavor) for animals that eat the fruit and thereby help to disperse the seeds. Other chemicals serve cellular functions that are unique to the particular plant in which they occur (e.g. resistance to salt or drought) (Dudareva and Pichersky, 2000; Mol *et al.*, 1998; Mert-Turk, 2002). Secondary metabolites are formed in only specific organisms, or groups of organisms, and are an expression of the individuality of species. They have a very restricted distribution than primary metabolites in the whole plant kingdom i.e. they are often found only in one plant species or a taxonomically related group of species.

High concentrations of secondary metabolites might result in a more resistant plant. Their production is thought to be costly and reduces plant growth and reproduction (Simms and Fritz, 1992; Karban and Baldwin, 1997; Stotz *et al.*, 2000; Siemens *et al.*, 2002). Therefore, defense metabolites can be divided into constitutive substances, also called prohibitions or phytoanticipins and induced metabolites formed in response to an infection involving *de novo* enzyme synthesis, known as phytoalexins (Van Etten *et al.*, 1994; Woodall and Stewart, 1999). Phytoanticipins are high energy and carbon consuming and exhibit fitness cost under natural conditions (Mauricio, 1998), but recognized as the first line of chemical defense that potential pathogens have to overcome. In contrast, phytoalexin production may take two or three days, as by definition first the enzyme system needs to be synthesized (Woodall and Stewart, 1999). The majority of secondary metabolites belong to one of a number of families, each of which has particular structural characteristics arising from the way in which they are built up in nature (biosynthesis). The three major classes of secondary metabolites are, polyketides and fatty acids, terpenoids and steroids, phenylpropanoids and alkaloids (Rosenthal *et al.*, 1992).

2.7.1 Surface exudates

Plants represent a rich source of nutrients for many organisms including bacteria, fungi, protists, insects, and vertebrates. Although lacking an immune system comparable to animals, plants have developed a stunning array of structural, chemical, and protein-based defenses designed to detect invading organisms and stop them before they are able to cause extensive damage (Brian and Gwyn, 2008).

Trichomes are simple or glanded epidermal appendages that occur on most plants. Glandular secreting trichomes are found on approximately 30% of vascular plants (Dell and McComb, 1978; Fahn, 1988), and they produce surface-accumulated exudates that usually contain hydrophobic isoprenoids and phenylpropanoids, the latter including flavonoids, phenolics, tannins, quinones, etc. (Wagner *et al.*, 1991). In some plants, amphipathic sugar esters are also commonly found in glandular trichome exudates (Wagner *et al.*, 2004). Such compounds have been associated with insect resistance in many plants, and pest resistance is often correlated with glandular trichome density (Spring, 2000). Two well-studied cases of glandular trichome-based insect resistance are found in the plant family Solanaceae. Sugar esters produced by tall glandular trichomes (TGSTs) of primitive tomato (*Lycopersicon pennellii*) and potato (*Solanum berthaultii*) species, and the diterpenoid cembratriene-ol produced by tobacco TGSTs, have been shown to inhibit aphid infestation (Goffreda *et al.*, 1990; Wang *et al.*, 2004). Antimicrobial activities of trichome exudate compounds (particularly monoterpenoids and sesquiterpenoids) have also been reported, but these are less studied than insect resistance (Wagner *et al.*, 2004).

Considerable recent research has focused on glandular secreting trichomes (GSTs) to understand and exploit their ability to secrete phytochemicals that might improve resistance to insects, microbes and herbivores. This research is also looking at the possibility of modifying gland metabolism towards improving properties of exudates (e.g. flavour and aroma in herbs) and allows commercial production of useful compounds (molecular farming). Plant sources are increasingly being exploited for new drug development, and there is increased interest in validating traditional medicines and herbal remedies (Wagner, 1991). Many of the phytochemical principals involved are surface secretions. Leaf surface protuberances include simple trichomes, glandular secreting trichomes, hydathodes, and other structures that extend above the epidermal plane, and the morphologies of these vary widely between plant species (Wagner *et al.*, 2004). In general, surface structures increase the distance between invaders and the leaf interior, and because of this they are mechanical deterrents to pathogens (Bennet and Wallsgrave, 1994). Trichomes were among the first anatomical features of plants to be recognized by early microscopists, and they have played a key role in plant taxonomy (Bennet and Wallsgrave, 1994). Simple trichomes are present on aerial surfaces of most angiosperms and on some gymnosperms and bryophytes (Uphof, 1962; Johnson, 1975).

In Angiosperms, trichomes may occur on leaves, petals, stems, petioles, peduncles and seed coats, depending on the species. GSTs are found on perhaps 30 % of vascular plants (Dell and McComb, 1978; Fahn, 1988). The amount of exudate produced by GSTs may reach 30 % of mature leaf dry weight (Dell and McComb, 1978). Terpenoids (mono- terpenes, sesquiterpenes, diterpenes and triterpenes) are clearly the most common and structurally diverse compound group found in GST exudates (Kellogg, 2001). Certain plants produce copious amounts of mixed-terpenoids, such as the terpenophenolic cannabinoids of *Cannabis* (Mahlberg and Kim, 1992). Flavonoids occur as aglycones in trichome exudates of a number of species, sometimes as crystalline deposits (Wang *et al.*, 2004). The occurrence of flavonoids as aglycones is not surprising since they are lipophilic and would be highly soluble in mixed exudates containing terpenoids.

In contrast, flavonoid glycosides, as accumulated in many plants, occur in vacuoles and chromoplasts. Examples of trichome-secreted phenolics include primin (*Primula sp.*), chlorogenic acid and rutin (*Solanum sp.*). The accumulation of toxic compounds at the surface allows their direct contact with insects, pathogens and herbivores. Thus trichome exudates are ideally positioned to provide a first line of defence against attacking organisms, perhaps providing time for activation of induced defences (Harborne, 1999; Wagner *et al.*, 2004).

2.7.2 Biosynthesis of Flavonoids

Flavonoids are synthesized by extension of *p*-hydroxycoumaroyl CoA with three molecules of malonyl CoA in a head-to-tail manner forming a tetraketide intermediate (Figure 12.14). This step is catalyzed by the enzyme chalcone synthase (CHS). The intermediate then folds and condenses further to give a chalcone. This reaction is the first step in flavonoid biosynthesis and is also catalyzed by chalcone synthase (CHS). Chalcone isomerase (CHI) is the second enzyme involved in the biosynthesis of flavonoids and it catalyzes the stereospecific intramolecular cyclization of the chalcones isoliquiritigenin and naringenin-chalcone into the flavanones (2*S*)-liquiritigenin and (2*S*)-naringenin, respectively (Winkel-Shirley, 2001).

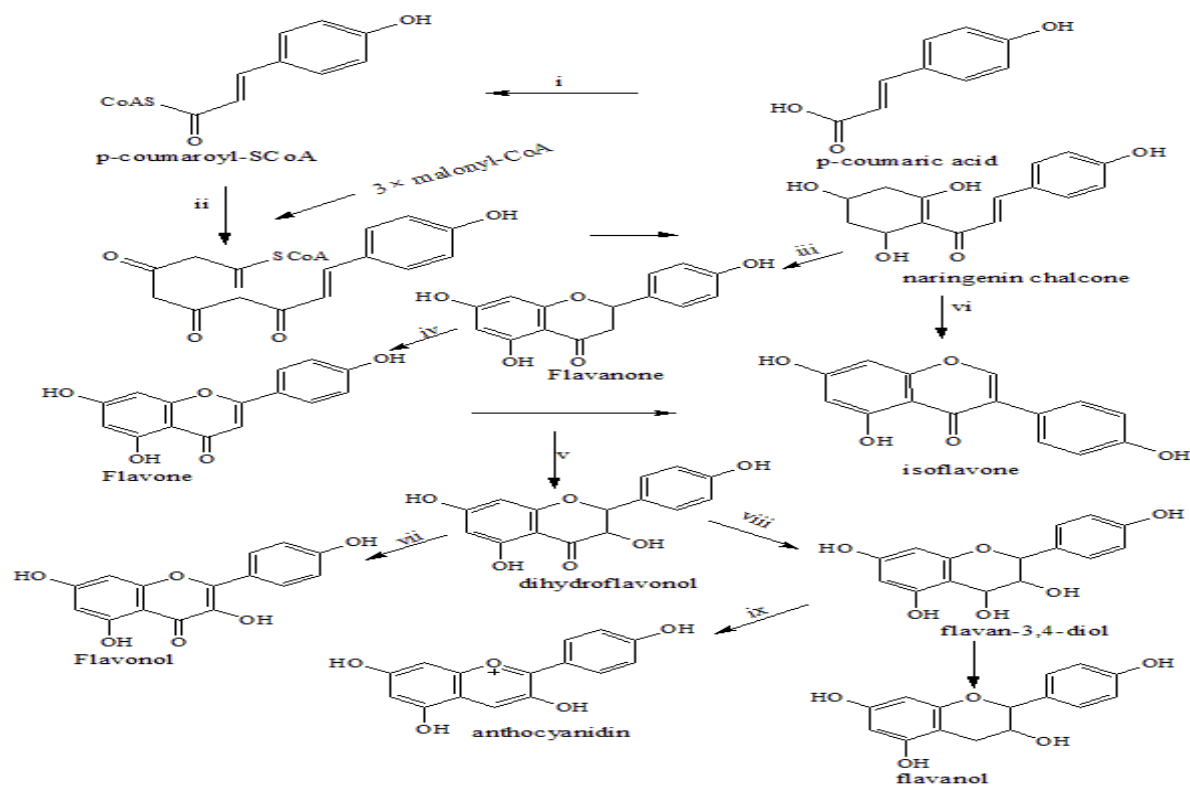


Figure 2.13: General Biosynthetic pathway of flavonoids

Enzymes: (i) p-coumarate-CoA ligase, (ii) chalcone synthase, (iii) chalcone isomerase, (iv) flavone synthase II, (v) flavanone 3- β -hydroxylase, (vi) isoflavone synthase, (vii) flavonol synthase, (viii) dihydroflavonol 4-reductase, (ix) anthocyanidin synthase (Davies and Schwinn, 2005)

2.7.3 Biosynthesis of Terpenoids

Terpenoids constitute the largest family of natural plant products with over 30,000 members (Sacchetti and Poulter, 1997; Dewick, 2002). Over 30,000 terpene hydrocarbons and oxidized or modified derivatives, called terpenoids, have been isolated to date from a wide variety of biological sources. These metabolites serve crucial roles in primary metabolism and ecological interactions of their organisms as well as display medically and economically valued characteristics (Sacchetti and Poulter, 1997).

For instance, several monoterpenes and sesquiterpenes are important volatile organic compounds that are important in the food and perfume industries (Schulz and Dickschat, (2007), whereas some microbial terpenoids are phytotoxic metabolites that have significant impact on agriculture (Collado *et al.*, 2007).

Terpenoids are classified by the homologous series of number of five carbon isoprene units in their structure: hemiterpenes C_5 (1 isoprene unit), monoterpenes C_{10} (2 isoprene units), sesquiterpenes C_{15} (3 isoprene units), diterpenes C_{20} (4 isoprene units), tri-terpenes C_{30} (6 isoprene units), tetraterpenes C_{40} (8 isoprene units), polyterpenes $(C_5)_n$ where 'n' may be 9–30,000 (McGarvey and Croteau, 1995). Monoterpenes (C_{10}) and sesquiterpenes (C_{15}), the constituents of essential oils, are important flavouring and fragrance agents in foods, beverages, cosmetics, perfumes, soaps and exhibit specific biological, pharmaceutical and therapeutical activities as well (Singh *et al.*, 1989; Mahmoud and Croteau, 2002).

Metabolically speaking, the biosynthesis of isoprenoids can be divided into three stages:

(i) formation of the C_5 isoprenoid unit, (ii) condensation of the C_5 units and chain elongation, and (iii) cyclization and modification of linear isoprenoid precursors. The extensive family of isoprenoid compounds is derived from two universal precursors, isopentenyl diphosphate (IPP) and dimethylallyl diphosphate (DMAPP). The mevalonate (MVA) pathway and the methylerythritol phosphate (MEP) pathway are the two independent pathways associated with the biosynthesis of these C_5 building blocks.

The classical mevalonate pathway (Fig: 2.14A and 2.16) begins with acetate molecules activated as acetyl-CoA, forming 3-hydroxy-3-methylglutaryl-CoA (HMG-CoA) which is reduced to mevalonate. Mevalonate is then activated via three phosphorylation reactions followed by a decarboxylation and elimination to yield IPP (Qureshi and Porter, 1981). DMAPP is generated subsequently from IPP in an equilibration reaction catalyzed by IPP-DMAPP isomerase that exploits a protonation-deprotonation mechanism (Leyes, *et al.*, 1999). Studies of this pathway have led to the development of the statins, or HMG-CoA reductase inhibitors, which have been widely used as therapeutics and preventive medicine for human cardiovascular diseases by lowering blood cholesterol levels (Stancu and Sima, 2001). An alternative mevalonate independent pathway, called the MEP pathway or deoxyxylulose phosphate pathway was discovered in certain bacteria and the plastids of both lower and higher plants in the 1990's (Rohmer, 1999; Eisenreich, *et al.*, 2004). The pathway starts with the condensation of pyruvate and glyceraldehyde-3-phosphate to form 1-deoxyxylulose-5-phosphate which undergoes a rearrangement coupled with a reduction resulting in MEP.

MEP is converted into its cyclic diphosphate, 2-C-methyl-D-erythritol-2, 4-cyclodiphosphate (MECDP), by sequential phosphocytidyl transfer and phosphorylation. MECDP is then reduced twice and finally transformed into IPP and DMAPP (Fig: 2.14 B and 2.15) (Eisenreich, *et al.*, 2004).

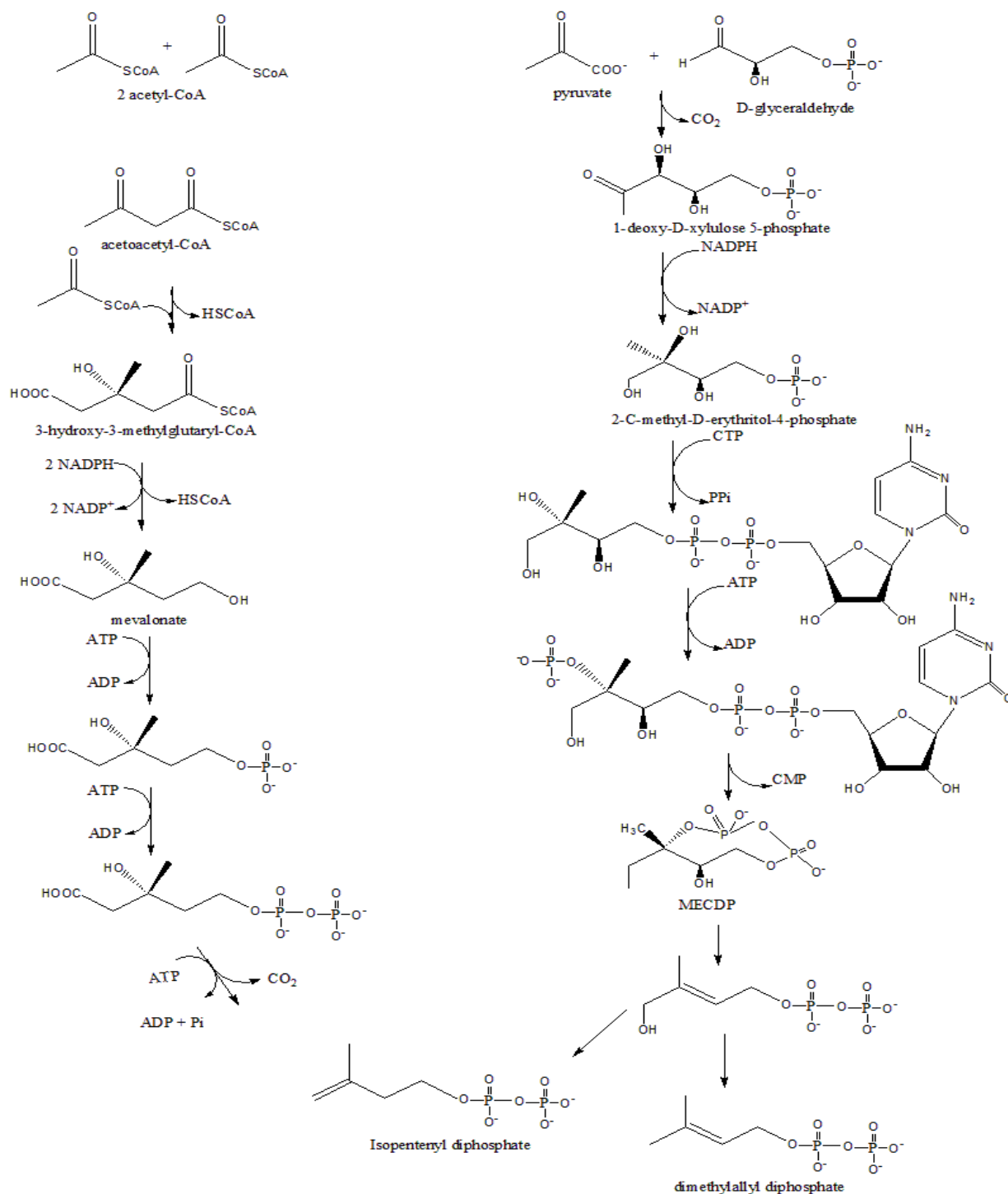


Figure 2.14: Two independent pathways for biosynthesis of IPP and DMAPP in plants

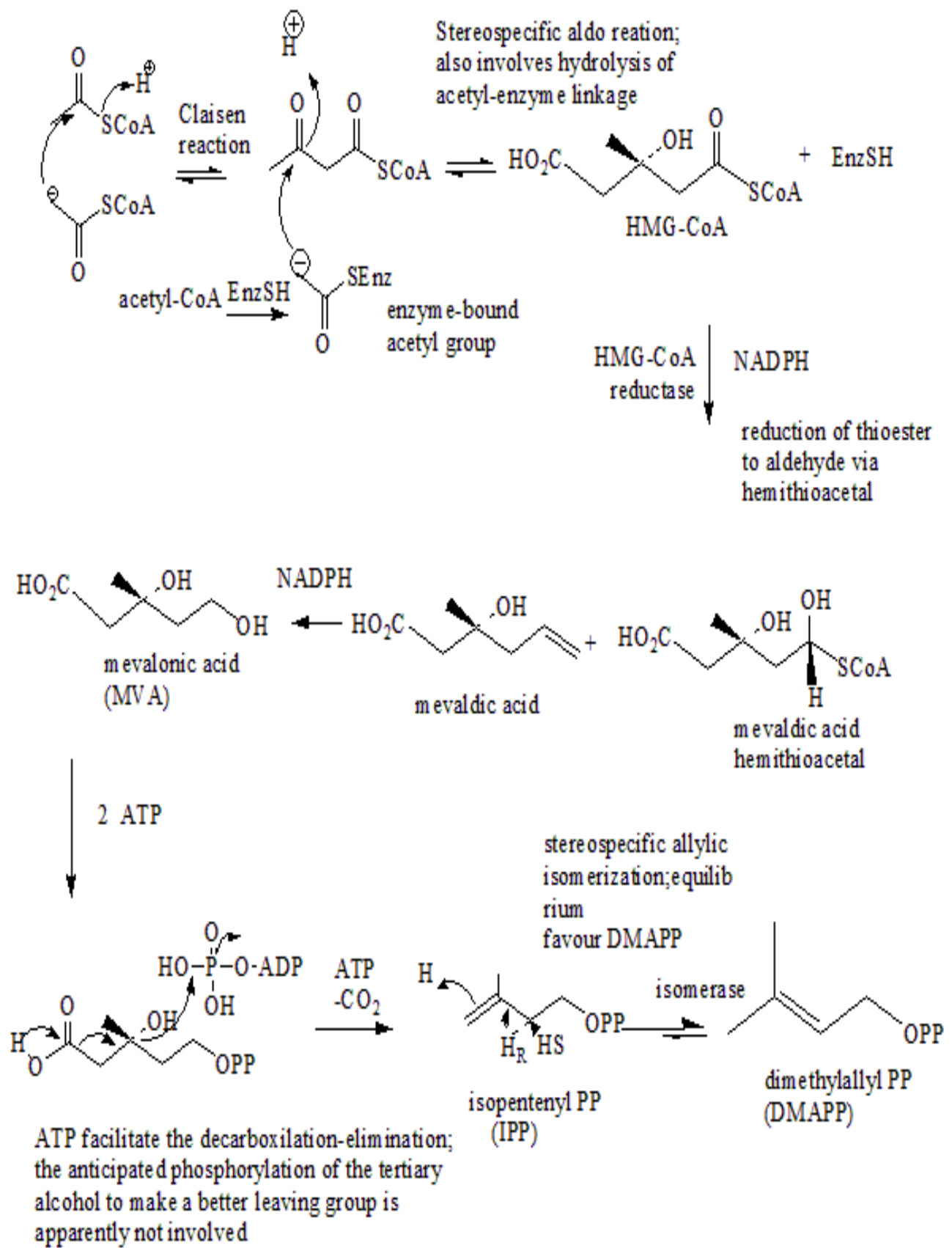


Figure 2.15: The mechanistic details of mevalonic acid pathway

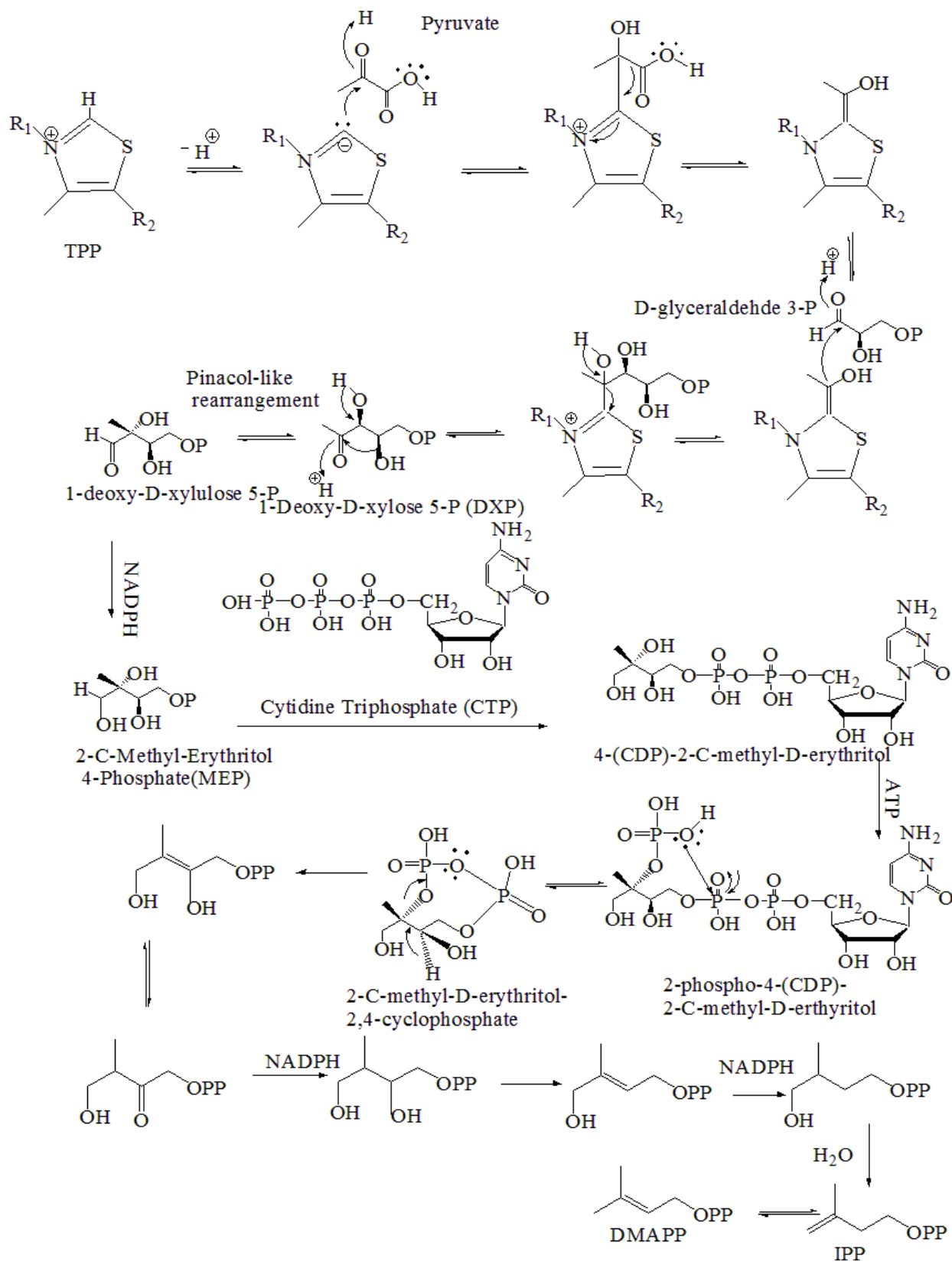


Figure 2.16: The mechanistic details of Deoxyxylulose phosphate (Methylerythrol Phosphate) pathway

All carbon skeletons of linear isoprenoids are assembled by repetitive head-to-tail condensation of IPP with an allylic diphosphate through the catalysis of prenyltransferases, a group of prenyl chain elongating enzymes (Takahashi and Koyama, 2006). Coupling of IPP to DMAPP by geranyl diphosphate synthase generates geranyl diphosphate (GPP, C₁₀) while addition of IPP to GPP by farnesyl diphosphate synthase gives farnesyl diphosphate (FPP, C₁₅), which can again couple to IPP by geranylgeranyl diphosphate synthase to yield geranylgeranyl diphosphate (GGPP, C₂₀), although FPP synthase and GGPP synthase can also utilize only IPP and DMAPP as the initial substrates in multistep elongation sequences via bound intermediates (Poulter and Rilling, 1981).

The reaction proceeds through an ionization-condensation-elimination mechanism, which involves an initial ionization of the allylic diphosphate ester substrate to form a charge delocalized carbocation which undergoes C4'-C1 coupling via the terminal double bond of IPP to generate a tertiary carbocation followed by deprotonation to complete the reaction (Fig.2.17). (Poulter and Rilling, 1978). The resulting acyclic intermediates are the key precursors from which monoterpenes (from GPP), sesquiterpenes (from FPP), and diterpenes (from GGPP) are derived. Terpenoid synthases, also called cyclases since the reaction products are most often cyclic, are the family of enzymes that catalyzes the conversion of the three acyclic branch point intermediates to the parent skeletons of the various monoterpene, sesquiterpene, and diterpene types and largely contributes to the great structural diversity encountered in this class of natural products. (Davis and Croteau, 2000). Reaction of terpenoid synthases may be considered as the intramolecular analogue of the intermolecular electrophilic coupling reaction catalyzed by the prenyltransferases, although the former case comprises far more diversity in type and mechanistic complexity (Davisson *et al* 1985). Terpenoid cyclases, responsible for catalyzing some of the most complex chemical reactions occurring in nature, are functionally soluble, modestly lipophilic and acidic enzymes with native sizes in the 35–80 kDa range; a divalent metal ion is the only cofactor required for enzyme activity, which exhibits Michaelis constants for the prenyl substrate that rarely exceed 10 μM and turnover numbers typically ranging from 0.03 to 0.3 s⁻¹ (Davisson *et al* 1985). About two-thirds of the carbon atoms of a linear terpenoid substrate undergo changes in bonding, hybridization, or configuration during the course of an elaborated cyclization cascade (Christianson, 2006).

Formation of a highly reactive carbocation intermediate is the initiation step of a terpenoid cyclase reaction which is either ionization-dependent or protonation-dependent.

Typical ionization-dependent cyclizations involve metal-triggered ionization of the prenyl diphosphate substrate yielding an allylic carbocation-pyrophosphate ion pair followed by an intramolecular attack from a double bond on the prenyl chain to generate a cyclic carbocation intermediate. Subsequent cationic transformations may involve further cyclizations and rearrangements, including hydride shifts, methyl migrations and Wagner-Meerwein rearrangements, which can create a potentially vast and diverse array of products. The reactions are ultimately terminated by quenching of the positive charge by deprotonation or capture of an external nucleophile such as water or the original pyrophosphate anion.

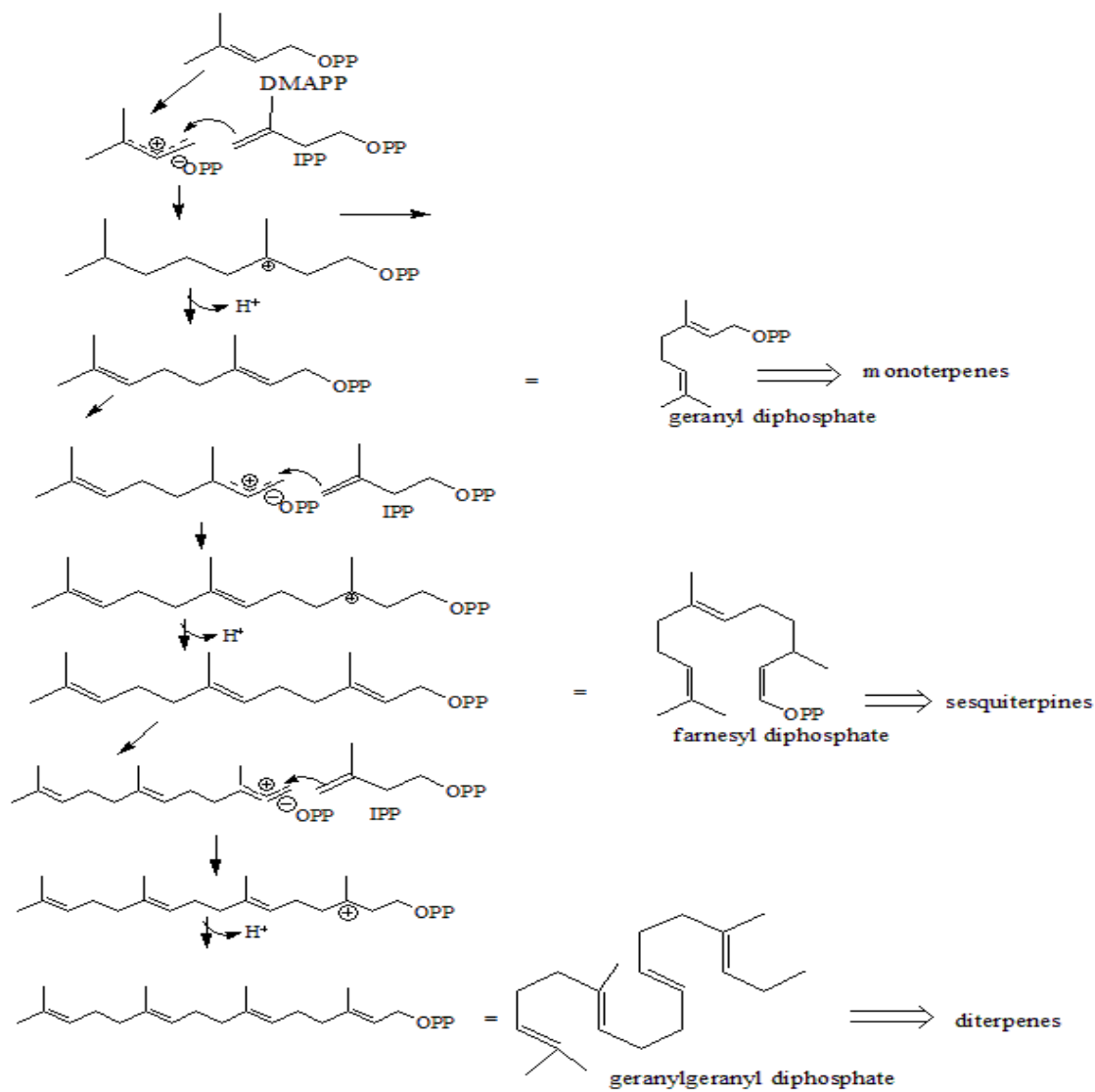


Figure 2.17: Chain elongations by prenyltransferases

CHAPTER THREE

MATERIALS AND METHODS

3.1 General

Column chromatography used Merck Silica gel 60 (0.063-0.200 mm) and Fluka Sephadex LH-20 as stationary phases; PTLC on Merck Silica gel 60 PF254+366, coated on glass plates (20 x 20 cm) to make 1.0 mm layers; Analytical TLC was carried out using factory prepared aluminum plates (0.25 mm) coated with Silica gel (60 F254, Merck) and spots visualized by observing under UV light at 254 or 365 nm, followed by spraying with 1% vanillin-H₂SO₄ spray reagent. These were carried out at the University of Nairobi, department of Chemistry.

1D data were acquired on Bruker Avance 600 MHz or Bruker BioSpin instrument 400 MHz for ¹H and Bruker Avance 100 MHz or Varian-Mercury 150 MHz for ¹³C, while 2D data were acquired on Bruker Avance 600 MHz or Bruker BioSpin instrument 400 MHz for HSQC and HMBC. NMR measurements were done in deuterated solvents using the residual undeuterated chemical shifts as internal standard. HREI-MS spectra were recorded on GC-MS TRACE DSQII single quadrupole mass.spectrometer. These were done at the National Center for Natural Product Research, School of Pharmacy, University of Mississippi.

3.2 Plant material

The aerial parts of *Microglossa pyrifolia* were collected from Ngong forest Nairobi in February 2010. A Voucher specimen was deposited at the University Herbarium, Department of Botany, University of Nairobi (No AAI/2010/10 *Microglossa pyrifolia*).

3.3 Extraction and isolation

Crude leaf surface extract was obtained by dipping about 400 g of fresh leaves into about 5 litres of ethyl acetate for a short period of 15 seconds thus avoiding the extraction of internal tissue components (Midiwo *et al.*, 1990). This process was repeated until the colour of ethyl acetate changed to yellowish at which fresh ethyl acetate was used. The surface extract was then filtered and evaporated off in a rotary evaporator under reduced pressure at a temperature not exceeding 45°C to give 190 g of yellowish black gummy extract.

Of this, 180 g portion of the extract was adsorbed on 200 g of silica gel and subjected to column chromatography over Silica gel (2 kg, 110 x 90 cm column), eluted with *n*-hexane/CH₂Cl₂ mixtures (100:0, 16 L; 9:1, 14 L; 4:1, 12 L; 3:2, 12 L; 1:1, 16 L; 1:3, 12 L; 0:100, 18 L) and CH₂Cl₂/MeOH mixtures (99:1, 14 L; 98:2, 12 L; 97.5:2.5, 12 L; 97:3, 10 L) to give a total of 127 fractions each 500 mL. The fractions were monitored by thin layer chromatography and similar fractions were combined to give a total of twelve fraction summarized in Table 3.1.

Table 3.1: Combined fractions and their weights

Fraction No	New fraction Code	Weight of Fraction (g)
1-3	1A	13.88
4-6	12B	13.28
7-16	12C	2.60
17-21	12D	5.87
22-25	12E	2.23
26-30	12F	5.08
31-36	12G	2.6
37-60	12H	3.17
61-69	12I	1.9
70-88	12J	7.68
89-103	12K	6.88
104-127	12L	16.75

Fraction 12E which was eluted with *n*-hexane/CH₂Cl₂ mixtures (100:0, 116 L) yielded epoxyisochiliolide lactone (**86**) (80 mg) which crystallized in 1:1 *n*-hexane/CH₂Cl₂. Fraction 12I (1.9g) was subjected to column chromatography over silica gel (20 g, 2.0 x 30 cm) eluted with *n*-hexane/CH₂Cl₂ mixture (9:1, 0.1 L; 4:1, 0.1 L; 7:3, 0.1 L; 1:1, 0.15 L; 1:3, 0.12 L; 0:100, 0.15 L) to give 36 fractions of 20 ml each. Combined fractions 22-29 (120 mg) were chromatographed over sephadex LH-20 (20 g, 2.5 x 30 cm; eluent: 1:1 CH₂Cl₂/MeOH) to give 8-acetoxyisochiliolide lactone (**87**) (210 mg).

Fraction 12L (16.76 g) was subjected to column chromatography on a silica gel column (200 g, 4×30 cm) eluted with increasing gradient of ethyl acetate in n-hexane (100:0, 1.6 L; 9:1, 1.4 L; 8.5:1.5, 1.2 L; 8:2, 1.2 L; 7.5:2.5, 1.6 L; 7:3, 1.2 L; 6.5:3.5, 1.8 L) to give 48 fractions each 20 mL.

These gave compounds **81** (80 mg), **82** (46 mg), **83** (35 mg) and **84** (52 mg) that crystallized from fractions 1-4, 5-6, 7-15 and 19-34 respectively. The other fractions contained mixtures of compounds with very close R_f values. Fraction 12K (6.88 g) was adsorbed onto 12 g of silica gel and subjected to column chromatography (120 g, 4×30 cm) eluted with increasing gradient of ethyl acetate in n-hexane to (100:0, 1.0 L; 9:1, 0.9 L; 8.5:1.5, 0.8 L; 8:2, 0.7 L; 7.5:2.5, 0.5 L; 7:3, 0.5 L; 6.5:3.5, 0.5 L) give thirty fractions of each 150 mL. Fractions 1-2 (58 mg) were combined and passed through sephadex LH-20 column (20 g, 2.5 x 30 cm) eluted with 1:1 CH₂Cl₂/MeOH and the fractionated eluents concentrated and re-crystallized in n-hexane/CH₂Cl₂ (1:1) to give compound **85** (47 mg).

3.4 *In vitro* anti-plasmodial assay

The *in vitro* anti-plasmodial activities were measured at the National Center for Natural Products Research of School of Pharmacy, University of Mississippi. All the compounds isolated (**81**, **82**, **83**, **84**, **85**, **86**, and **87**) were tested using the colorimetric assay method that determines the parasitic lactate dehydrogenase (pLDH) activity (Makler *et al.*, 1993; Samoylenko *et al.*, 2009). The assay was performed in 96-well microplate and included two *P. falciparum* strains [Sierra Leone D6 (chloroquine-sensitive) and Indochina W2 (chloroquine-resistant)].

For the assay, a suspension of red blood cells infected with *P. falciparum* (D6 or W2) strains (200 µL, with 2% parasitemia and 2% hematocrit in RPMI - 1640 medium supplemented with 10% human serum and 60 µg/mL amikacin) was added to the wells of a 96-well plate containing 10 µL of test samples at various concentrations. The plate was flushed with a gas mixture of 90% N₂, 5% O₂, and 5% CO₂, in a modular incubation chamber (Billups-Rothenberg, 4464 M) and incubated at 37 °C, for 72 h. Plasmodial LDH activity was determined by using Malstat™ reagent (FlowInc., Portland, OR). The IC₅₀ values were computed from the dose response curves generated by plotting percent growth against test concentrations. DMSO, artemisinin and chloroquine were included in each assay as vehicle and drug controls, respectively.

The selectivity index (SI) of anti-plasmodial activity was determined by measuring the cytotoxicity of samples towards mammalian cells (VERO; monkey kidney fibroblasts).

3.5 Cytotoxicity Assay

All the compounds isolated (**81**, **82**, **83**, **84**, **85**, **86**, and **87**) were tested. These were measured at the National Center for Natural Products Research of School of Pharmacy, University of Mississippi. The activity was determined against monkey kidney fibroblasts (VERO) obtained from the American Type Culture Collection (ATCC, Rockville, MD). The assay was performed in 96-well tissue culture-treated microplates. Cells were seeded at a density of 25000 cells / well and incubated for 24 hours. Samples at different concentrations were added and plates were again incubated for 48 hours. The number of viable cells was determined using Neutral Red (Borenfreund *et al.*, 1990). IC₅₀ values were determined from dose response curves of percent growth inhibition against test concentrations. Doxorubicin was used as a positive control, while DMSO was used as the negative (vehicle) control.

3.6 Antioxidant test

A preliminary anti-oxidant test was done by spotting the compounds (**81**, **82**, **83**, **84**, **85**, **86** and **87**) on a TLC plate followed by spraying with 0.2 mg/ml DPPH solution to view the active compounds which displayed white or yellowish spots on a purple background. The active compounds were quantitatively analysed by UV-VIS spectrometry method adopted from Hou *et al.*, 2002 with modifications made on the concentrations of the samples. For each compound, the concentration of the sample was varied by serial dilutions to give concentrations of 160, 80, 40, 20, 10, 5.0, 2.5, 1.25 and 0µg/ml while the concentration of DPPH was kept constant at 100 µg /ml (Figure 3.1). The reaction mixture consisted of adding 0.5 mL of sample, 3 mL of absolute ethanol and 0.3 mL of 100µg/mL DPPH radical solution in ethanol. These solutions were then measured for UV-VIS absorbance at DPPH absorbing wavelength (517 nm) half an hour after adding the DPPH.

The absorbance measured at each of these intervals were converted into percentages of scavenged DPPH radicals using the following equation

$$\% \text{ of scavenged DPPH} = \left(\frac{A_{\text{blank}} - A_{\text{sample}}}{A_{\text{blank}}} \right) \times 100\%$$

Where A_{blank} is the absorbance of DPPH solution without sample. The percentages of scavenged DPPH were then plotted against concentration of the compound to give graphs from which concentrations at half inhibition (IC_{50}) were determined. The tests were done in triplicates.



Figure 3.1: Samples with varied concentration of compounds but constant concentration of DPPH.

3.7 Antimicrobial Assay

The organisms were obtained from the American Type Culture Collection (Manassas, VA) and include the fungi *Candida albicans* ATCC 90028, *Candida glabrata* ATCC 90030, *Candida krusei* ATCC 6258, *Cryptococcus neoformans* ATCC 90113 and *Aspergillus fumigatus* ATCC 90906 as well as the bacteria *Staphylococcus aureus* ATCC 29213, methicillin-resistant *Staphylococcus aureus* ATCC 43300 (MRS), *Escherichia coli* ATCC 35218, *Pseudomonas aeruginosa* ATCC 27853 and *Mycobacterium intracellulare* ATCC 23068. Susceptibility testing was performed at the National Center for Natural Products Research, School of Pharmacy, University of Mississippi, using a modified version of the CLSI methods (Samoylenko *et al.*, 2009). All the isolated compounds were tested. Drug controls ciprofloxacin (ICN Biomedicals, Ohio) for bacteria and amphotericin B (ICN Biomedicals, Ohio) for fungi were included in each assay. The number of microbial cells of 10^7 CFU/ml was prepared by using McFarland standards.

This was done by adjusting the absorbance at 550 nm to 0.125 of microbial cells suspension with sterile saline and making an additional 10-fold diluted cells suspension of it with sterile saline. The number of microbial cells was 10^7 CFU/ml. The compounds, drug controls ciprofloxacin (ICN Biomedicals, Ohio) for bacteria and amphotericin B (ICN Biomedicals, Ohio) for fungi were serially diluted to obtain concentrations of 64, 32, 16, 8, 4, 2, 1, 0.5, 0.25, 0.12, 0.06 $\mu\text{g} / \text{ml}$ in Mueller-Hinton broth. Of the serially diluted compounds and drug controls, 180 μl of each were dispensed to each well and then 10 μl of the cell suspension (10^7 CFU/ml) was inoculated to each well (final cell density: 10^4 CFU/ml). The plate was incubated at 47°C for 6 hours, after which 10 μl of Coloring reagent was added to each well and the microplate incubated for an additional 2 hours. Absorbance was measured at 450 nm using the microplate reader

3.8 Anti-leishmanial Assay

Antileishmanial activity of the compounds was tested *in vitro* on a culture of *Leishmania donovani* promastigotes at the National Center for Natural Products Research of School of Pharmacy, University of Mississippi. In a 96 well microplate assay compounds with appropriate dilution were added to the *Leishmania* promastigotes culture (2×10^6 cells/mL). The plates were incubated at 26°C for 72 hours and growth of *Leishmania* promastigotes was determined by Alamar blue assay (Mikus and Steverding, 2000). All the compounds isolated were tested. Pentamidine and amphotericin B were used as standard antileishmanial agents. 10^6 cells/ml promastigotes were seeded in 96-well flat-bottom plates in final volume of 200 μl . Various concentrations of experimental compounds and pentamidine and amphotericin B drugs were made in three-fold dilutions and 180 μl of each were then added to triplicate wells and plates incubated for 72 h at 22°C . At the end of incubation, cell viability was measured by counting the number of motile cells using a haemocytometer. For each drug, the concentration-response curve was plotted from which IC_{50} values (the concentration of the agent that reduce cell viability by 50% compared to controls) were determined. All experiments were repeated at least three times. IC_{50} values for each compound were computed from the growth inhibition curve.

CHAPTER FOUR

RESULTS AND DISCUSSION

4.1 Secondary metabolites isolated from surface exudates of *Microglossa pyrifolia*

Chromatographic separation of the surface exudates of *Microglossa pyrifolia* led to the isolation and characterization of five flavonoids and two diterpenoids, of which three (**82**, **84** and **86**) are new compounds.

4.1.1 Flavonoids

In this study five flavonoids were isolated and successfully characterized. Of these flavonoids, four were trihydroxylated and methoxylated flavones while one was trihydroxylated and methoxylated flavanone. The characterization of these compounds are discussed as follows.

4.1.1.1 5, 7, 4'-Trihydroxy-3, 8, 3'-trimethoxyflavone (81)

Compound **81** was isolated as a yellow solid with HR-EIMS molecular ion at m/z 359.1009 $[M-H]^+$ (Appendix1E) consistent with the molecular formula $C_{18}H_{16}O_8$. The HR-EIMS fragmentation pattern (Figure 4.2) had fragment at m/z 344 (M-CH₃) and the C-ring degraded by cleavage of bonds 0, 2 and produced retro Diels-Alder (RDA) fragment $^{0.2}A^+$ which lost a CO to produce a peak at m/z 154.98. This fragment then lost a CH₃ resulting to a peak at m/z 112.99 (Tsimogiannis *et al.*, 2007) (Appendix1E). Another cleavage of ring A resulted in a peak at m/z 248.97. Its ¹H NMR (Table 4.1) (Appendix 1A) revealed the presence of four aromatic protons of which one is singlet at δ_H 6.27 on ring A and the other three showed an ABX spin system [δ_H 6.98 (1H, *d*, *J* = 8.4 Hz), δ_H 7.26 (1H, *dd*, *J* = 8, 1.6 Hz) and δ_H 7.66 (1H, *d*, *J* = 2 Hz)]. The singlet at δ_H 6.27 showed HMBC correlations with two oxygenated carbons δ_C 157.4 and 156.3 (Appendix1B) and long range HMBC correlation with the α,β -conjugated ketone (δ_C 178.5, C-4) as shown by the blue and yellow arrows respectively in figure 4.1, suggesting that the proton be assigned to H-6. In agreement with this, 5-hydroxy group did not correlate with any of the carbons carrying the methoxyl groups and hence any of the methoxyl could not be placed at position C-6. This led to exclusion of the possibility of 5, 6, 7-oxygenation pattern in ring A. According to the HSQC spectrum, we had three methoxyl groups attached to carbons resonating at δ_C 55.9, 60.1 and 60.4.

The downfield chemical shift of one of the hydroxyl group proton at δ_H 12.32 was due to its *peri* position to the carbonyl carbon and hence the hydroxyl group was at C-5. The HMBC correlations between H-6 and C-10 (δ_C 104.47) and a weak 2J interaction between H-6 and C-5 (δ_C 157.4) were all evident from HMBC spectrum (Appendix 1D).

The presence of two oxygenated carbons at δ_C 149.0 and 150.2 was in agreement with 1,2 oxygenation pattern in ring B of flavone skeleton and the three aromatic protons with ABX spin pattern showed HMBC cross peaks with these oxygenated carbons (Table 4.1) which was in agreement with the placement of the two oxygenated carbons at C-3',4' positions.

One of the methoxyl protons at δ_H 3.80/ δ_C 55.2(Appendix 1C) showed HMBC correlation with one of the above oxygenated carbons at δ_C 149.0 and hence assigned to be at position C-3'. If ring A and B are tri- and di-substituted respectively, then the remaining methoxyl group δ_H 3.85/ δ_C 60.1 which showed HMBC correlation with the carbon at δ_C 138.2, is at C-3 position. Thus, the above NMR data in comparison with literature NMR data (Horie *et al.*, 1998) suggested that the compound is 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) as shown in figure 4.1.

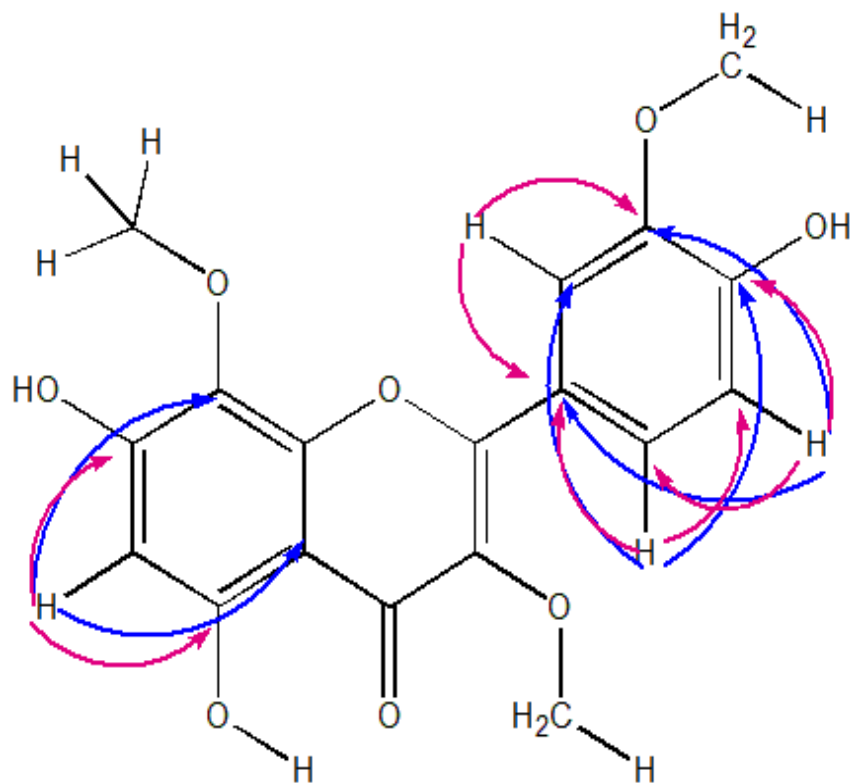


Figure 4.1: Structure of 5, 7, 4'-Trihydroxy-3, 8, 3'-trimethoxyflavone (**81**)

Table 4.1: $^1\text{H}/^{13}\text{C}$ NMR (400/100 MHz) of Compound (**81**)

Position	δ_{H}	δ_{C}	HMBC
2	-	155.6	-
3	-	138.2	-
4	-	178.5	-
5	-	157.4	-
6	6.27, <i>s</i>	99.2	C-4,5,6,7,8,10
7	-	156.3	-
8	-	127.9	-
9	-	147	-
10	-	104.5	-
1'	-	122.5	-
2'	7.66 (<i>d</i> , $J=2$)	112.0	C-1',3',4',5',6'
3'	-	149.0	-
4'	-	150.2	-
5'	6.98 (<i>d</i> , $J=8.4$)	116.2	C-1',3',4',5',6'
6'	7.26 (<i>dd</i> , $J=8, 1.6$)	121.5	C-1',2',4',5',6'
3'-OCH ₃	3.80, <i>s</i>	55.2	C-3'
3-OCH ₃	3.82, <i>s</i>	60.4	C-3
8-OCH ₃	3.85, <i>s</i>	60.1	C-8,9
5-OH	12.33, <i>s</i>	-	C-5,6,10

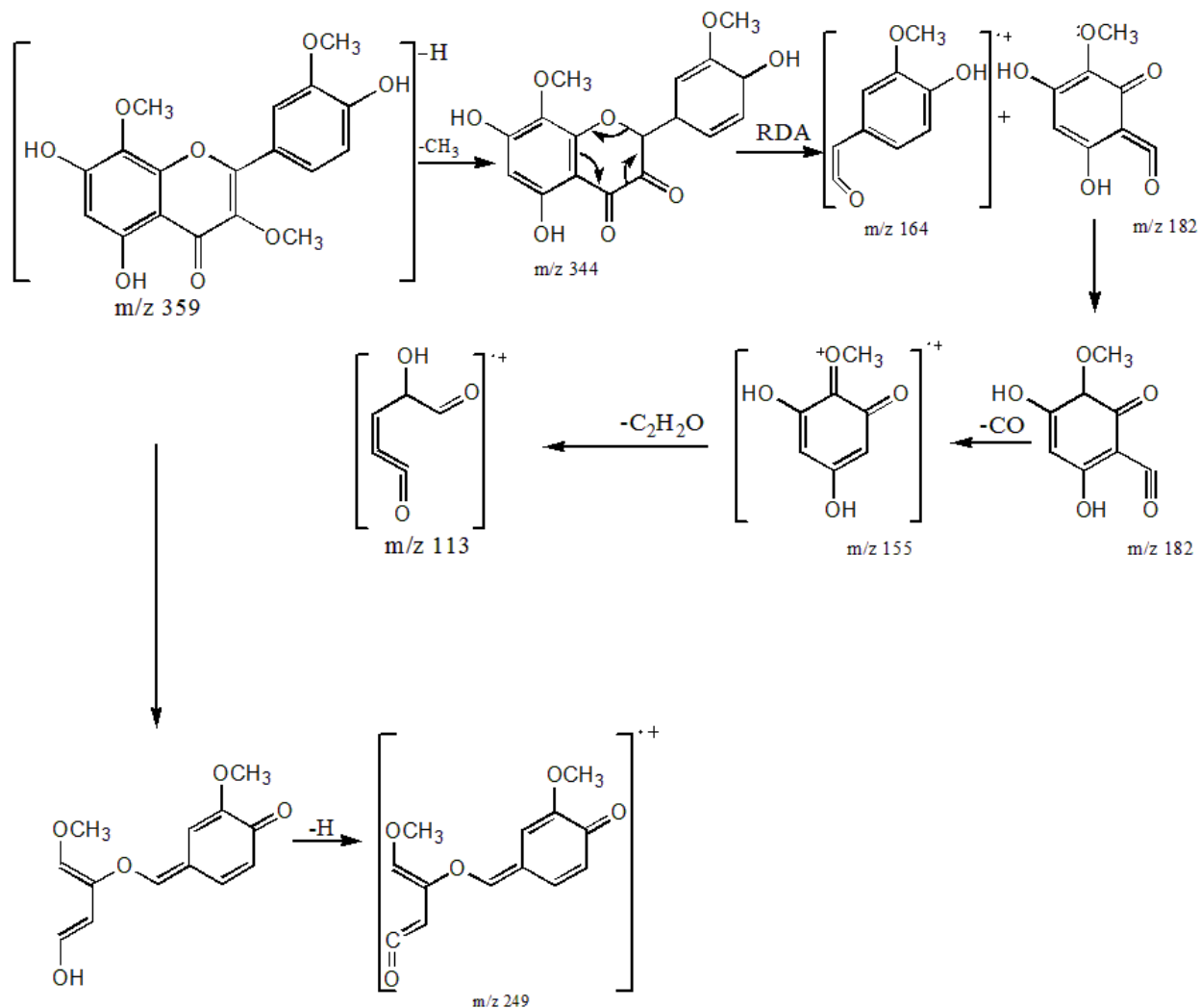


Figure 4.2: Fragmentation pattern for 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**)

4.1.1.2 5, 7, 4'-Trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**)

Compound **82** was isolated as a yellow solid and gave a molecular ion peak of 389.102 on HR-EIMS analysis (Appendix2E). The HR-EIMS fragmentation pattern (Figure 4.4) showed fragments at m/z 374 $[\text{M}-\text{CH}_3]^+$ and m/z 290.95 $[\text{M}-3\text{CH}_3-\text{CO}-\text{CHO}+\text{H}]^+$.

The fragment at m/z 290.95 lost a CH_2COH to form a peak at 248.96, which then underwent a series of cleavages to form a fragment at m/z 112.98. (Tsimogiannis *et al.*, 2007) (Appendix2E). Its $^1\text{H-NMR}$ (Table 4.2) displayed the presence of a singlet at δ_{H} 6.27 (1H, s) and δ_{H} 7.42 (2H, s). The singlet proton at δ_{H} 6.27 showed HSQC correlation (Appendix 2C) with carbon with δ_{C} 99.2 and HMBC correlations with carbon δ_{C} 104.4, 127.9, 157.5 and δ_{C} 178.4 (Appendix2B) suggesting that this proton be assigned to H-6 (Appendices 2C and 2D).

Furthermore, the above spectroscopic data suggests that ring A of flavone skeleton is tri-substituted.

The presence of two protons appearing as singlet at δ_H 7.42 suggest that there is symmetry in ring B of the flavone skeleton. The presence of δ_C 138.4 and 139.4 suggested the possibility of a methoxy placed at C-3 and a 1, 2, 3 tri-oxygenated pattern in ring B. The methoxy resonating at δ_H 3.81/ δ_C 60.1 showed HMBC correlation with the quaternary carbon at δ_C 138.4 suggesting that ring B is tri-substituted and taking into consideration the symmetry and HMBC correlation of the two protons with δ_C 106.1, 120.1, 139.4 and 155.3 suggesting that the two protons (at δ_H 7.42) belong to H-2' and 6'. Furthermore, the two methoxyl of ring B were placed on C-3' and C-5'. Thus, the above spectroscopic NMR data suggests that the compound is 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**) as shown in figure 4.3. This compound was found to be a new natural product.

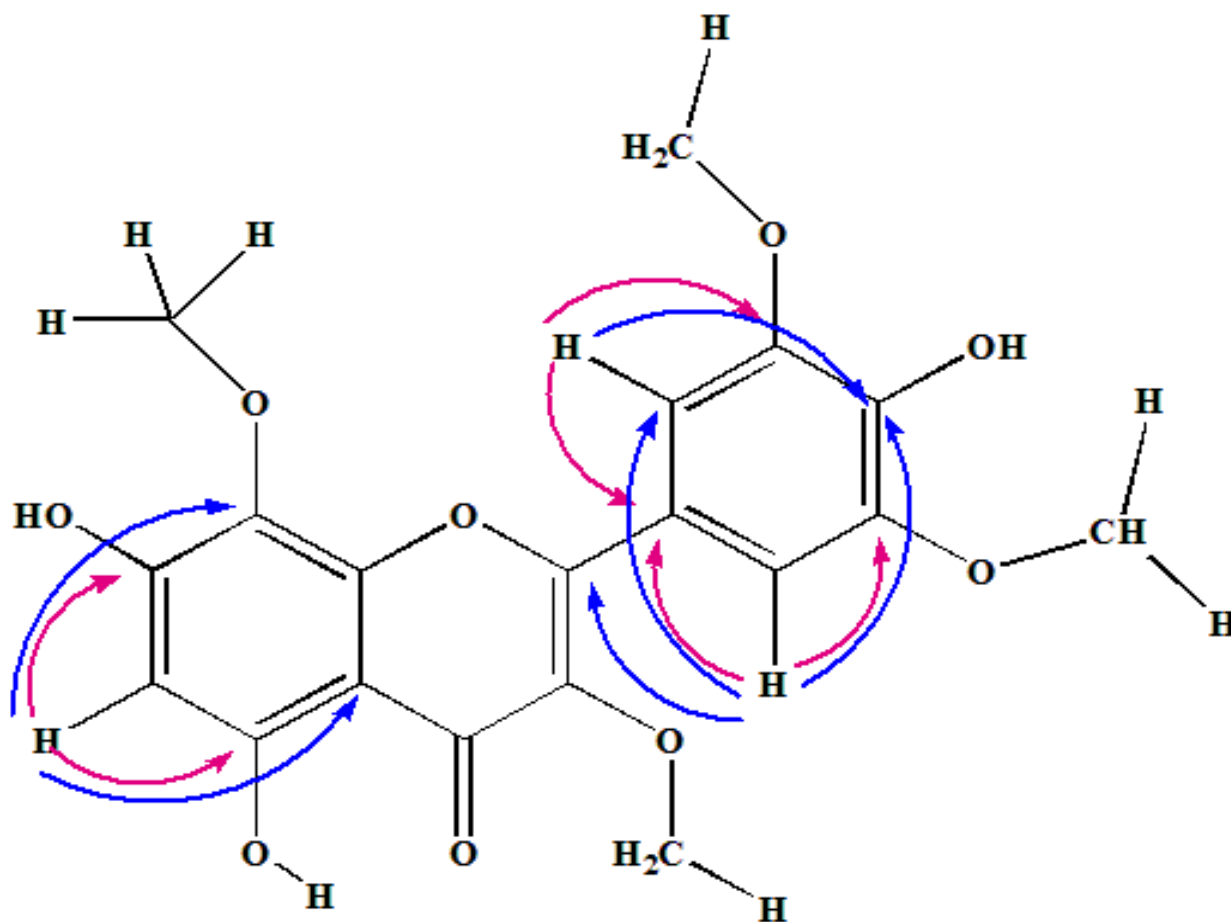


Figure 4.3: Structure of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**).

Table 4.2: $^1\text{H}/^{13}\text{C}$ NMR (400/100 MHz) of Compound (82).

Position	δ_{H}	δ_{C}	HMBC
2	-	155.3	-
3	-	138.4	-
4	-	178.5	-
5	-	157.5	-
6	6.26, <i>s</i>	99.2	C-4,5,6,7,8,10
7	-	156.3	-
8	-	127.9	-
9	-	148.9	-
10	-	104.4	-
1'	-	120.0	-
2'	7.42, <i>s</i>	106.1	C-1',2',3',4',6'
3'	-	148.2	-
4'	-	139.4	-
5'	7.42, <i>s</i>	148.2	-
6'	-	106.1	C-1', 2',4',5',6'
5-OH	12.39(5-OH)	-	C-5,6,10
3',5'-OCH ₃	3.82, <i>s</i>	58.2	C-3',5'
3-OCH ₃	3.81, <i>s</i>	60.1	C-3

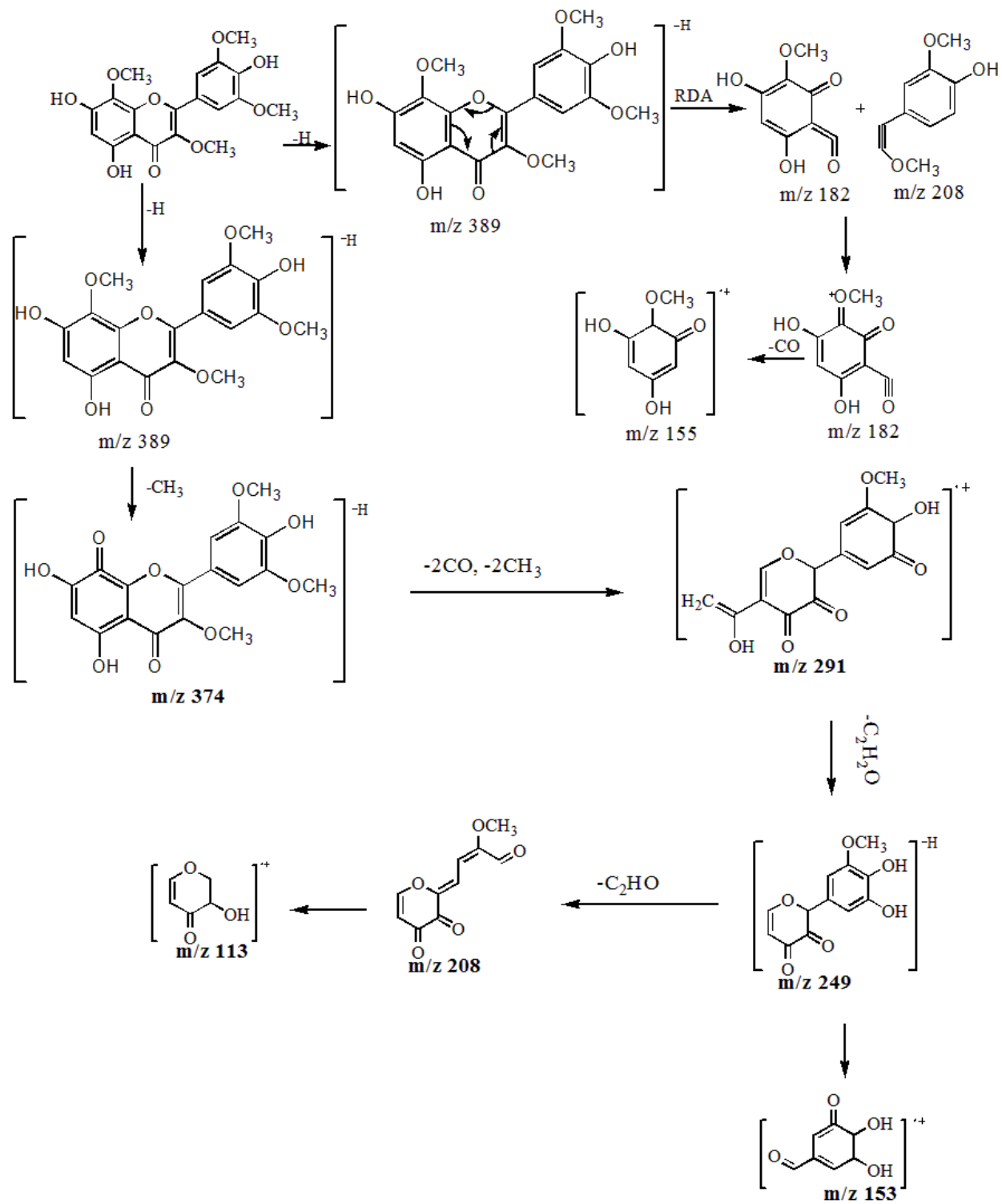


Figure 4.4: Fragmentation pattern for 5, 7, 4'-trihydroxy-3, 8, 3',5'-tetramethoxyflavone (82)

4.1.1.3 5, 3', 4'-Trihydroxy-7-methoxyflavanone (83)

Compound **83** was isolated as a light yellow solid and gave a molecular ion peak of 303.08 [M+H]⁺ on HR-EIMS analysis ((Appendix 3E). The protonated flavanone dehydrated to give [M+H-H₂O]⁺ (*m/z*= 253) and also may have lost the whole B-ring to produce a [M+H-B-ring]⁺ (*m/z*= 192) (Figure 4.6) (Hughes *et al.*, 2001).

The HRMS fragmentation pattern (Figure 4.6, Appendix 3E) further confirmed the structure with molecular ion peak at *m/z* 303.08 (C₁₆H₁₄O₆), a loss of a the C-ring and methyl giving a peak at *m/z* 191.08, loss of a methoxyl and water molecules giving a fragment at *m/z* 253.06

Its ¹H-NMR (Table 4.3) (Appendix 3A) displayed a pair of doublet aromatic protons at δ_H 6.08 (1H, *d*, *J*=2.0 Hz) and δ_H 6.06 (1H, *d*, *J*=2.0 Hz) being placed meta to each other. Based on HMQC correlations (Appendix 3C), these two protons belonged to δ_C 94 and δ_C 95 respectively. The proton at δ_H 6.06 showed HMBC correlations with the quaternary carbons at δ_C 103.0, 163.3 and 167.8 and the carbonyl carbon (197.4) (Appendix 3B) and hence the two protons were assigned to H-6 (δ_H 6.06) and H-8 (δ_H 6.08). The ¹H NMR further revealed that ring B of flavanone skeleton is di-substituted. The presence of two aromatic protons at δ_H 6.07 (1H, *d*, *J*=8.0) and 6.77(1H, *d*, *J*=8.0) coupling to each other and showing HMBC correlations with C-5' and C-6' suggest that ring B is di-substituted at C-3' and C-4'. Based on HSQC correlations (Appendix 3c), two diastereotopic protons at δ_H 3.23(1H, *dd*, *J*=17.2Hz, 12.8Hz) and δ_H 2.71 (1H, *dd*, *J*=17.2Hz, 3.2Hz) were assigned to carbon at δ_C 42.6 (C-3). Likewise, the proton at δ_H 5.41 (1H, *dd*, *J*=12.8Hz, 3.2Hz) was assigned to carbon at δ_C 79.1 (C-2). The HMBC correlations further revealed that the methylene protons (H-3) and the methine proton (H-2) showed HMBC correlations with the carbonyl carbon (C-4) and a quaternary carbon δ_C 129.7 (C-1'). The ¹³C NMR (Table 4.3) spectrum showed two oxygenated carbons in a 1,2-dioxygenation pattern at δ_C 145.6 and 146.2 and the HMBC correlation of all the three protons (ring B) with these two carbons further support that ring B is di-oxygenated at C-3' and C-4' positions. Thus, based on the above spectroscopic evidence the compound was identified to be 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**). The NMR data matched those reported in literature (Ibrahim *et al.*, 2003).

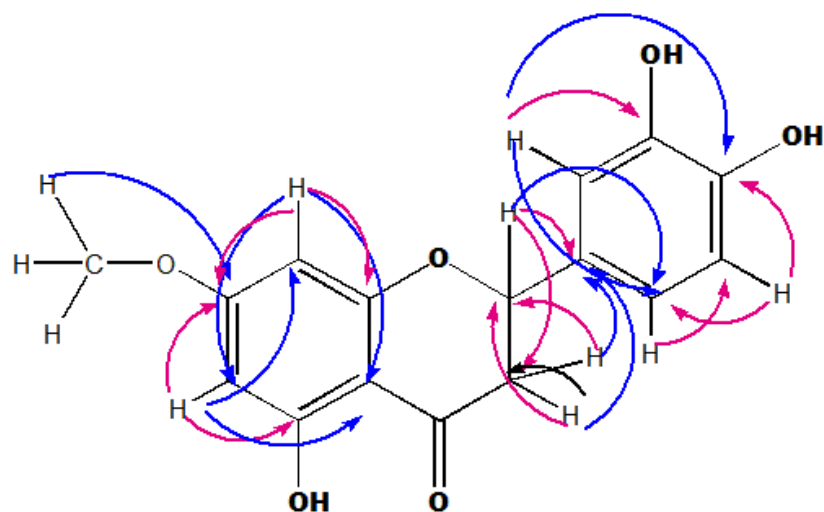


Figure 4.5: Structure of 5, 3', 4'-Trihydroxy-7-methoxyflavanone (**83**)

Table 4.3: ¹H/¹³C NMR (400/100 MHz) of Compound (**83**)

Position	δ_{H}	δ_{C}	HMBC
2	5.41, <i>m</i>	79.1	C-2',6',3,1'
3	2.71, 3.22, <i>dd</i>	42.6	C-2,3,1',6'
4	-	197.4	-
5	-	163.3	-
6	6.07 (<i>d</i>)	95.0	C-4,5,6,7,8,9,10
7	-	167.8	-
8	6.08 (<i>d</i>)	94.2	-
9	-	163.6	-
10	-	103.0	-
1'		129.7	-
2'	6.77	114.8	C-2',3',4', 6'
3'		146.2	-
4'		145.6	-
5'		115	C-5',6',4'
6'		118	C-1', 2', 5',6'
5-OH	12.11(5-OH)	-	C-5
7-OCH ₃	3.78, <i>s</i>	56.2	C-7

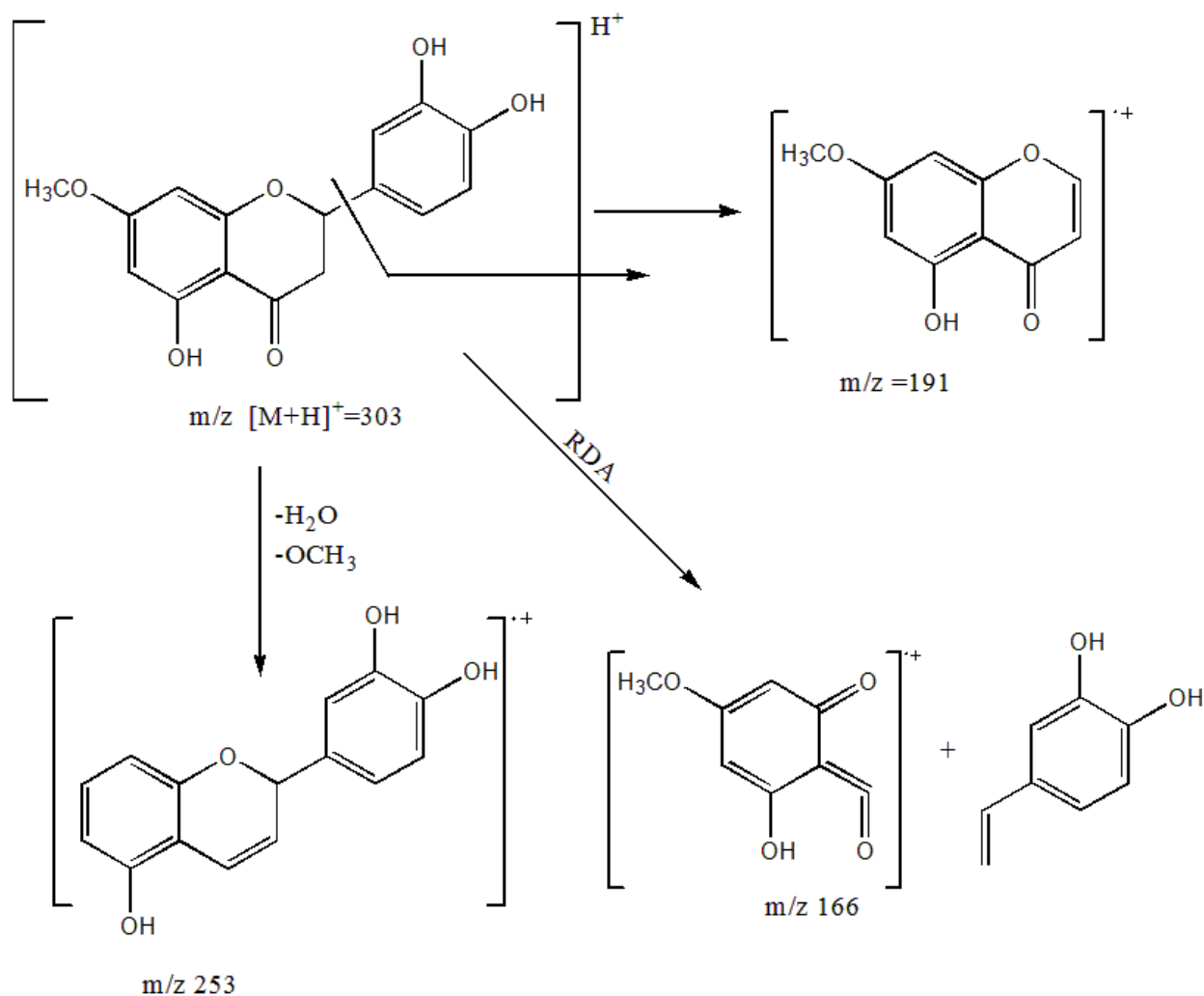


Figure 4.6: Fragmentation pattern for 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**)

4.1.1.4 5, 7, 3'-Trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**).

Compound **84** was isolated as yellow solid. Its HR-EIMS molecular ion (Appendix 4E) was observed at m/z 389.12 $[M-H]^+$ corresponded to the molecular formula of $C_{19}H_{18}O_9$. The HR-EIMS fragmentation pattern (Figure 4.8) had fragment at m/z 359 $[M-CH_3]^+$ and the C-ring degraded by cleavage of bonds 0,2 to produce RDA fragment $^{0,2}A^+$ which subsequently lost a CO to produce a peak at m/z 154.98. The other peaks were at m/z 248.98 and m/z 112.99 which were due to a series of cleavages (figure 4.8). Its 1H NMR (Appendix 4B and Table 4.4) gave seven signals of which three (δ_H 7.35, 7.21, 6.14) were singlet aromatic protons (Appendix 4A) and four (δ_H 3.84, 3.79, 3.76) were attributed to methoxyl groups (Appendix 4C).

Its ^{13}C NMR (Table 4.4) revealed nineteen signals of which fifteen were attributed for a flavonoid basic skeleton whereas the remaining four are methoxyl groups.

HMBC (Appendix D) correlation of the singlet proton at δ_{H} 6.14 with C-4, C-5, C-7, C-8, and C-10 suggested that this proton belong to H-6 and carbons 5, 7 and 8 are oxygenated quaternary carbons. The HMBC correlation (Appendix 4D) of the two methoxyl groups with oxygenated carbons at δ_{C} 129.1 (C-8) and δ_{C} 138.5 (C-8) suggested that the oxygenated quaternary carbons at C-3 and C-8 have methoxy substituents.

The presence of two singlet protons at δ_{H} 7.35 and δ_{H} 7.21, suggested that ring B of flavones skeleton is tri-substituted. The two singlet aromatic protons at δ_{H} 7.35/ δ_{C} 110.2 and δ_{H} 7.21/ δ_{C} 103.5 were attributed to H-4' and H-6' based on HMBC correlations of H-4' (δ_{H} 7.35) with carbons C-2', C-3', C-5', C-6' and H-6' (δ_{H} 7.21) with C-1', C-2', C-4' and C-5'. Furthermore, the HMBC correlation of the two methoxyl groups at C-2' and C-5' allowed the assignment of the two methoxyl groups at C-2' and C-5' positions. Thus, based on the above spectroscopic evidence the compound was identified to be 5, 7, 3'-trihydroxy-3, 8, 2', 5'-tetramethoxyflavone (**84**). This compound was found to be a new natural product.

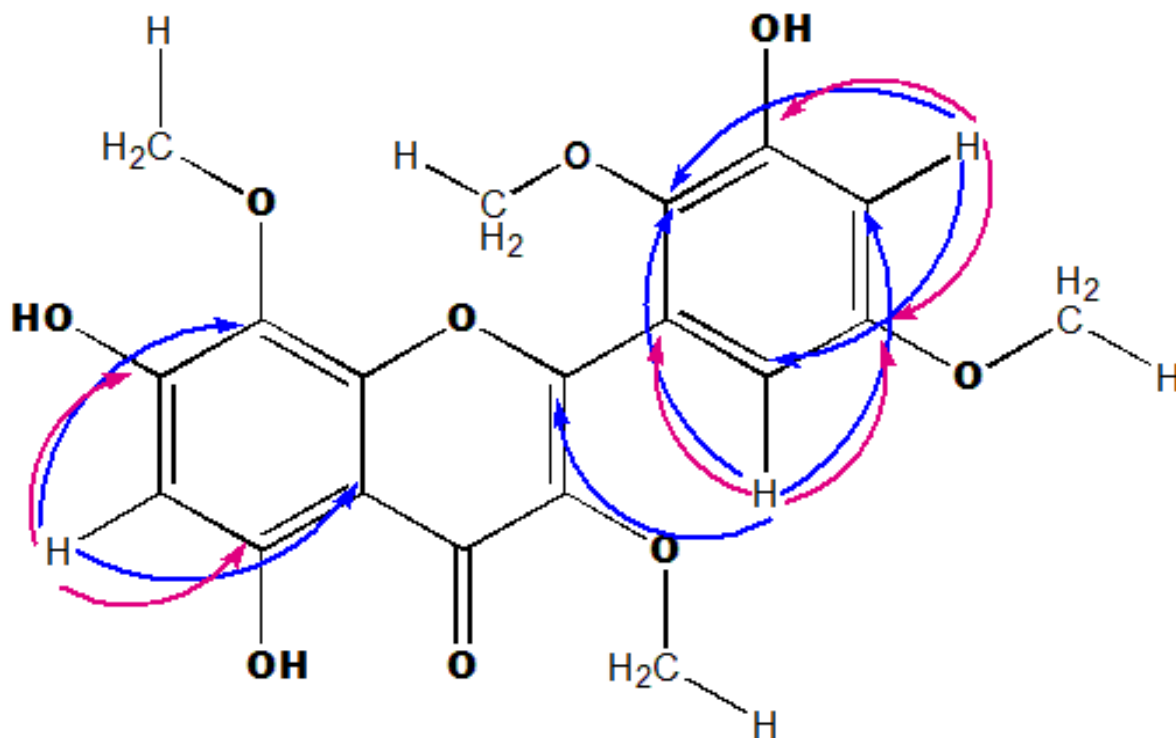


Figure 4.7: Structure of 5, 7, 3'-trihydroxy-3, 8, 2', 5'-tetramethoxyflavone (**84**)

Table 4.4: $^1\text{H}/^{13}\text{C}$ NMR (400/100 MHz) of Compound (**84**).

Position	δ_{H}	δ_{C}	HMBC
2	-	153.7	-
3	-	138.5	-
4	-	177.5	-
5	-	165	-
6	6.14, <i>s</i>	101.2	C-4,5,6,7,8,10
7	-	156.7	-
8	-	129.1	-
9	-	149.0	-
10	-	101.7	-
1'	-	126.0	-
2'	-	139.0	-
3'	-	151.5	-
4'	7.35, <i>d</i>	110.2	C-2', 3',4' 5', 6'
5'	-	153.4	-
6'	7.21, <i>d</i>	103.5	C-1', 2',3' 4', 5'
8-OCH ₃	3.79, <i>s</i>	60.25	C-8
2'-OCH ₃	3.81, <i>s</i>	60.7	C-2'
5'-OCH ₃	3.76, <i>s</i>	56.1	C-5'
3-OCH ₃	3.83, <i>s</i>	60.4	C-3

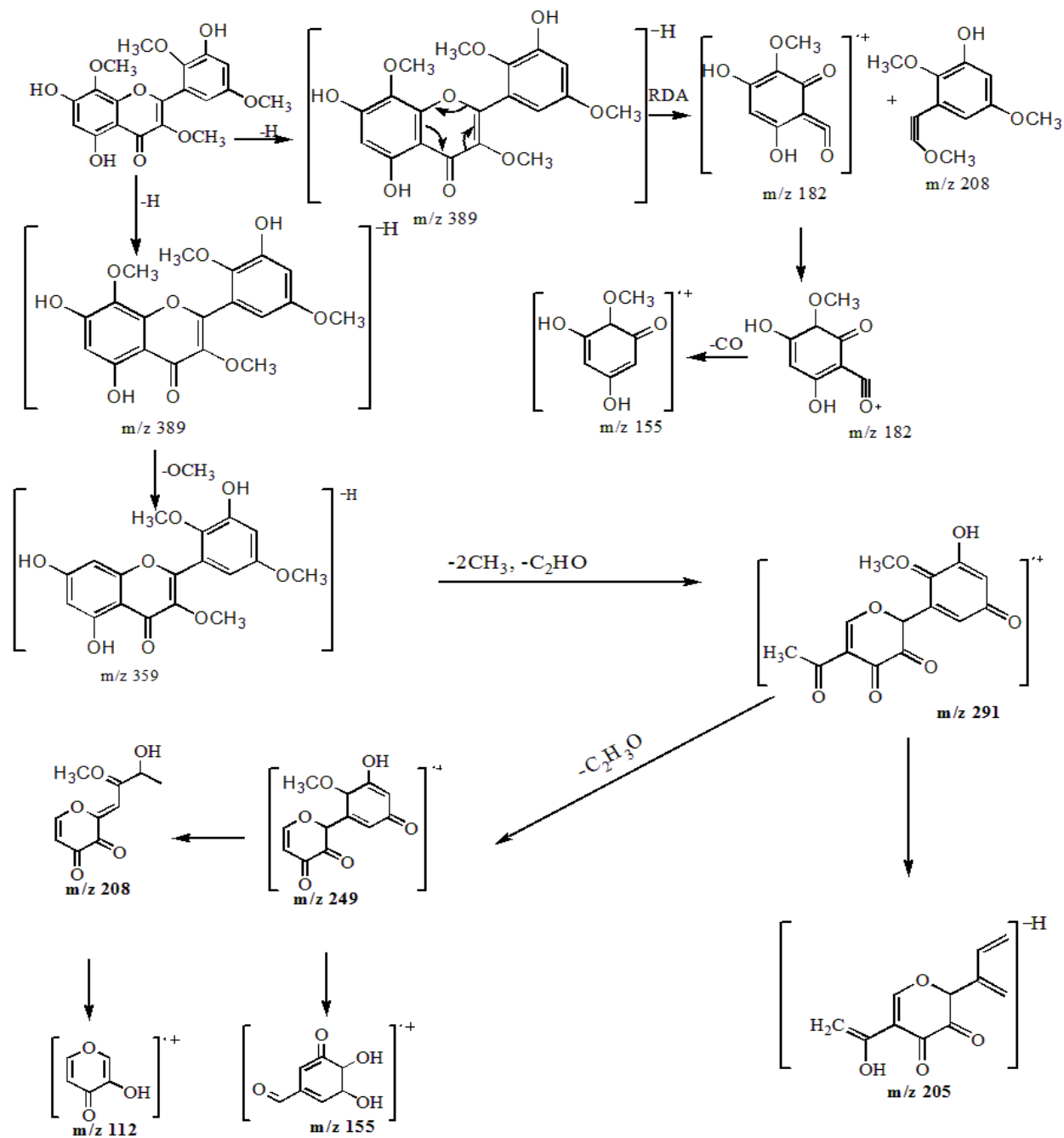


Figure 4.8: Fragmentation pattern for 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**)

4.1.1.5 5, 3', 4'-Trihydroxy-3, 7, 8-trimethoxyflavone (**85**)

Compound **85** was isolated as a yellow compound. Its HR-EIMS (Appendix 5E) molecular ion was observed at m/z 361 $[M+1]^+$ corresponded to the molecular formula of $C_{18}H_{16}O_8$ and the C-ring degraded by cleavage of bonds 0,2 to produce RDA fragment $^{0,2}A^+$ which subsequently lost a CO to produce a peak at m/z 169 (Figure 4.10).

Its ^1H NMR (Table 4.5 and Appendix 5A) spectrum revealed four aromatic signals δ_{H} 6.89 (*d*, $J=8.4$ Hz), δ_{H} (7.47, *dd*, $J=8.4$, 1.6 Hz) δ_{H} 7.58, (*d*, $J=1.6$ Hz) and a singlet δ_{H} 6.95. Its ^{13}C NMR (Table 4.5 and Appendix 5B) spectrum revealed a total of eighteen signals out of which fifteen belong to the flavonoid skeleton whereas the remaining three were attributed to methoxyl groups (Appendices 5A and 5B).

The downfield chemical shift of the hydroxyl groups at δ_{H} 12.62 suggests its *peri* position to the carbonyl carbon at C-4 and hence the hydroxyl group be assigned at C-5 position. The singlet aromatic proton at δ_{H} 6.95 showed HMBC correlation with C-4, C-5, C-7, C-8 and C-10 suggesting that this proton be assigned to H-6 (Appendices 5C and 5D). The above spectroscopic evidence suggests that positions C-5, C-7 and C-8 were oxygenated carbons belonging to δ_{C} 152.1, 159.0, 131.9 respectively. The HMBC correlation of the methoxy groups with C-8 (δ_{C} 131.9) and C-7 (δ_{C} 159.0) suggest that positions C-7 and C-8 are substituted with methoxy groups. Furthermore, the methoxyl group at δ_{H} 3.77/ δ_{C} 60.1 was placed at C-3 based on HMBC correlation with C-3 (δ_{C} 138.1).

The presence of two oxygenated quaternary carbons at δ_{C} 145.7 and 149.2 suggest that ring B of flavones skeleton is di-substituted. Taking into consideration the ABX spin pattern observed in ring B and HMBC correlation of these protons with the two oxygenated quaternary carbons suggest that ring B is oxygenated at C-3' and C-4' positions. The HRMS fragmentation pattern (Figure 4.10, Appendix 5E) showed a protonated molecular ion peak at m/z 361 and a fragment with m/z 169 (Wu *et al.*, 2003). Thus, based on the above spectroscopic evidence the compound was identified as 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**) as shown below. It was previously reported from *Ricinocarpos muricatus* (Henrick and Jefferies, 1965).

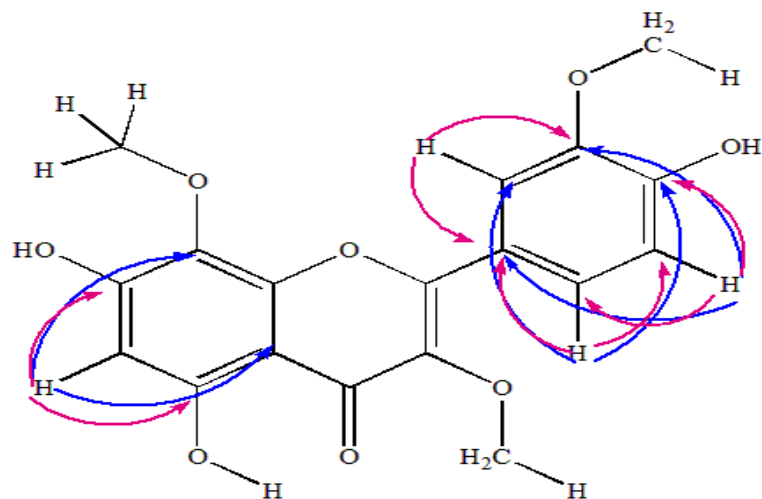


Figure 4.9: Structure of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**)

Table 4.5: $^1\text{H}/^{13}\text{C}$ NMR (400/100 MHz) of Compound (**85**).

Position	δ_{H}	δ_{C}	HMBC
2	-	156.4	-
3	-	138.1	-
4	-	178.6	-
5	-	152.1	-
6	6.95	91.7	C-4,5,6,7,8,10
7	-	159.0	-
8	-	131.9	-
9	-	149.2	-
10	-	105.9	-
1'	-	121.0	-
2'	7.58	115.9	C-1',2',3',4', 6'
3'	-	145.7	-
4'	-	149.2	-
5'	6.89	116.1	C-1',3', 4',5', 6'
6'	7.47	121.0	C-1', 2', 4', 5',6'
3-OCH ₃			C-3
7-OCH ₃			C-7
8-OCH ₃			C-8

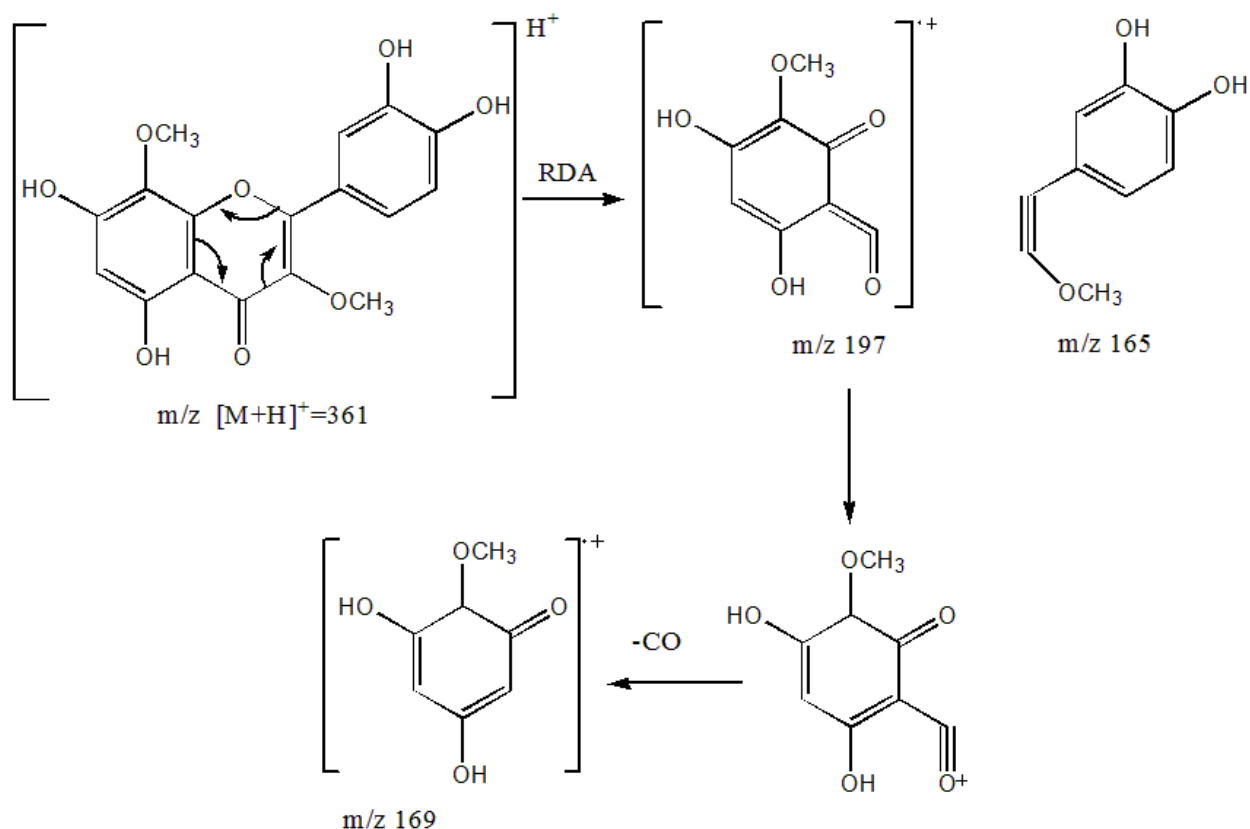


Figure 4.10: Fragmentation pattern for 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**)

4.1.2 Rearranged Clerodane diterpenoids

From the surface exudates of the aerial parts of *Microglossa pyrifolia* two rearranged clerodane diterpenoids, 8-acetoxyisochiliolide lactone (**86**) and epoxyisochiliolide lactone (**87**) were isolated of which **86** was found to be new compound.

4.1.2.1 8-Acetoxyisochiliolide lactone (**86**)

Compound **86** was isolated as white crystals. Its HR-EIMS (Appendix 6F) provided a molecular ion peak at 402.1700 which is attributed to $C_{22}H_{26}O_7$ from the (calc for 402.1679). Its ^{13}C NMR (Appendix 6B) revealed four olefinic carbons (δ_C 108.5, 126.0, 140.0, 144.5), three ester carbonyls (δ_C 169.6, 172.5, 177.5). The presence of two olefinic carbons, one ester carbonyl and five rings is in agreement with the DBE of 10 derived from its molecular formula.

The presence of two olefinic protons at δ_H 6.45 (*d*, 1.5 Hz) and 7.45 (*d*, 1.5 Hz) coupled with the singlet proton at 7.49 (*s*) and four olefinic carbons suggest the presence of a furan ring.

Two of the ester carbonyls (δ_C 172.5 and 177.5) were attributed to two lactone rings where as the one at δ_C 169.6 was an acetoxy carbonyl based on its HMBC (Appendix 6D) correlation with a methyl singlet at δ_H 1.79.

Three more methyl singlets were observed, two of which being vicinal methyls at δ_H 1.08 and 1.13 and the third methyl at δ_H 1.93. According to the HSQC, these methyl singlets were attached to carbons resonating at δ_C 9.4, 16.9 and 22.4 respectively (Appendices 6D and 6E). The presence of four olefinic carbons for a furanyl rings, two lactone carbonyls and three methyls suggests a clerodane diterpenoid of isochiliolide lactone skeleton (Zdero *et al.*, 1990b). Furthermore, these NMR data matched those reported for the pentacyclic isochiliolide lactone diterpenoid, 8-hydroxyisochiliolide lactone reported from *M. pyrrophappa* (Zdero *et al.*, 1990b). Thus, based on the above spectroscopic evidence the compound was identified to be 8-acetoxyisochiliolide lactone (**86**) as in figure 4.12.

4.1.2.2 7, 8-Epoxyisocholiolide lactone (**87**)

Compound **87** was isolated as white crystals from fraction 12I. The ^{13}C -NMR (appendix 7B) had twenty signals implying that it was a diterpenoid with twenty carbons. Among the carbons were four olefinic carbons at δ_C 144.8, 140.8, 124.5, and 108.5. [for a furanyl ring, δ_H 7.58 (*d*, $J = 1.2$ Hz), δ_H 7.51 (*br s*) and δ_H 6.51 (*br s*)], two lactone carbonyls at δ_C 172.2, 178.1 and three methyls at δ_C 9.7, 17.4 and 20.8, suggesting a clerodane diterpenoid of isochiliolide lactone skeleton (Zdero *et al.*, 1990b). The adjacent oxygen bearing carbons at positions 7 and 8 gave peaks at δ_C 62.6 and δ_C 60.8 and were deduced to be forming an epoxide from the EIMS molecular ion at m/z 360 (Appendix 7E). Compound **87** lacks one of the acetoxy moiety in the NMR spectrum and instead epoxy groups at positions 7 and 8 appeared at δ_C 62.6 and 60.8. The above spectroscopic data for compound **87** is the same as the data reported for 7, 8-epoxyisocholiolide lactone isolated from *M. pyrrophappa* (Zdero *et al.*, 1990b). So it is concluded that the compound is 7, 8-epoxyisocholiolide lactone (**87**) as shown in figure 4.11.

Table 4.6: $^1\text{H}/^{13}\text{C}$ NMR (400/100 MHz) of Compounds (**86**) and (**87**).

	86			87	
	^{13}C	^1H	HMBC	^{13}C	^1H
1	28.8	2.23, <i>ddd</i> (13.6, 10.5, 3.8) 1.84, <i>ddd</i> (13.6, 9.2, 4.2)	C-2 C-2, C-5	28.8	1.90, <i>m</i> 1.72, <i>m</i>
2	29.4	1.94, <i>m</i> 1.71, <i>ddd</i> (13.0, 9.2, 3.8)	C-1 C-1	28.5	1.90, <i>m</i> 2.26, <i>m</i>
3	177.5			178.4	
4	54.4			55.6	
5	51.2			50.0	
6	26.1	1.44, <i>td</i> (14.0, 3.2) 1.39, <i>dt</i> (14.0, 3.8)	C-5, C-7	28.4	1.93, <i>br d</i> (18.0) 1.87, <i>m</i>
7	31.5	2.54, <i>dt</i> (14.0, 3.6) 1.75, <i>m</i>	C-5, C-6, C-17	62.6	3.35, <i>br s</i>
8	86.0			60.8	
9	54.4			52.0	
10	93.3			91.0	
11	34.4	2.69, <i>dd</i> (13.8, 8.2) 2.61, <i>dd</i> (13.8, 8.8)	C-8, C-9, C-10, C-12, C-13 C-9, C-10, C-12, C-13, C-20	37.6	2.63, <i>dd</i> (14.4, 7.8) 2.67, <i>dd</i> (14.4, 9.4)
12	70.7	5.37, <i>dd</i> (8.8, 8.2)		72.1	5.40, <i>dd</i> (9.4, 7.8)
13	126.0			124.5	
14	108.5	6.45, <i>d</i> (1.5)	C-13, C-15, C-16	108.5	6.51, <i>br s</i>
15	144.5	7.47, <i>t</i> (1.5)	C-13, C-16	144.8	7.51, <i>d</i> (1.2)
16	140.0	7.49, <i>s</i>	C-14, C-15	140.8	7.58, <i>br s</i>
17	21.2	1.93, <i>s</i>	C-7, C-8, C-9	20.8	1.52, <i>s</i>
18	9.4	1.08, <i>s</i>	C-2, C-3, C-4, C-5	9.7	1.12, <i>s</i>
19	16.9	1.13, <i>s</i>	C-4, C-5, C-6, C-10	17.4	1.13, <i>s</i>
20	172.5			172.2	
OAc- CO	169.6				
OAc- Me	22.2	1.79, <i>s</i>	C-OAc CO		

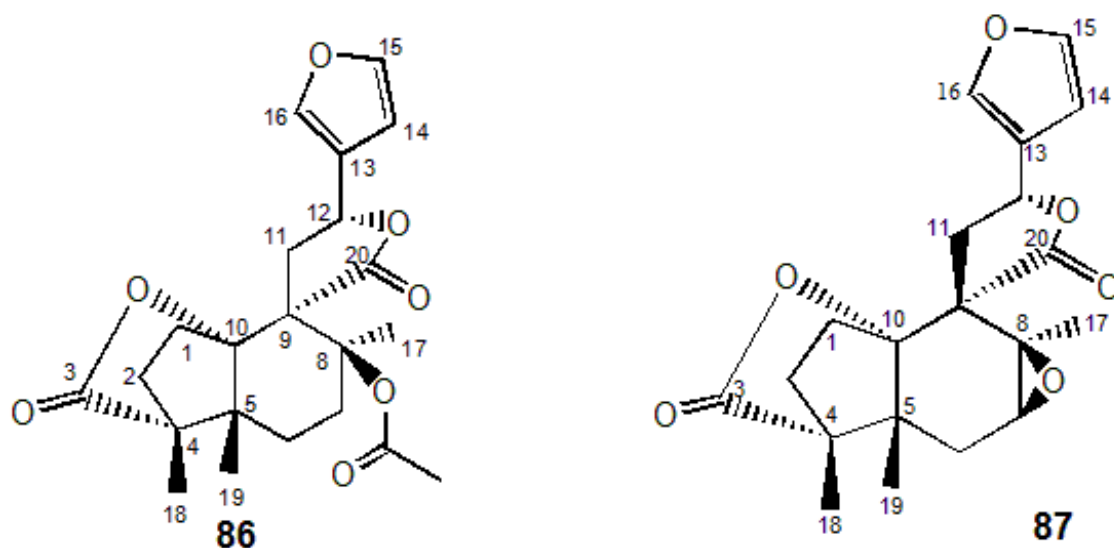


Figure 4.11: Structures of rearranged clerodane diterpenoids, 8-acetoxyisocholiolide lactone (**86**) and 7, 8-epoxyisocholiolide lactone (**87**)

4.2 Anti-plasmodial, anti-oxidant, anti-microbial and anti-leishmanial activities

All the compounds isolated were subjected to bio assay tests to establish their potential as antiplasmodial, antioxidant, antimicrobial and antileishmanial as follows.

4.2.1 Anti-plasmodial activity of compounds

As a follow up to the antiplasmodial activities shown by the extract of the investigated plants (Table 1.1), *in-vitro* antiplasmodial tests were done for the compounds isolated from the plant. The tests aimed at identifying active components which contributed to the observed activities of crude extract. The compounds tested were 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**), 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**), 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**), 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**), 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**), 8-acetoxyisocholiolide lactone (**86**) and 7, 8-epoxyisocholiolide lactone (**87**). Two *P. falciparum* strains were used, Sierra Leone D6 (chloroquine-sensitive) and Indochina W2 (chloroquine-resistant). Cytotoxicities of the compound were measured simultaneously. These results are summarized in table 4.7.

Generally, a compound is considered to be inactive when it shows an $IC_{50} > 200 \mu M$, whereas those with an IC_{50} of 100-200 μM have low activity; IC_{50} of 20-100 μM , moderate activity; IC_{50} of 1-20 μM good activity; and $IC_{50} < 1 \mu M$ excellent/potent antiplasmodial activity (Batista *et al.*, 200). All the compounds under test except crude extract did not show antiplasmodial activities. This can be interpreted that the activity of their extracts are either due to synergistic effects or contributed by trace compounds which were not isolated.

Table 4.7: In-vitro antiplasmodial activity and cytotoxicity

Compound	<i>In-vitro</i> antiplasmodial activity (IC_{50})		Cytotoxicity (IC_{50})
	D6 ($\mu g/ml$)	W2 ($\mu g/ml$)	($\mu g/ml$)
5, 7, 4'-Trihydroxy-3, 8, 3'-trimethoxyflavone (81)	NA	NA	NC
5, 7, 4'-Trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (82)	NA	NA	NC
5, 3', 4'-Trihydroxy-7-methoxyflavanone (83)	NA	NA	NC
5,7,3'-Trihydroxy-3,8,4',5'-tetramethoxyflavone (84)	NA	NA	NC
5, 3', 4'-Trihydroxy-3, 7, 8-trimethoxyflavone (85)	NA	NA	NC
8-Acetoxyisochiliolide lactone (86)	NA	NA	NC
7, 8-Epoxyisocholiolide lactone (87)	NA	NA	NC
Crude extract	8.00	13.00	NC
Chloroquine (standard)	<0.026	0.14	NC

NA = Not Active (IC_{50} values $> 50 \mu g/ml$); NC = Not Cytotoxic (up to the maximum dose tested 5.0 $\mu g/ml$).

4.2.2 Anti-oxidant activities of compounds

All the compounds isolated were tested for anti-oxidant activity. The compounds were first subjected to preliminary qualitative testing on a TLC, out of which Compounds **86** and **87** (rearranged clerodane diterpenoids) were inactive while compounds **81**, **82**, **83**, **84** and **85** (flavonoids) were active. Compounds which showed activities on TLC assays were quantitatively analysed by UV-VIS absorption measurements on the compound solutions on reaction with DPPH. The results are presented in Table 4.8 and Figure 4.8.

Table 4.8: Anti-oxidant activity of flavonoids with DPPH experiment

Compound	IC ₅₀	
	µg/ml	µM
5, 7, 4'-Trihydroxy-3, 8, 3'-trimethoxyflavone (81)	6.02±0.2	17.3
5, 7, 4'-Trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (82)	6.45±0.3	16.5
5, 3', 4'-Trihydroxy-7-methoxyflavanone (83)	8.54±0.3	28.3
5, 7, 3'-Trihydroxy-3,8, 4', 5'-tetramethoxyflavone (84)	8.79±0.3	22.5
5, 3', 4'-Trihydroxy-3, 7, 8-trimethoxyflavone (85)	6.2±0.2	17.2
Quercetin (standard)	6.02 ± 0.2	19.9

The most active compound was 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**). Its radical scavenging activity of 6.2µg/ml (17.3 µM) was less than that of the standard used, quercetin (IC₅₀ 6.0 µg/ml; 19.9 µM).

All of the compounds found to have radical scavenging activities had phenolic moiety oxygenated at either *ortho* or *para* positions, explained by the ability of hydroxyl groups at these positions to reduce radicals and form quinones (Chen *et al.*, 1996), as illustrated for compound **81** in figure 4.13.

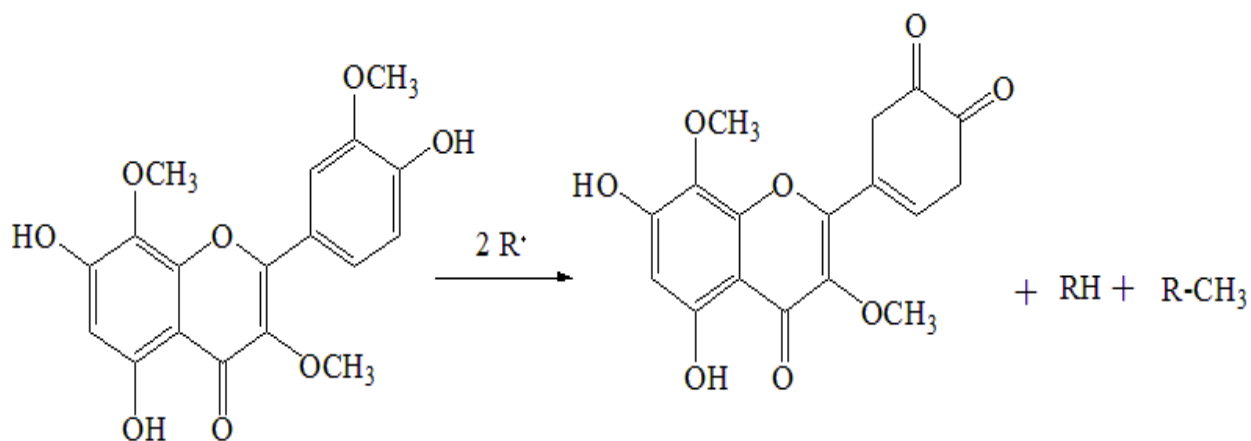


Figure 4.12: Reduction of radicals by a compound with phenolic ortho oxygenation.

4.2.3 Antimicrobial activities

The isolated compounds were tested for antifungal and antibacterial activities, using fungi *Candida albicans*, *Candida glabrata*, *Candida krusei*, *Cryptococcus neoformans*, *Aspergillus fumigatus* as well as the bacteria *Staphylococcus aureus*, methicillin-resistant *Staphylococcus aureus* (MRS), *Escherichia coli*, *Pseudomonas aeruginosa* and *Mycobacterium intracellulare*. Compounds showing % Inhibition < 50 are considered inactive at 15.9 $\mu\text{g}/\text{m}$. All the compounds tested showed no activity against any fungi or bacteria.

4.2.4 Anti-leishmanial activities

Antileishmanial activities of the compounds were tested *in vitro* on a culture of *Leishmania donovani* promastigotes. 5, 3', 4'-Trihydroxy-7-methoxyflavanone (**83**) exhibited mild antileishmanial activity (IC_{50} 13.13 $\mu\text{g}/\text{ml}$). Compounds which exhibited antileishmanial activities of IC_{50} >50 $\mu\text{g}/\text{ml}$ were considered inactive. Results for antileishmanial activity are summarized in Table 4.11.

Table 4.9: Antileishmanial activities

Compound	Activity against <i>L. donovani</i> ($\mu\text{g/ml}$)	
	IC ₅₀	IC ₉₀
5, 7, 4'-Trihydroxy-3, 8, 3'-trimethoxyflavone (81)	NA	NA
5, 7, 4'-Trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (82)	NA	NA
5, 3', 4'-Trihydroxy-7-methoxyflavanone (83)	13.13	NA
5, 7, 3'-Trihydroxy-3,8, 4', 5'-tetramethoxyflavone (84)	NA	NA
5, 3', 4'-Trihydroxy-3, 7, 8-trimethoxyflavone (85)	31.13	38.15
8-Acetoxyisochiliolide lactone (86)	NA	NA
7, 8-Epoxyisocholiolide lactone (87)	38	NA
Pentamidine (standard)	0.85	1.75
Amphotericin B (standard)	0.12	0.15

NA = Not Active

CHAPTER FIVE

CONCLUSIONS AND RECOMMENDATIONS

5.1 Conclusions

Chromatographic separation of the surface exudates of *Microglossa pyrifolia* led to the isolation and characterization of five flavonoids and two diterpenoids, of which three of them, 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**), 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**) and 8-epoxyisocholiolide lactone (**86**), are new compounds.

In-vitro anti-plasmodial testing revealed that all compounds were found to be inactive against D2 and W6 strains of *Plasmodium falciparum*. The internal tissues of the aerial parts of *M pyrifolia* studied previously by Zdero *et al* afforded clerodanes, sec-clerodane, rearranged clerodane and geranylgeranol derivatives. During the preliminary investigations, the crude extract was found to have IC₅₀ of 8.0 µg/ml (against *P.falciparum* D6) and 13.0 µg/ml (against *P.falciparum* W2). In this analysis the compounds isolated were from surface exudates. None of these compounds resembled those isolated by Zdero *et al* and the bulk of them were flavonoids. This therefore suggests that either the preliminary result was due to the compounds found in the internal tissues which are different from those on the surface or that it was due to synergistic effect or to trace compounds which were not isolated.

Anti-oxidant testing revealed that phenolic compounds with the phenolic hydroxyl groups and other hydroxyl or methoxyl groups at ortho or para positions were reasonably active. These were the flavonoids and the most active of them was 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) with IC₅₀ of 6.02 µg/ml (16.7 µM) which was less than that of quercetin (IC₅₀ 6.0 µg/ml; 19.9 µM), the standard used. Other compounds that had good anti-oxidant activities were, **82** (5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone) and compound **85** (5, 3', 4'-Trihydroxy-3, 7, 8-trimethoxyflavone) which had IC₅₀ of 6.45 ± 0.3 µg/ml (16.7 µM) and 6.2 ± 0.2 µg/ml (17.2 µM) respectively. This gives a revelation for a possibility of having an anti-malarial which is also an anti-oxidant, and hence giving the possibility of fighting malaria while minimizing the chances of developing complications from oxidative stress.

All the compounds were inactive to the *Leishmania donovani promastigotes*, except for 5, 3', 4'-Trihydroxy-7-methoxyflavanone (**83**) which exhibited mild antileishmanial activity (IC₅₀ 13.13µg/ml).

5.2 Recommendations

- i. In taxonomic terms, other related species within the genus have to be phytochemically investigated so as to establish detailed botanical grouping of the genus.
- ii. Careful analysis of the antiplasmodial activity of all compounds of the genus and their structure-activity relationship followed by rational synthetic modifications has potential for identifying useful agents with low cytotoxicity and high anti-plasmodial activity in the fight against malaria.
- iii. The anti-plasmodial activity of the mixture of compounds should be performed to establish which combination would give a better activity than the individual compounds
- iv. This mixture of non-polar compounds (flavonoids and diterpenoids) could be playing an ecological role for *M. pyrifolia*. The methoxylation of flavonoid hydroxyl groups is an established anti-herbivore, anti-microbiol feature in other species. They also play a role in sunscreen. The existence of the modified clerodanes, **86** and **87** are an interesting feature of this exudate and the potential activities of these compounds should be pursued to help speculate their potential biological function

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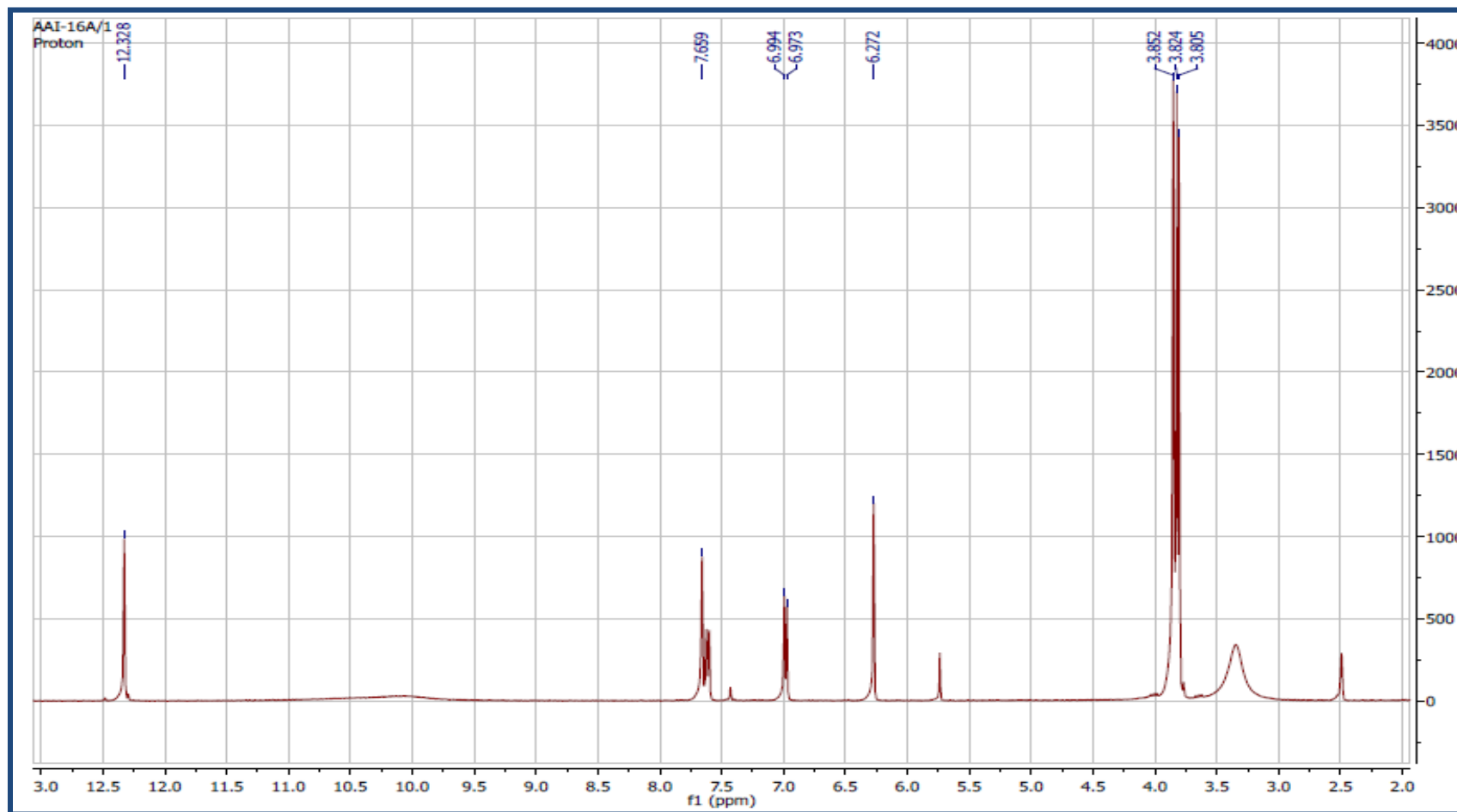
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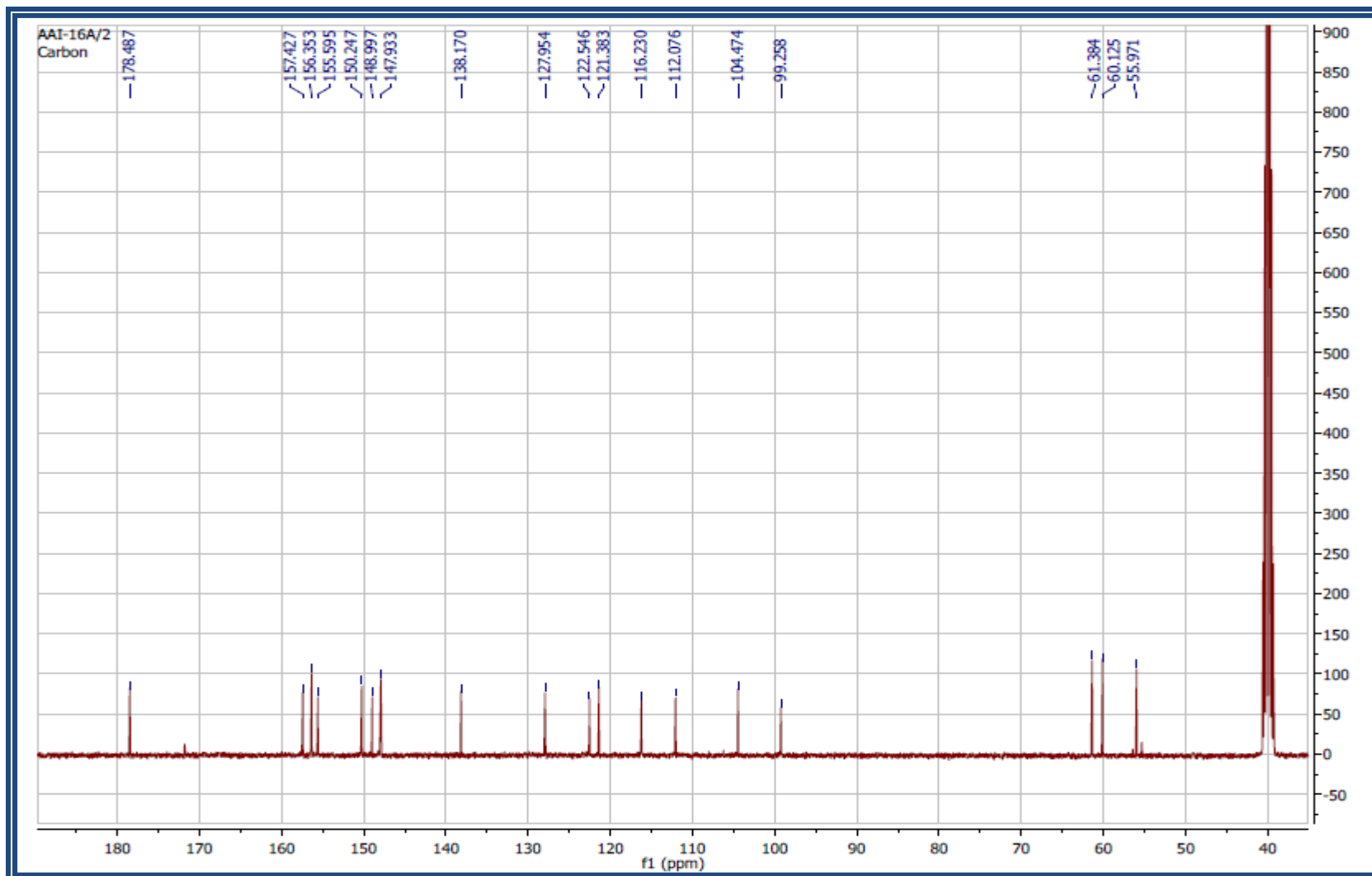
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APPENDICES

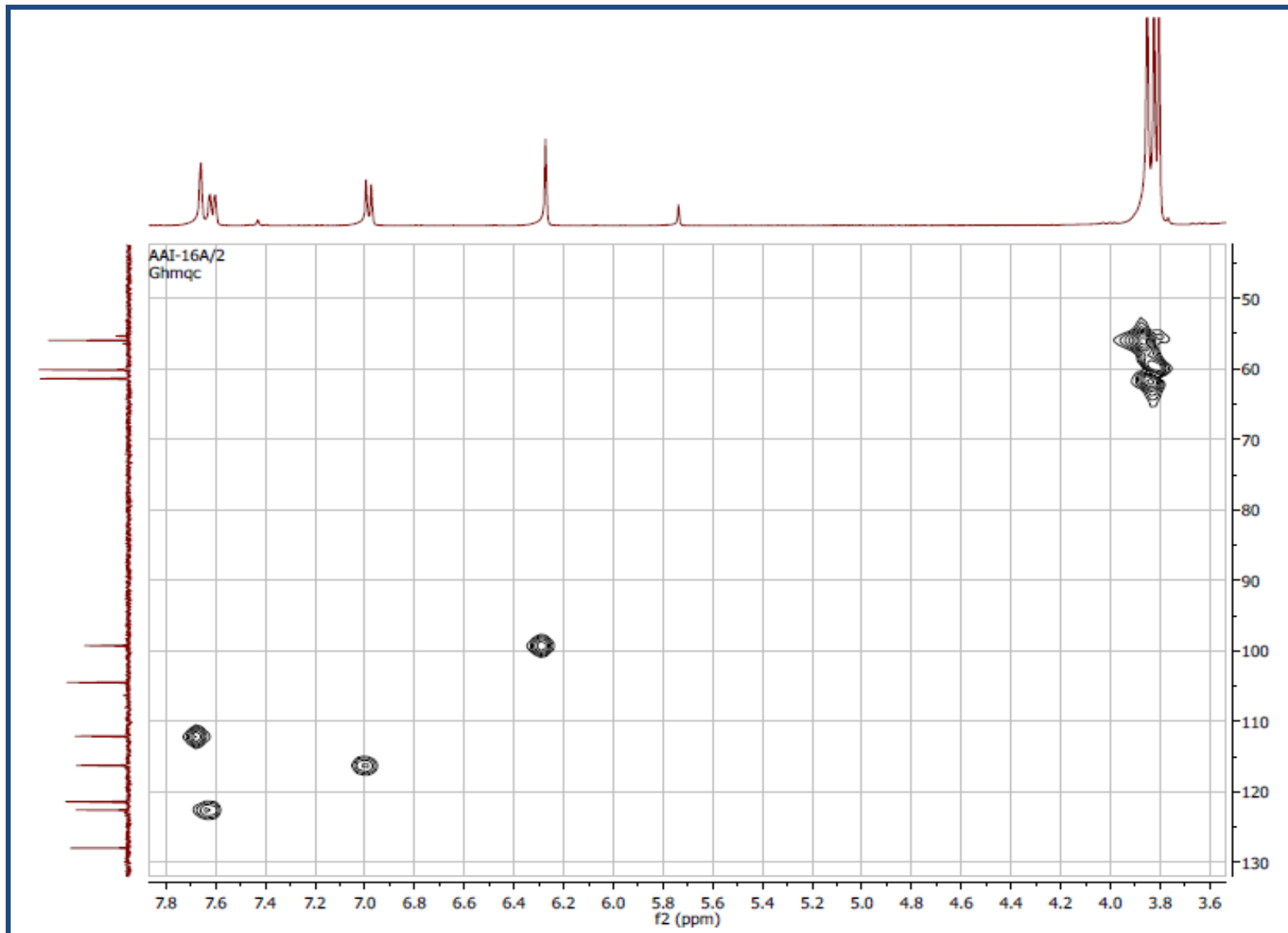
Appendix 1A: ^1H NMR (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) in DMSO-d_6



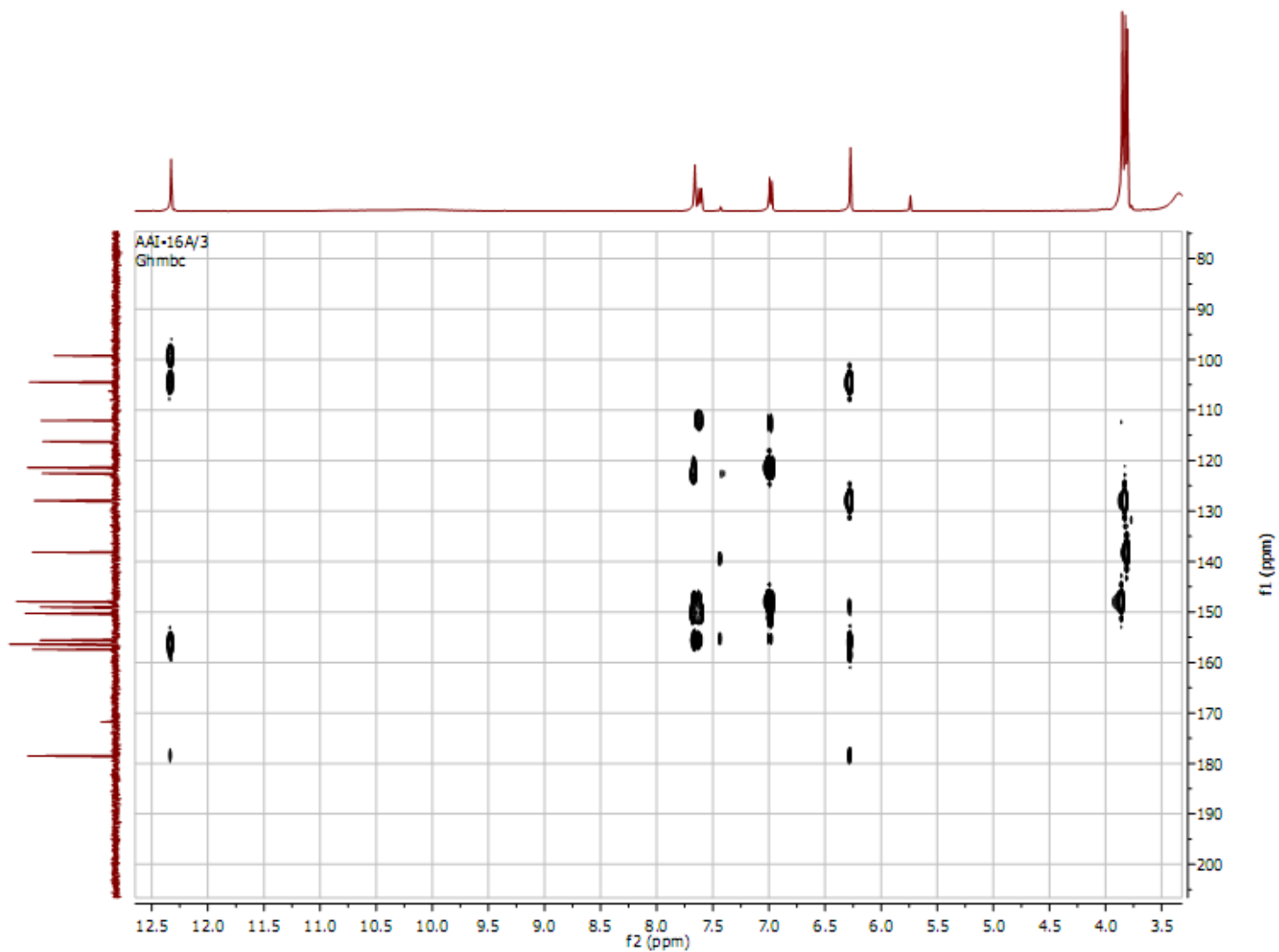
Appendix 1B: ^{13}C NMR (100 MHz) spectrum 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) in DMSO-d_6



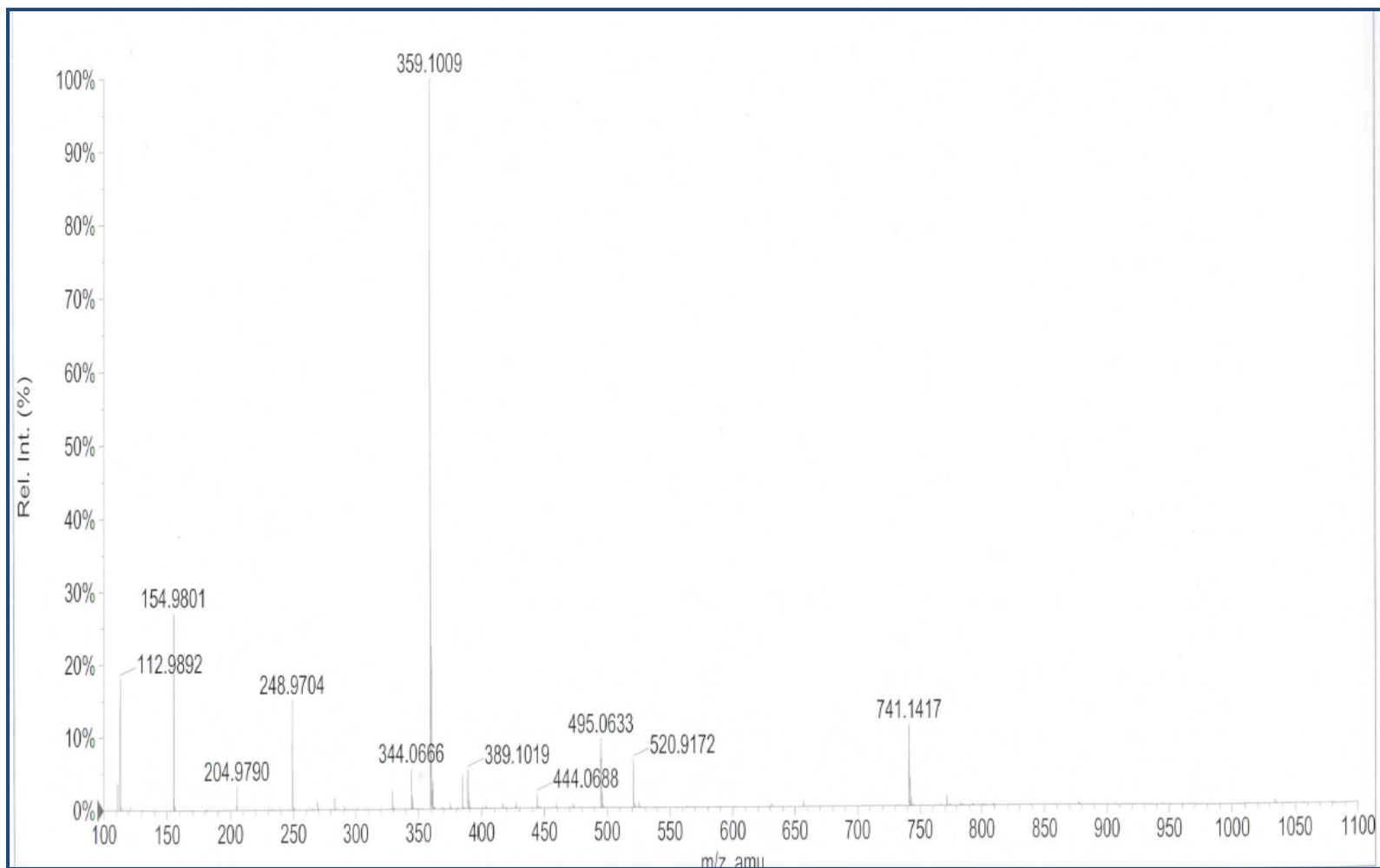
Appendix 1C: HSQC (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) in DMSO-d₆



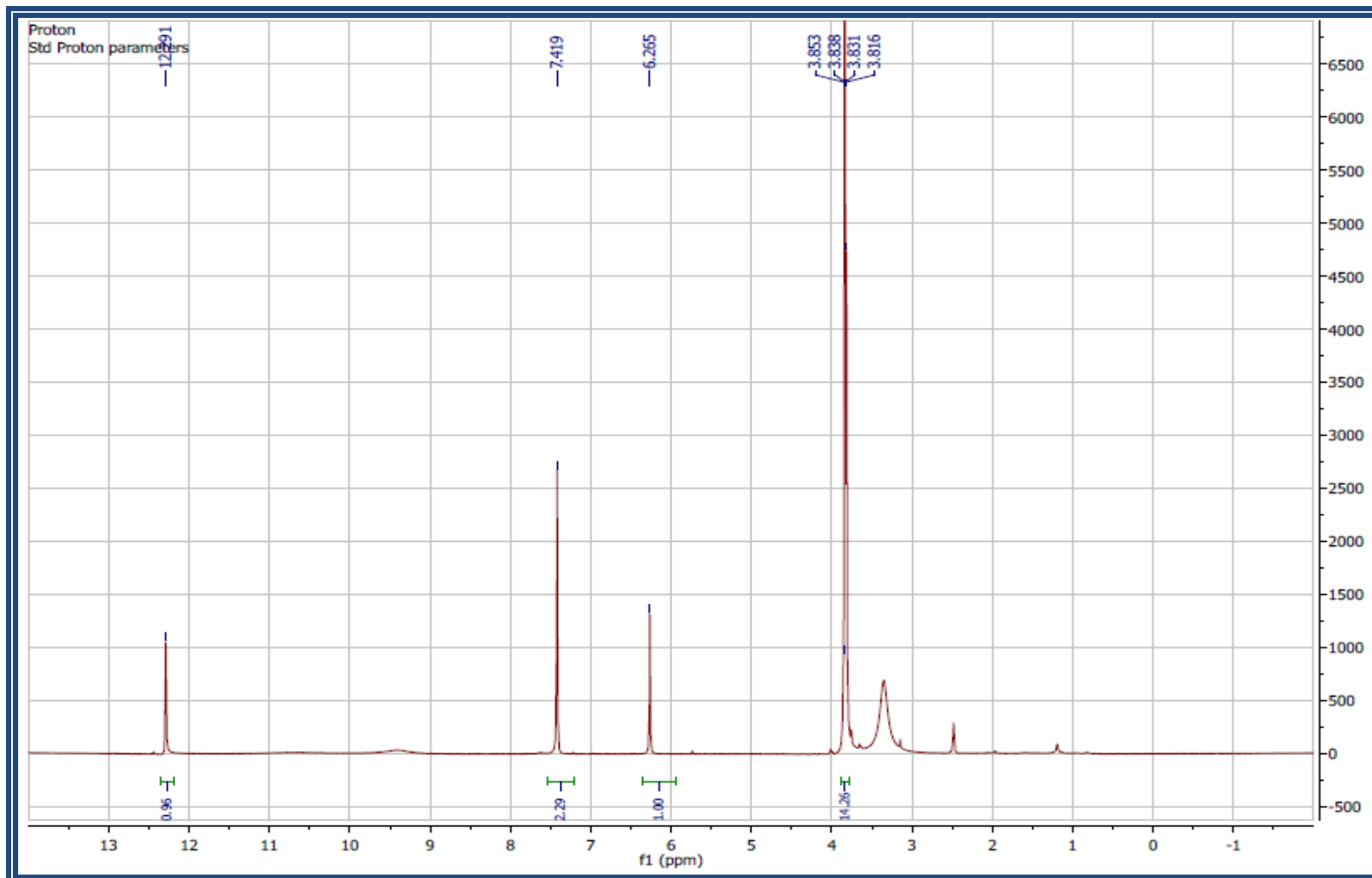
Appendix 1D: HMBC (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**) in DMSO-d₆



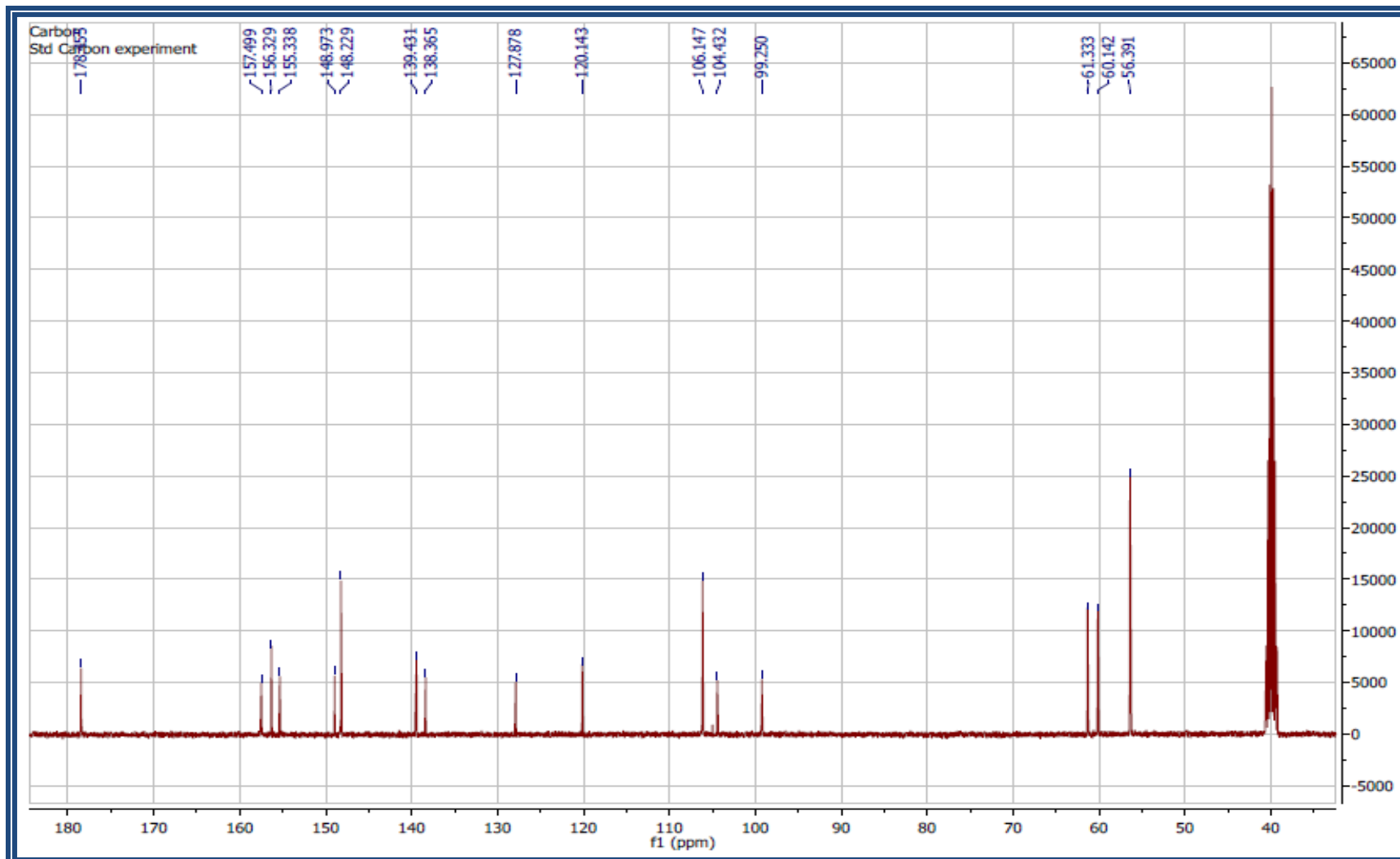
Appendix 1E: HR-EIMS spectrum of 5, 7, 4'-trihydroxy-3, 8, 3'-trimethoxyflavone (**81**)



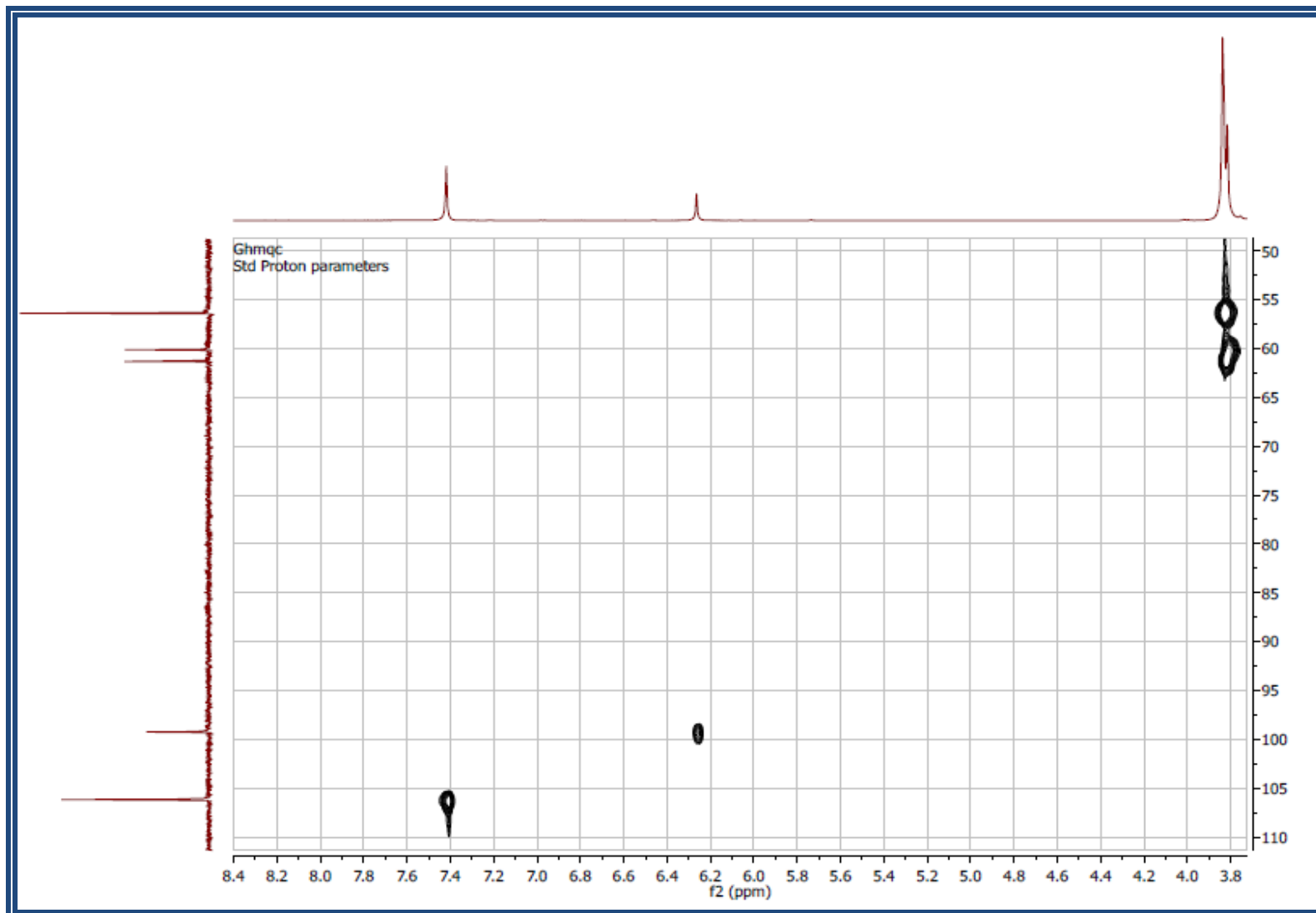
Appendix 2A: ^1H NMR (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**) in DMSO-d₆



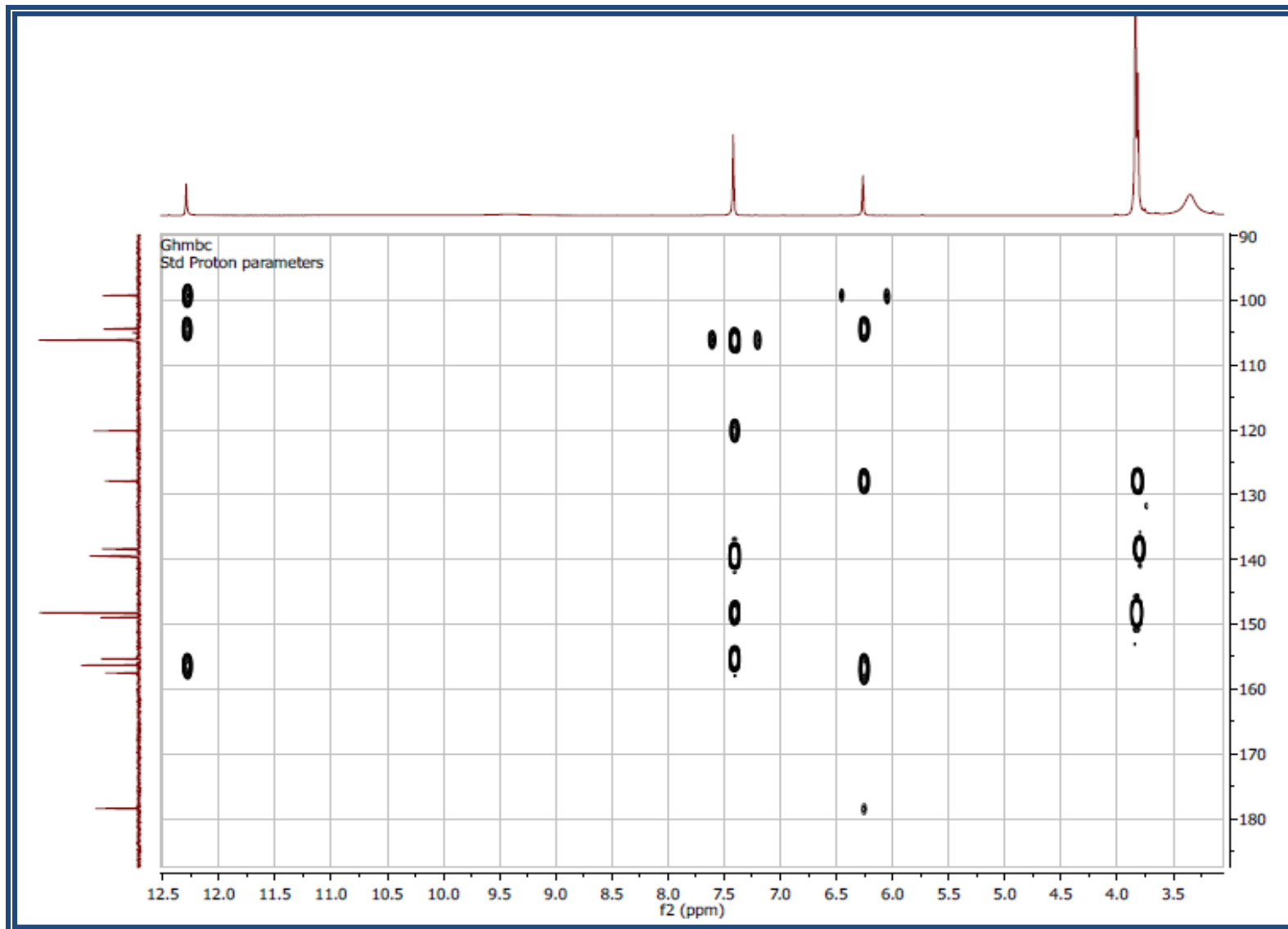
Appendix 2B: ^{13}C NMR (100 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**) in DMSO-d_6



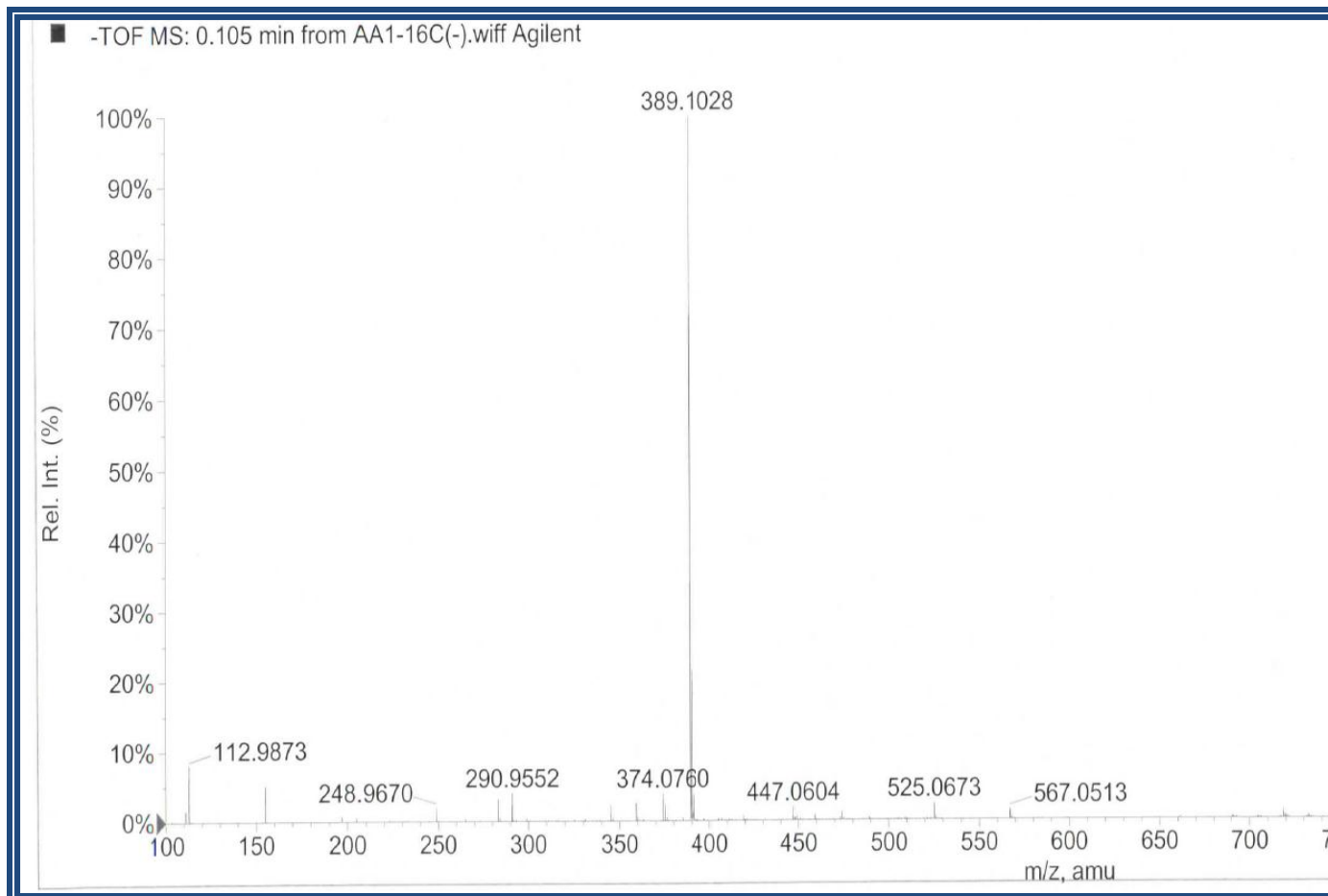
Appendix 2C: HSQC (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**) in DMSO-d₆



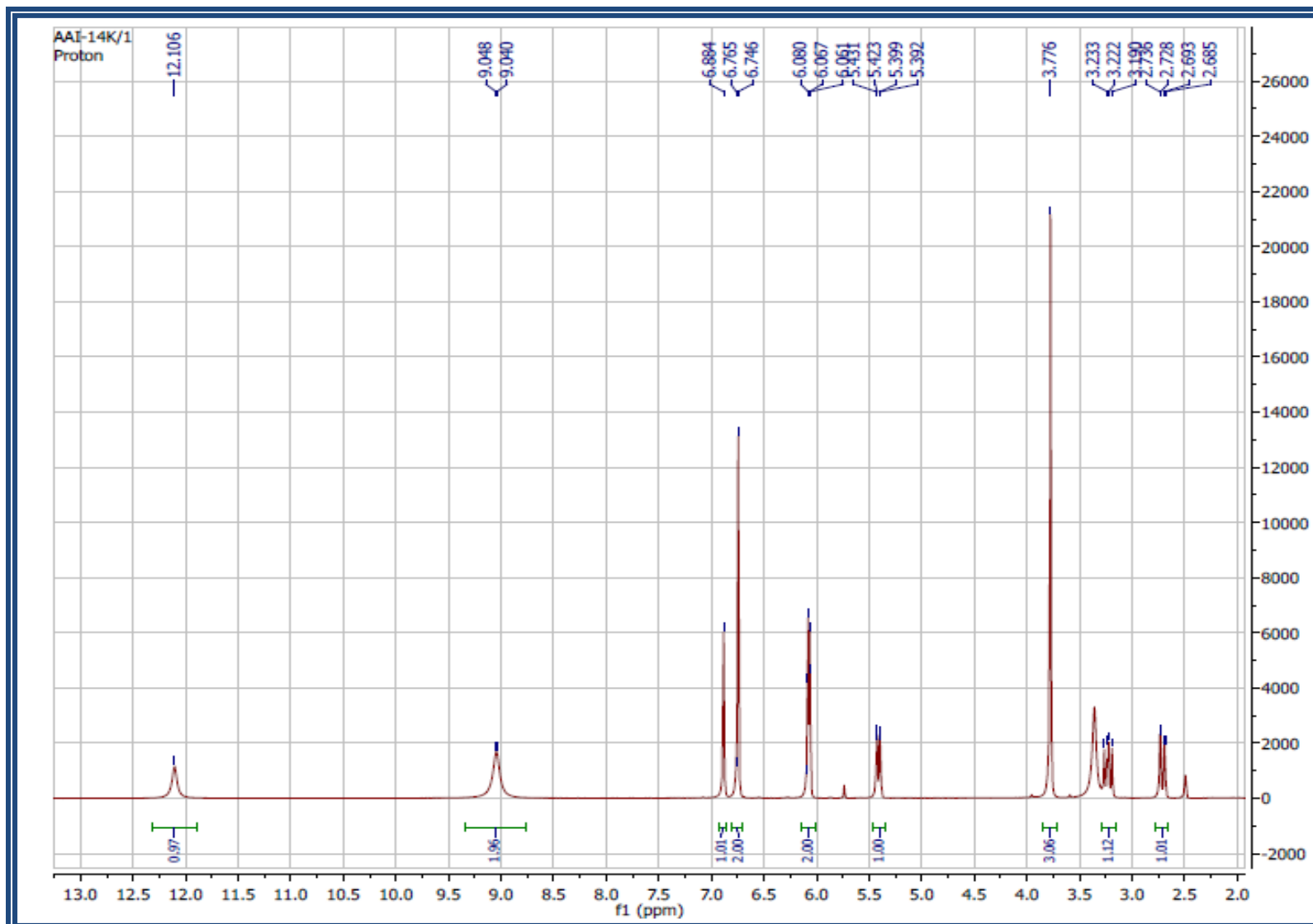
Appendix 2D: HMBC (400 MHz) spectrum of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**) in DMSO-d₆



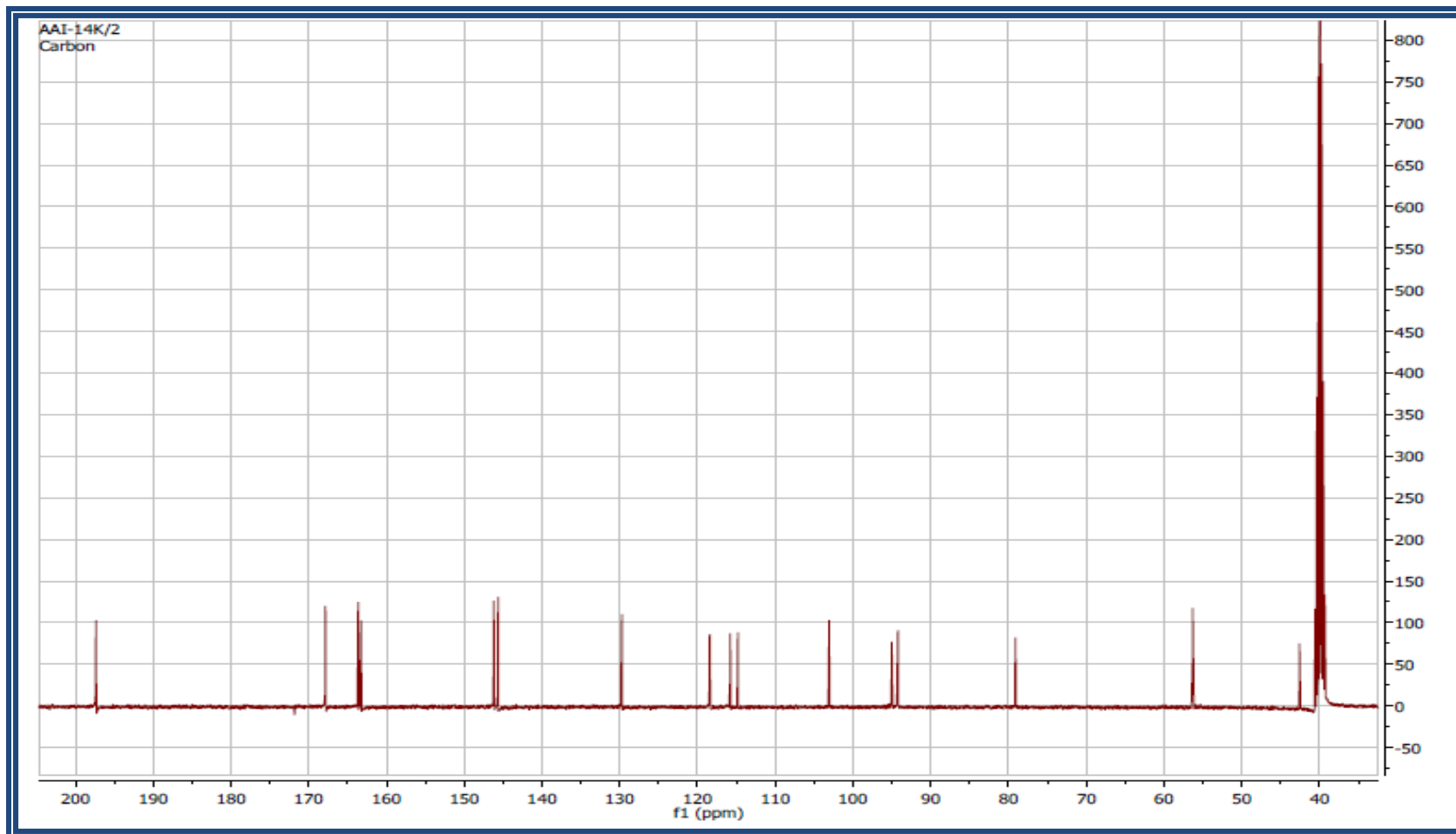
Appendix 2E: HR-EIMS spectrum of 5, 7, 4'-trihydroxy-3, 8, 3', 5'-tetramethoxyflavone (**82**)



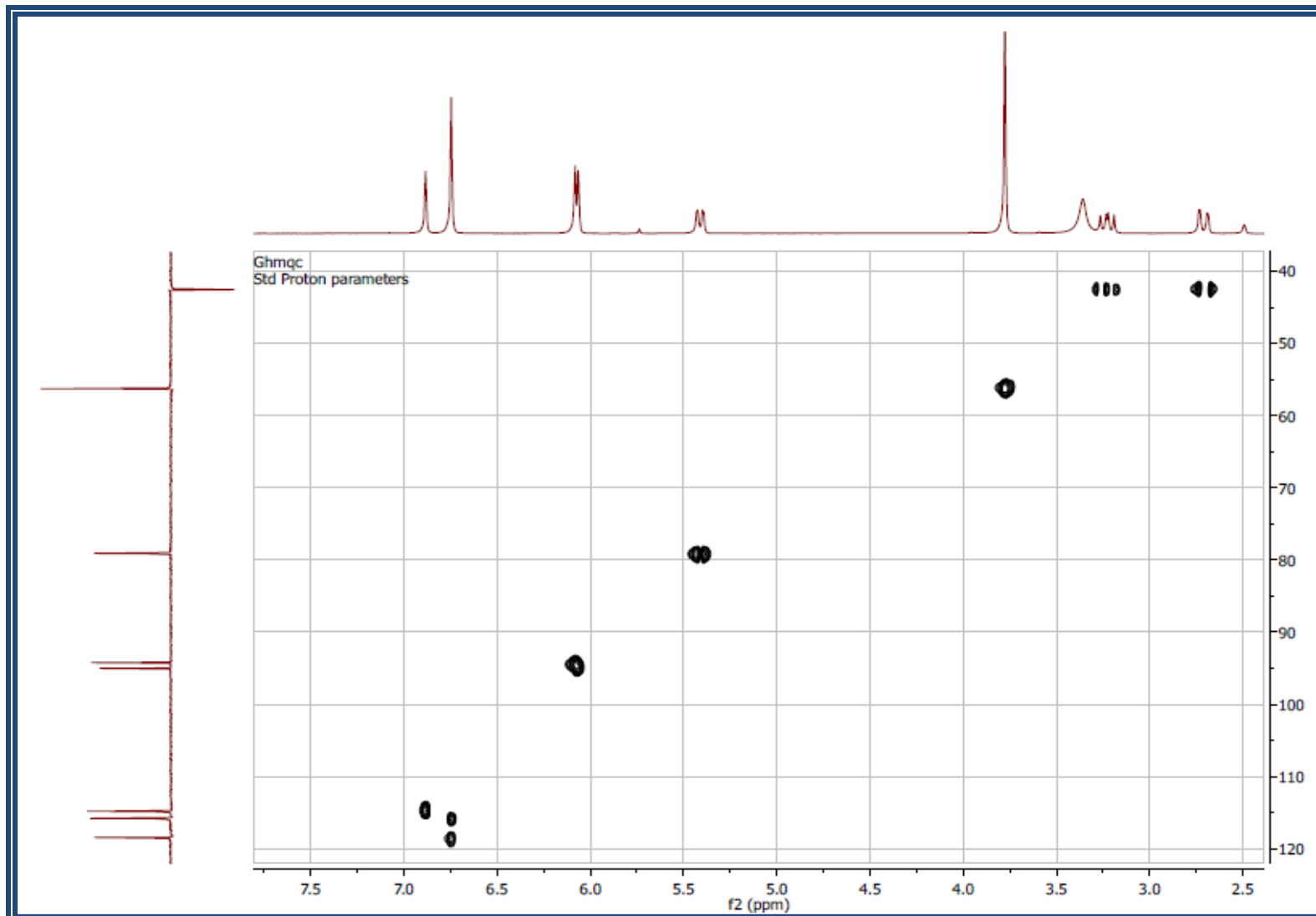
Appendix 3A: ^1H NMR (400 MHz) spectrum of 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**) in DMSO-d_6



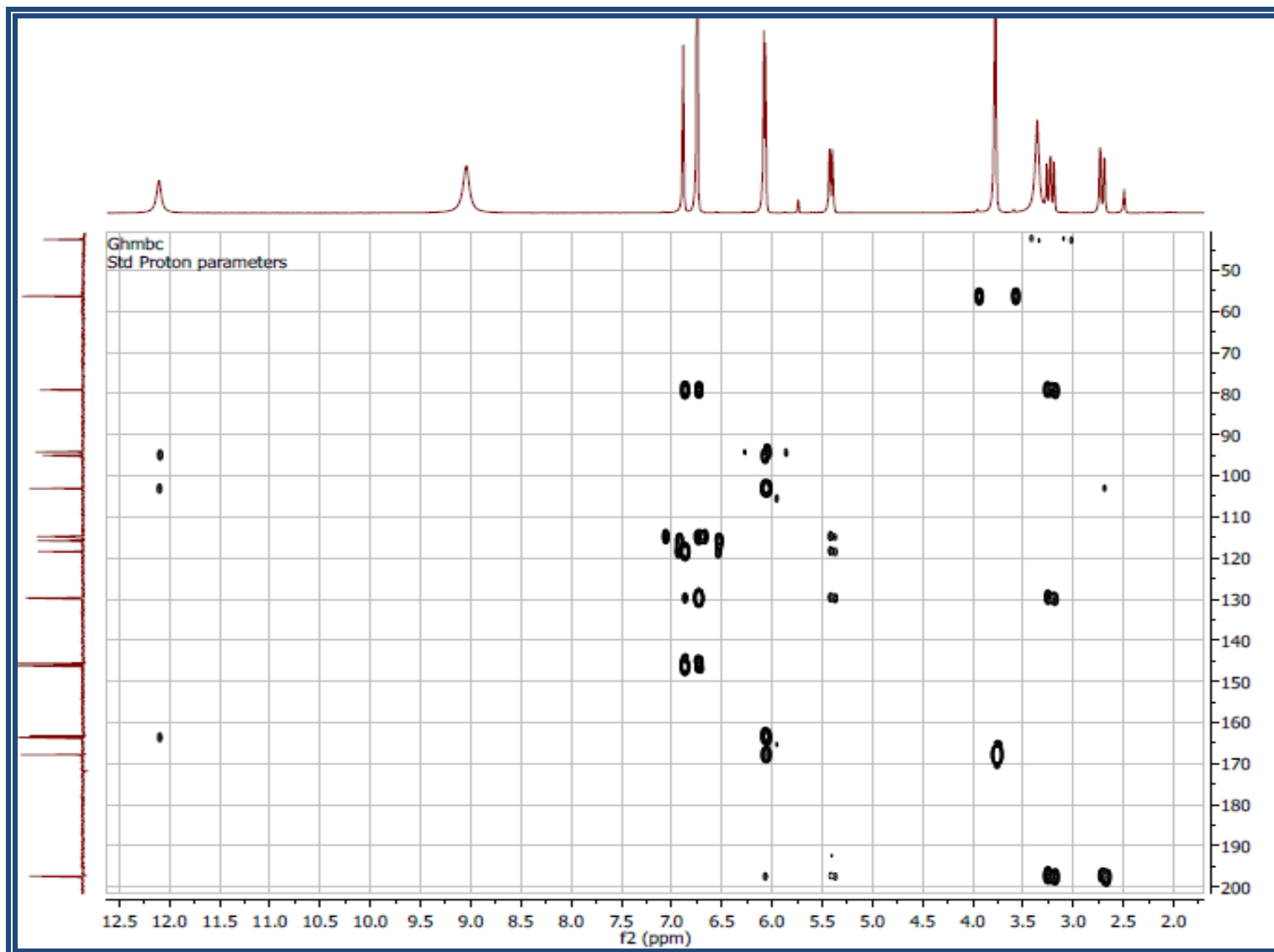
Appendix 3B: ^{13}C NMR (100 MHz) spectrum of 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**) in DMSO-d_6



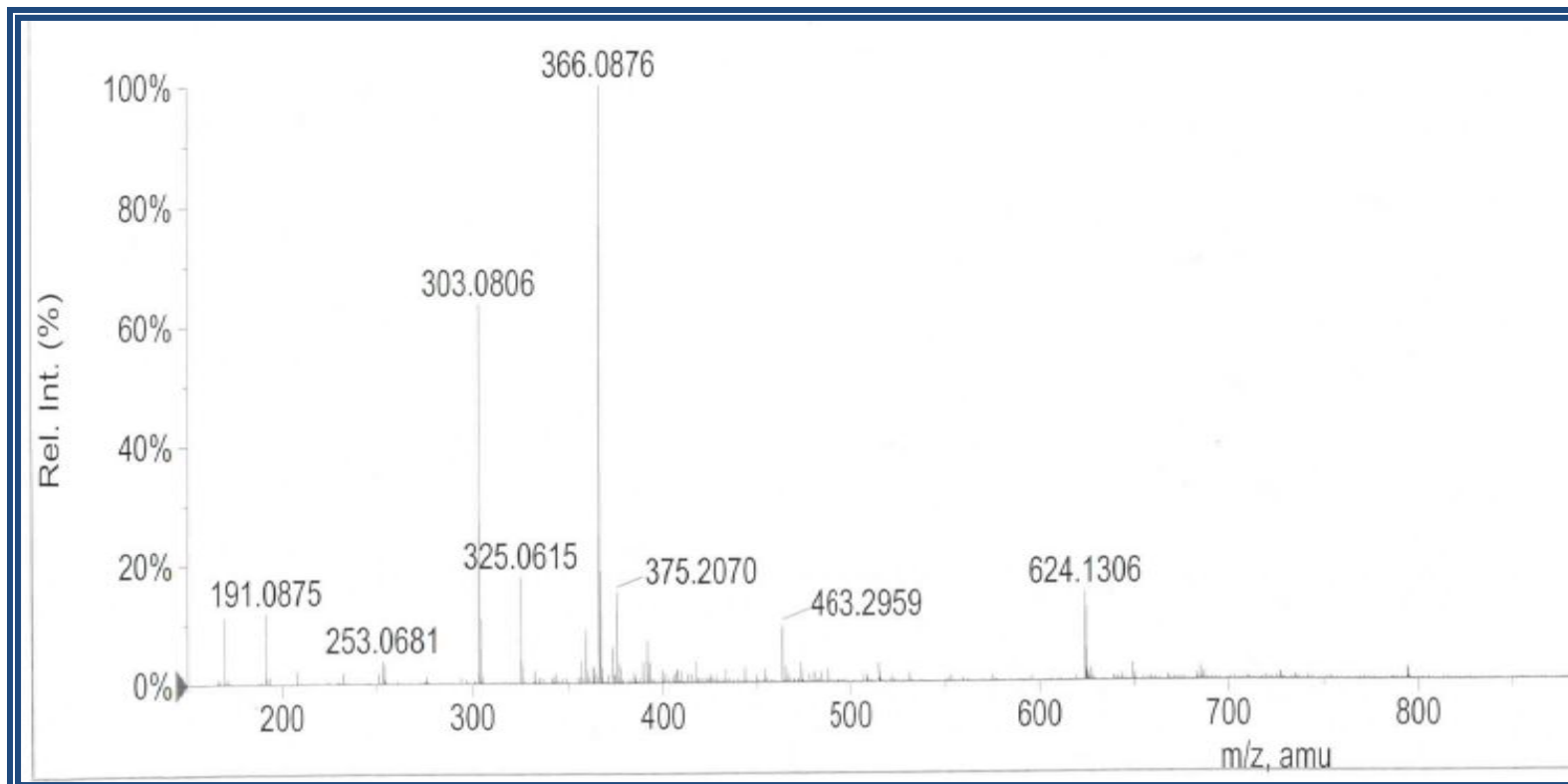
Appendix 3C: HSQC (400 MHz) spectrum of 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**) in DMSO-d₆



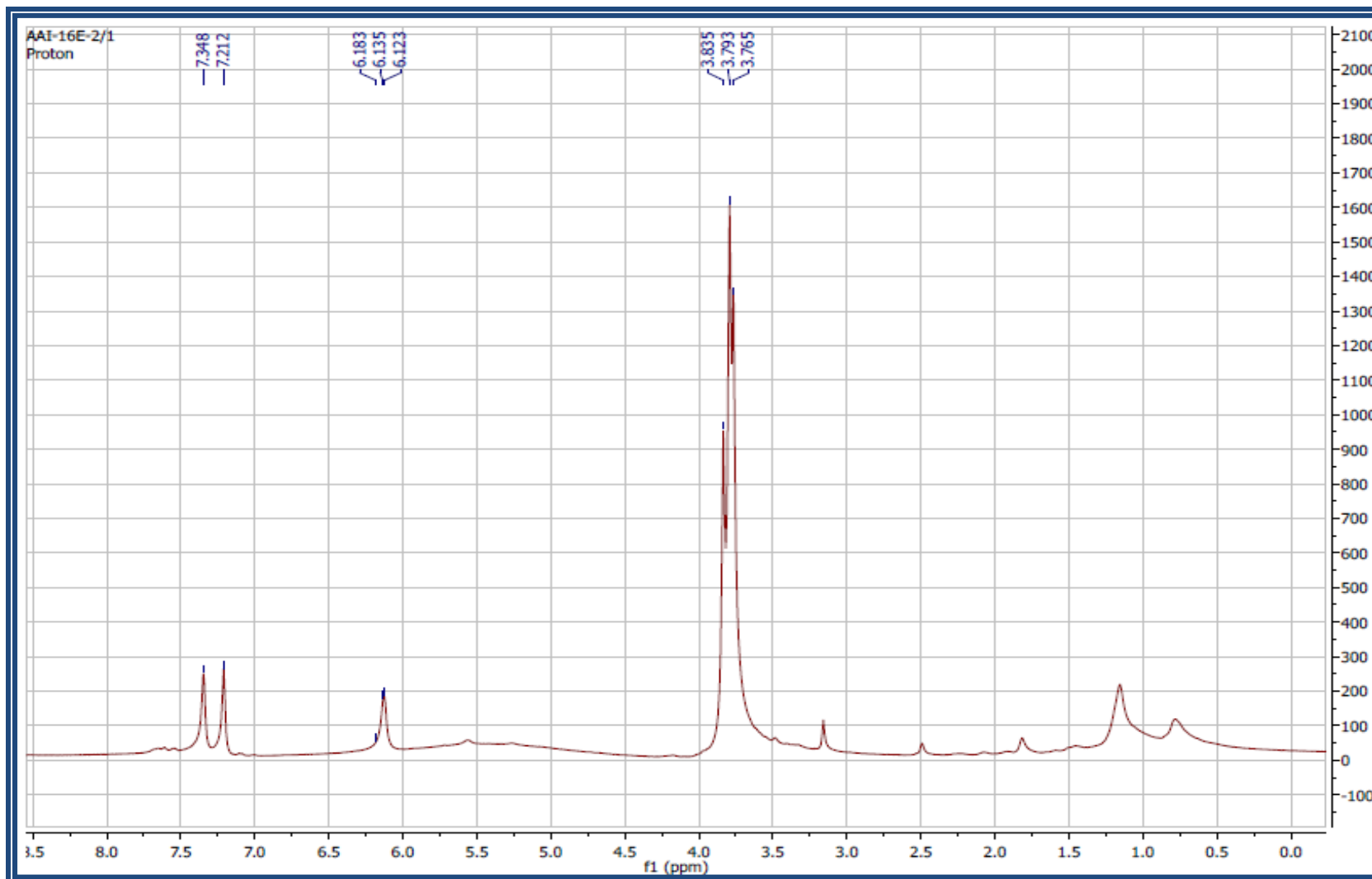
Appendix 3D: HMBC (400 MHz) spectrum of 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**) in DMSO-d₆



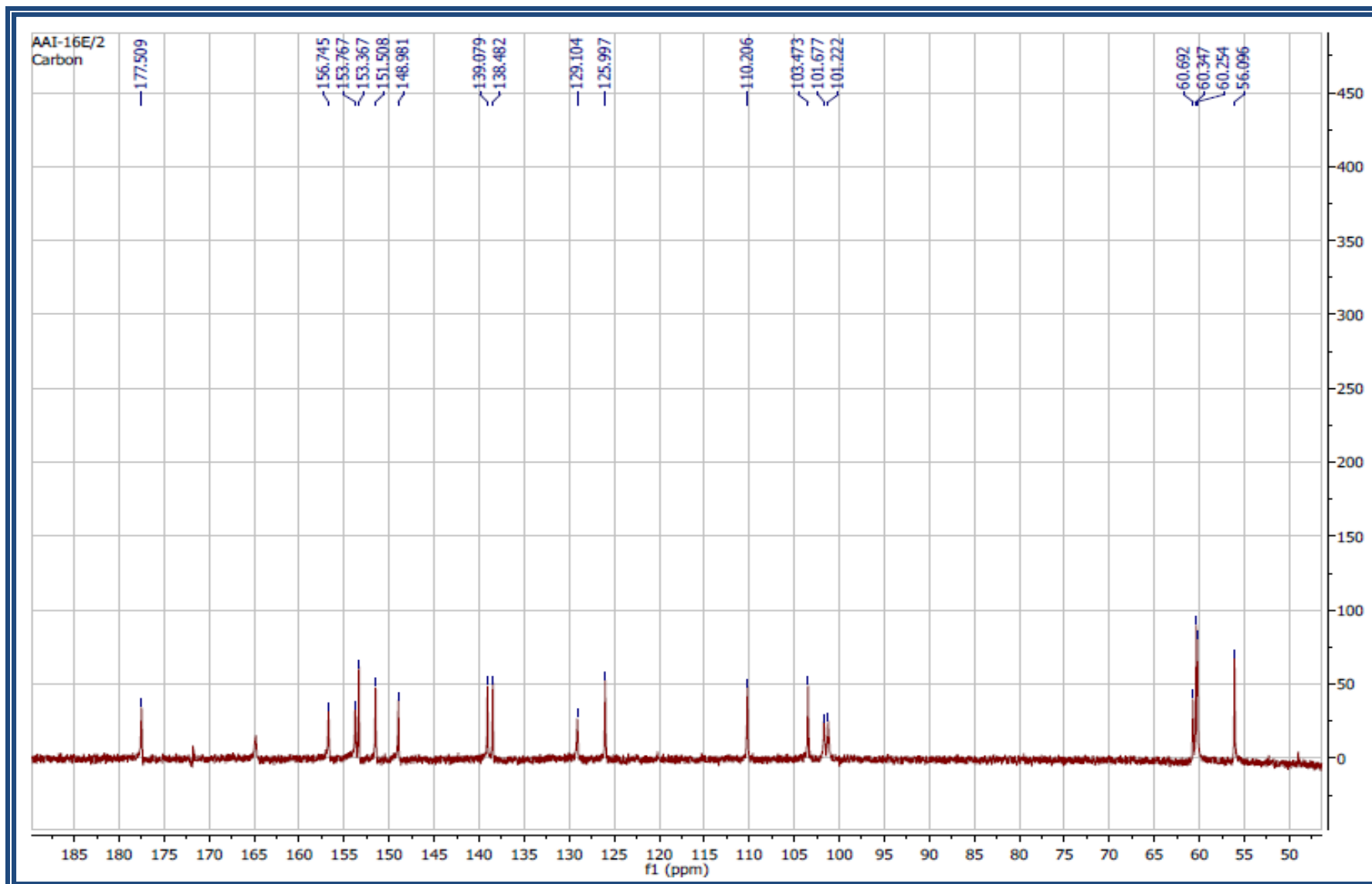
Appendix 3E: HR-EIMS spectrum of 5, 3', 4'-trihydroxy-7-methoxyflavanone (**83**)



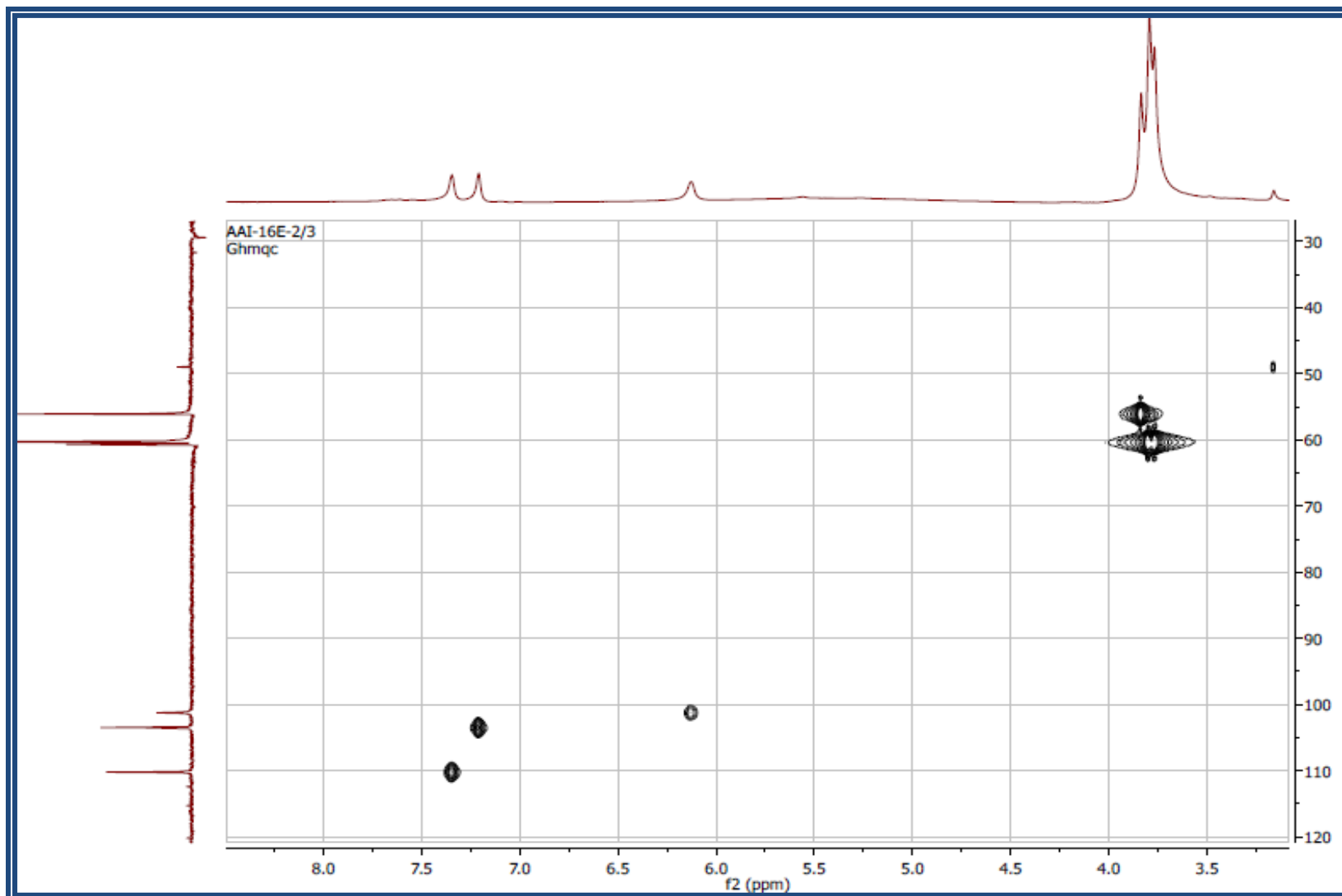
Appendix 4A: ^1H NMR (400 MHz) spectrum of 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**) in DMSO-d_6



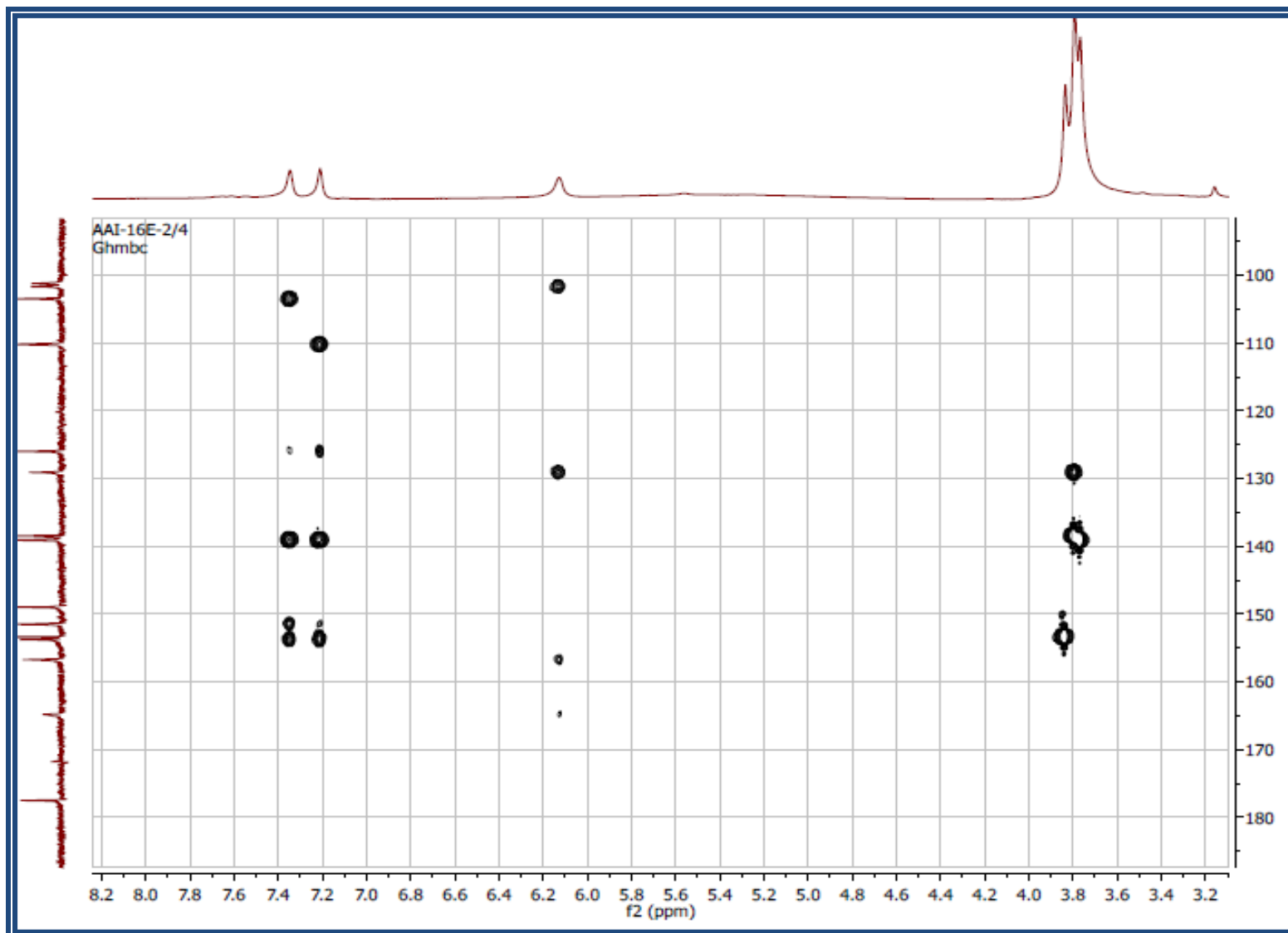
Appendix 4B: ^{13}C NMR (100 MHz) spectrum of 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**) in DMSO-d_6



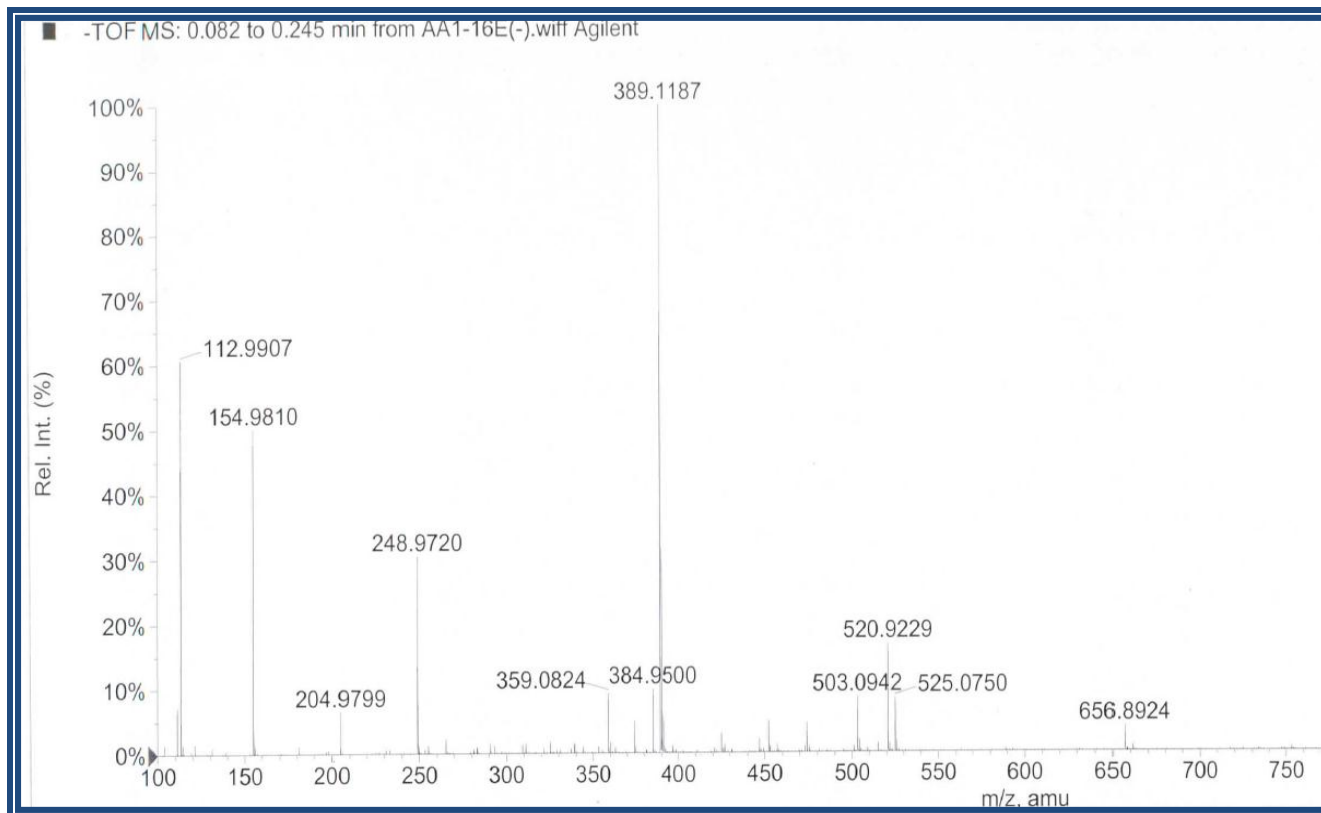
Appendix 4C: HSQC (400 MHz) spectrum of 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**) in DMSO-d₆



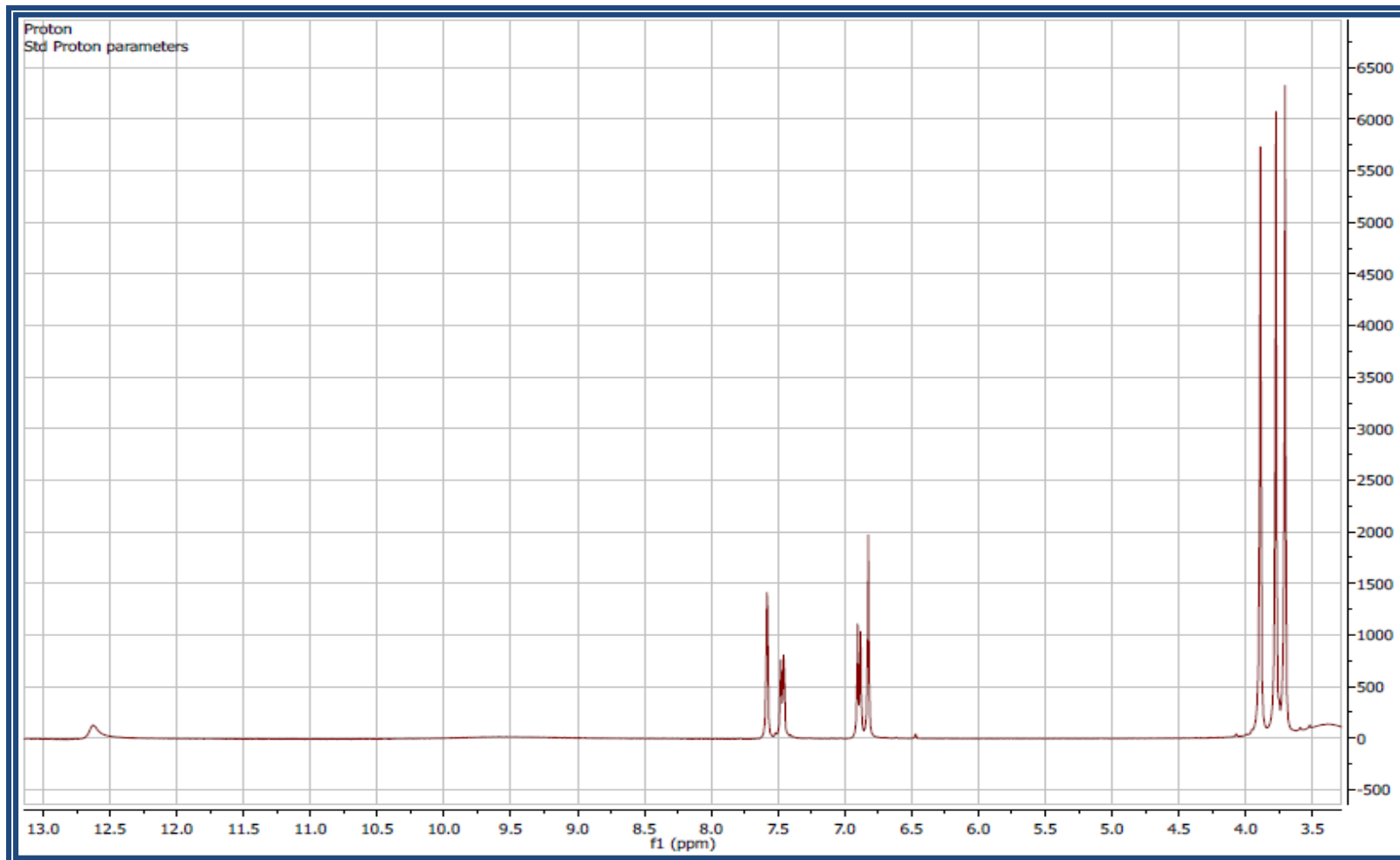
Appendix 4D: HMBC (400 MHz) spectrum of 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**) in DMSO-d₆



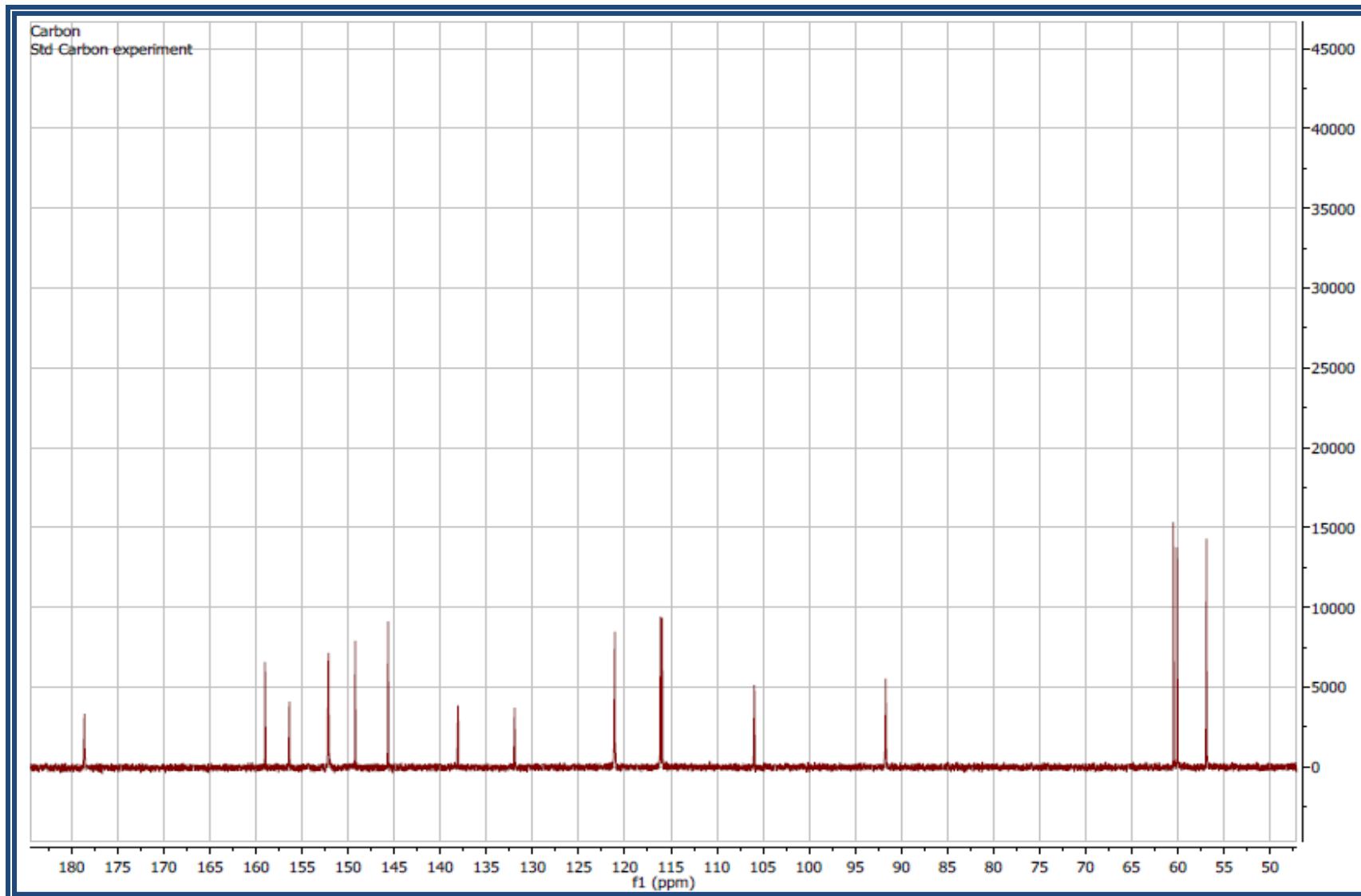
Appendix 4E: HR-EIMS spectrum of 5, 7, 3'-trihydroxy-3, 8, 4', 5'-tetramethoxyflavone (**84**)



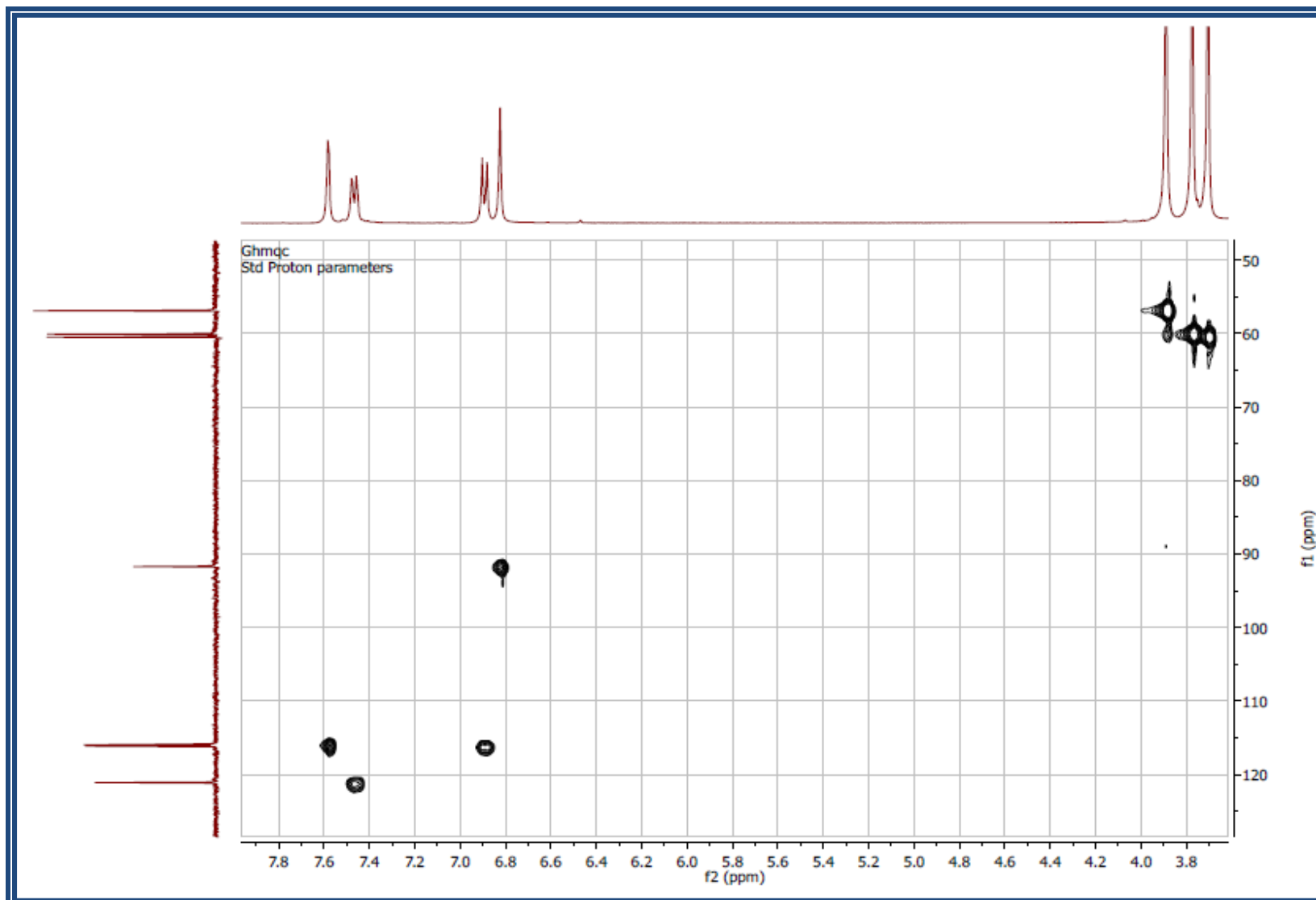
Appendix 5A: ^1H NMR (400 MHz) spectrum of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**) in DMSO-d_6



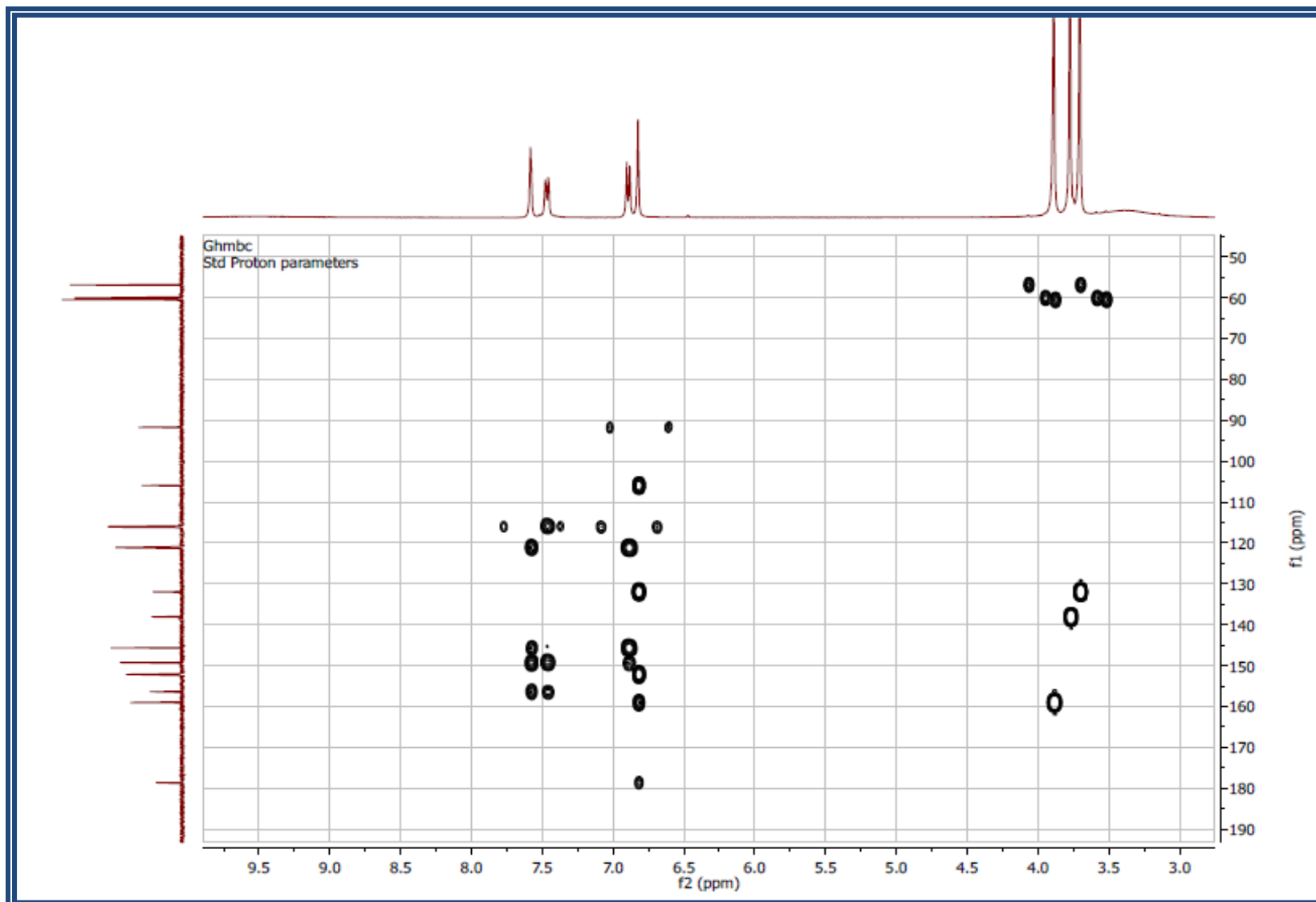
Appendix 5B: ^{13}C NMR (100 MHz) spectrum of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**) in DMSO-d_6



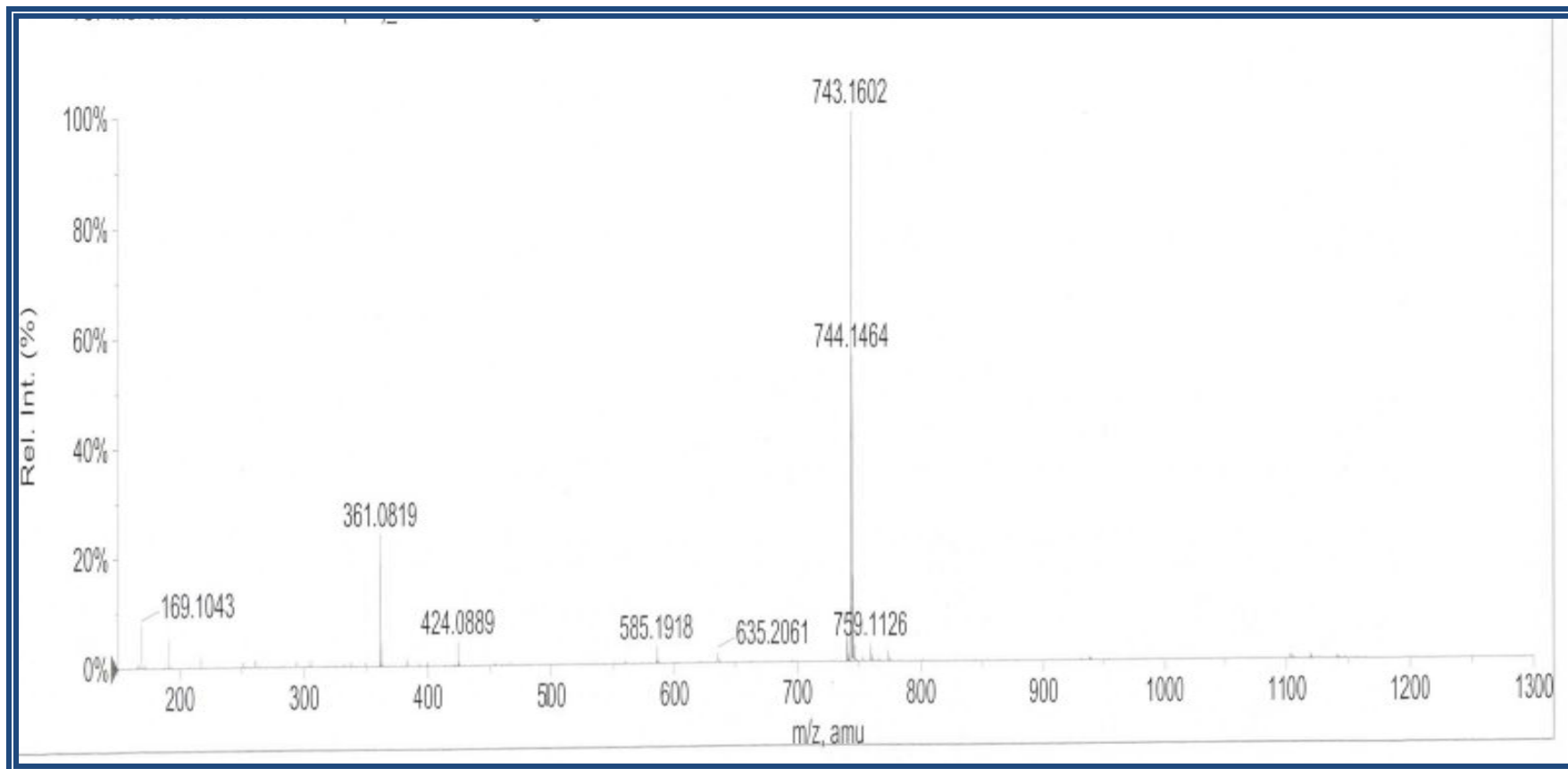
Appendix 5C: HSQC (400 MHz) spectrum of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**) in DMSO-d₆



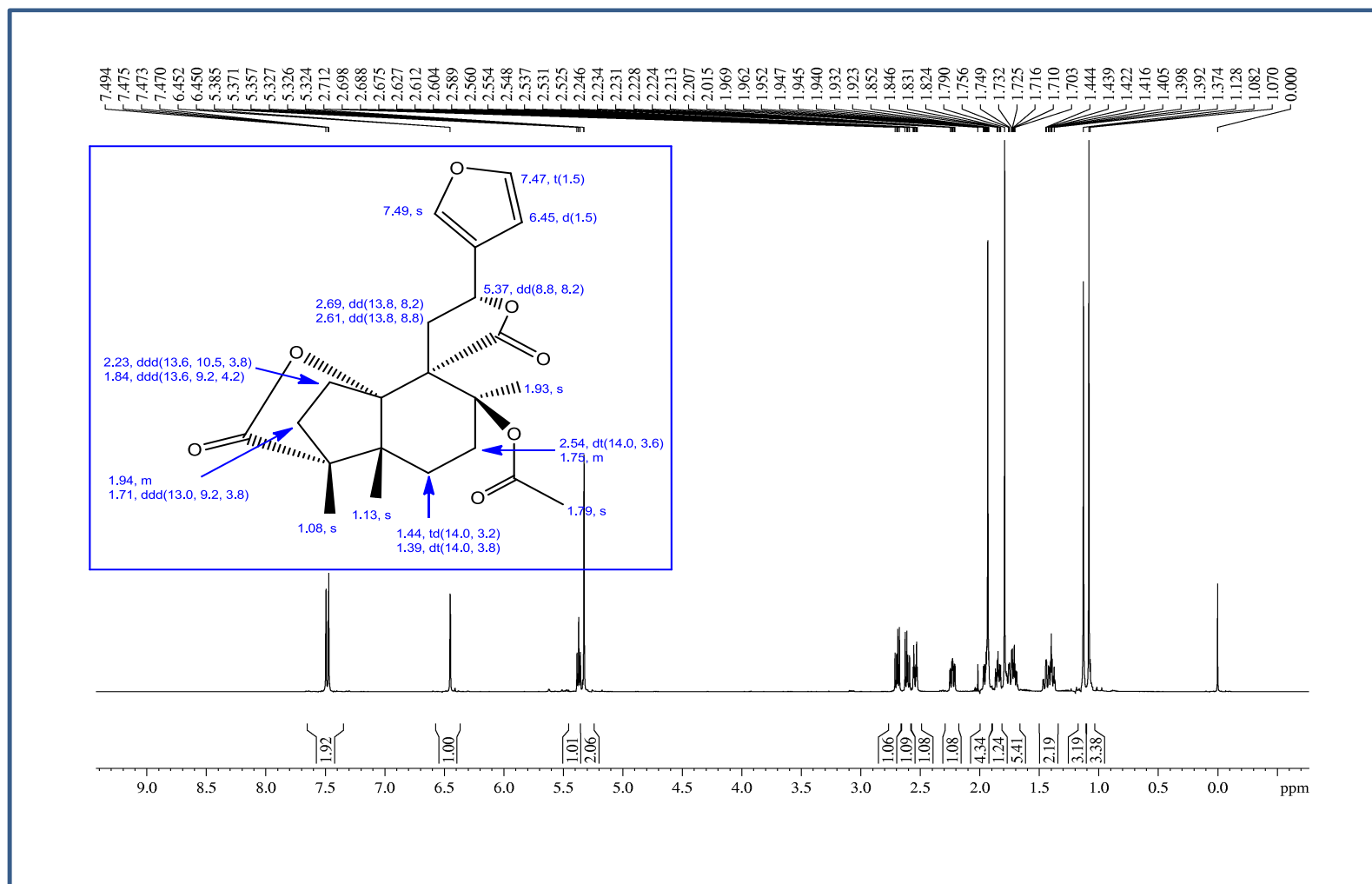
Appendix 5D: HMBC (400 MHz) spectrum of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**) in DMSO-d₆



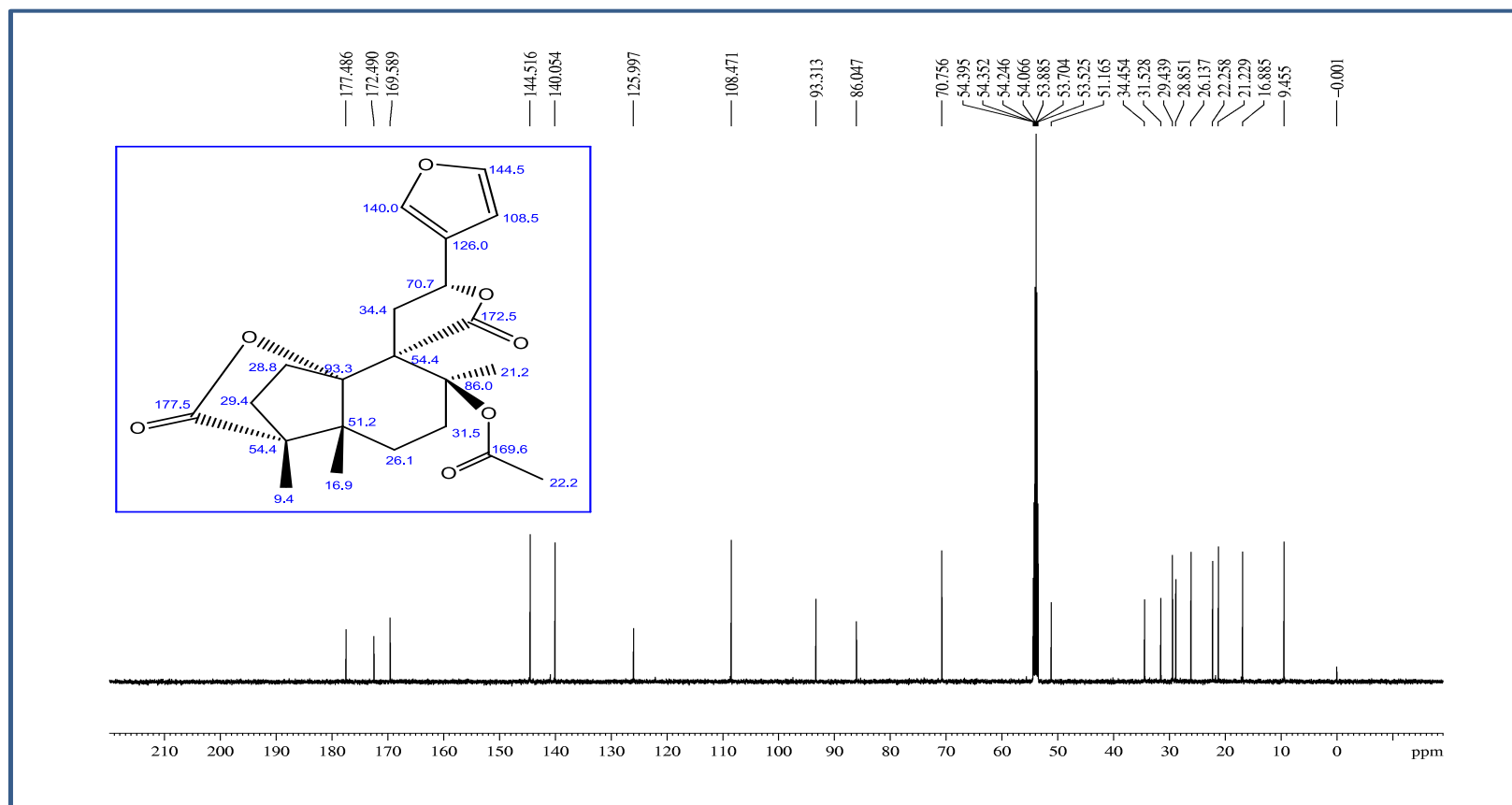
Appendix 5E: HR-EIMS spectrum of 5, 3', 4'-trihydroxy-3, 7, 8-trimethoxyflavone (**85**)



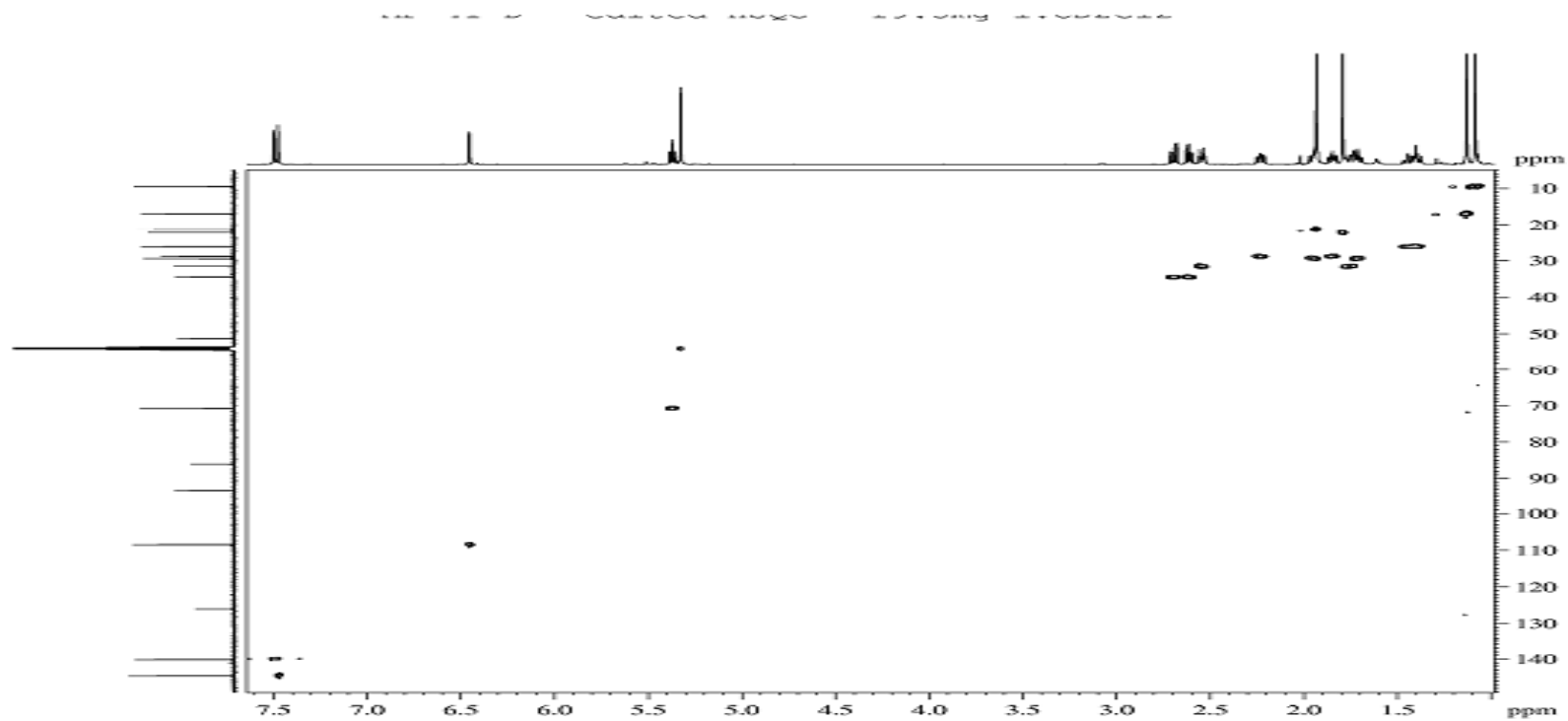
Appendix 6A: $^1\text{H-NMR}$ (600 MHz) spectrum of 8-acetoxyisochiliolide lactone (**86**), CD_2Cl_2 .



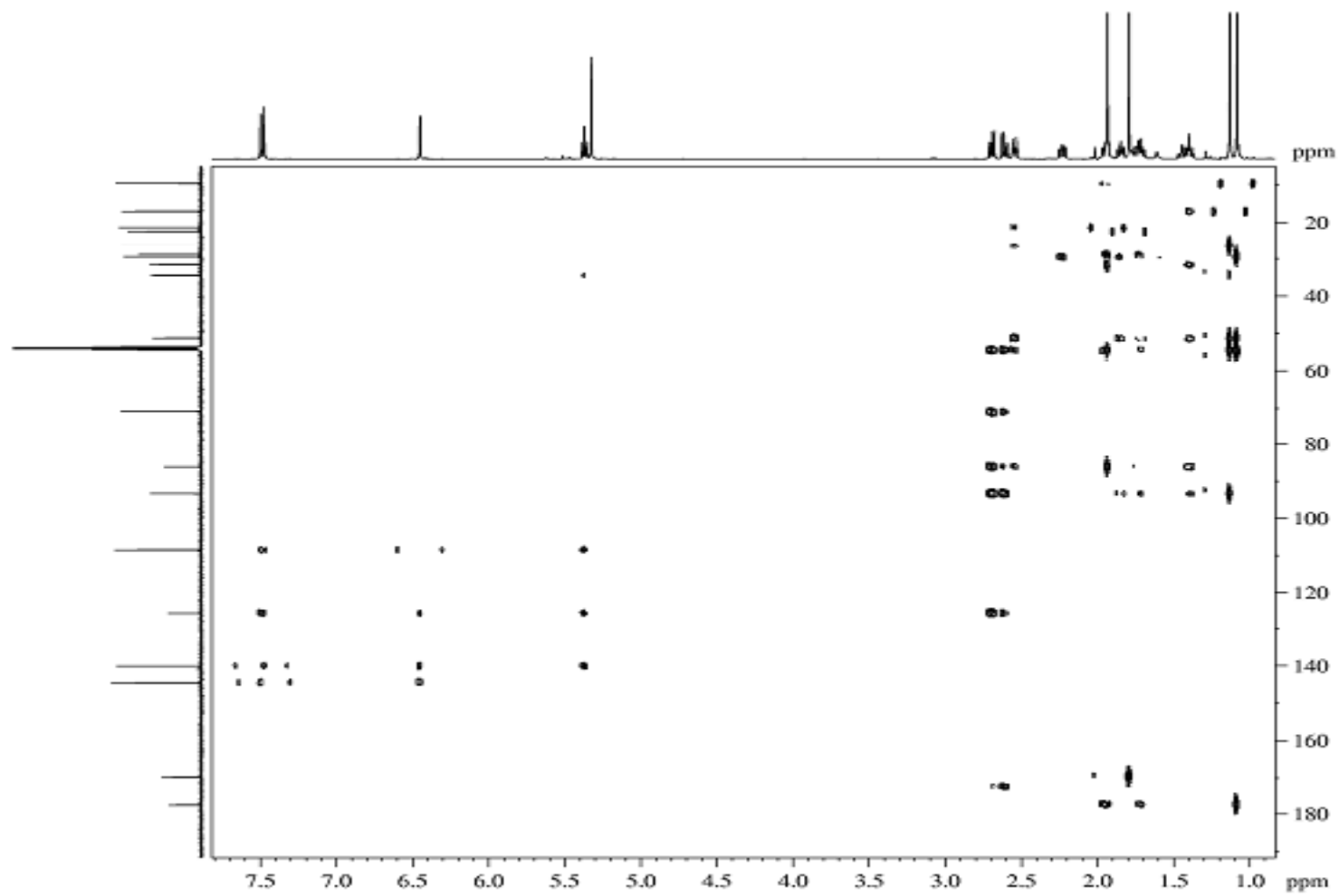
Appendix 6B: ^{13}C NMR (150 MHz) spectrum of 8-acetoxyisochiliolide lactone (**86**), CD_2Cl_2 .



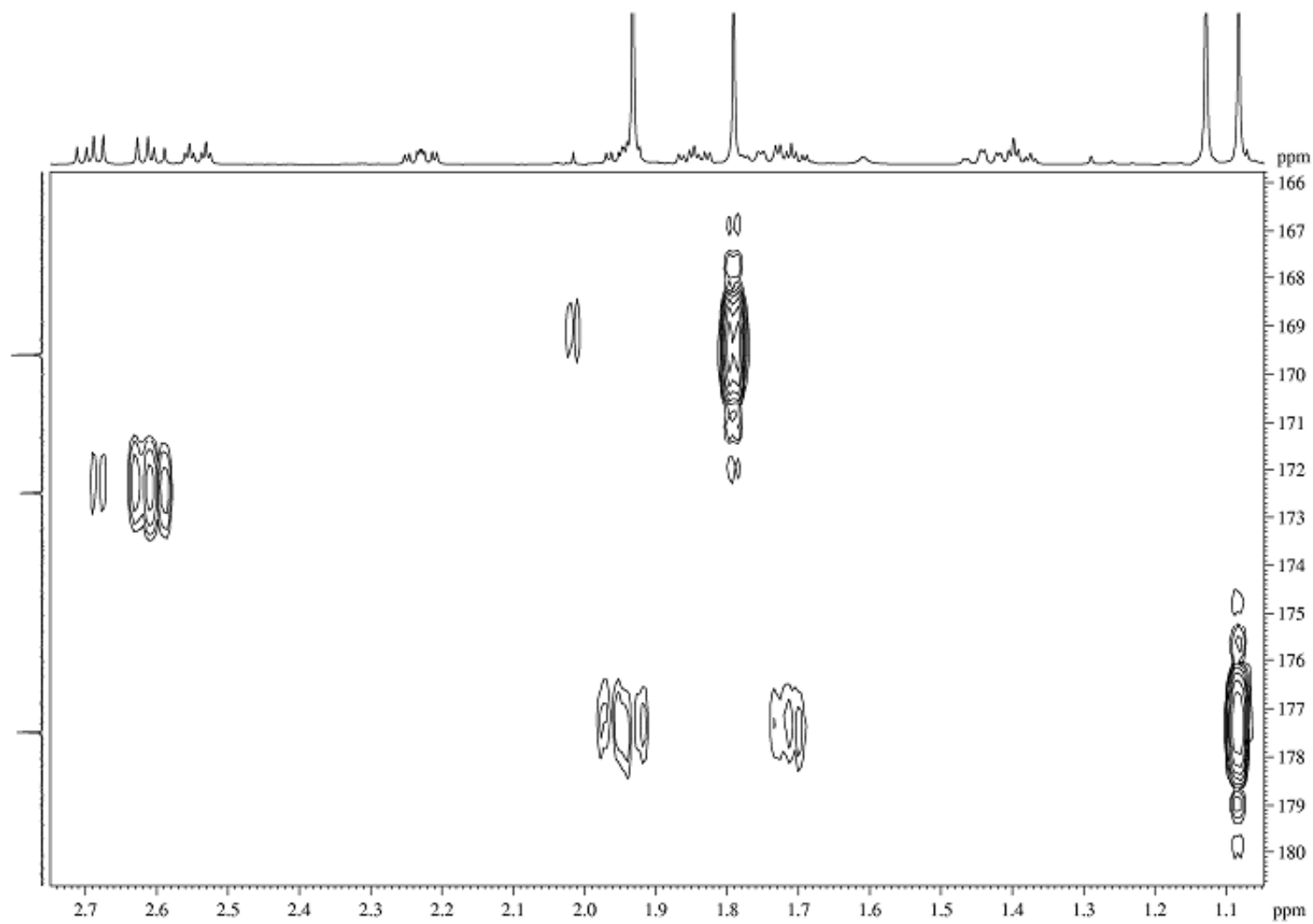
Appendix 6C: HSQC (600 MHz) spectrum of 8-acetoxyisochilolide lactone (**86**), CD₂Cl₂.



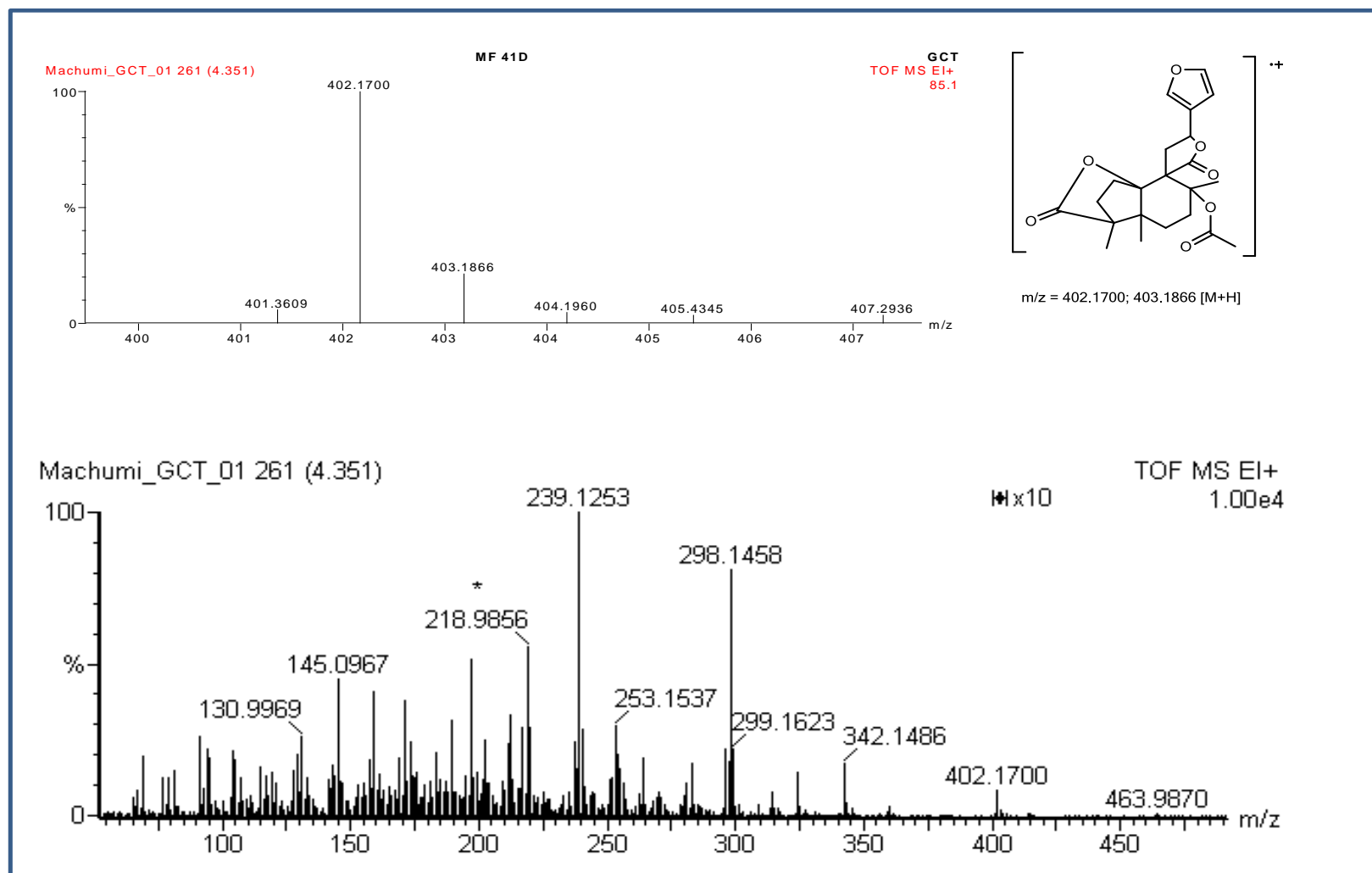
Appendix 6D: HMBC (400 MHz) spectrum of 8-acetoxyisochiliolide lactone (**86**), CD₂Cl₂.



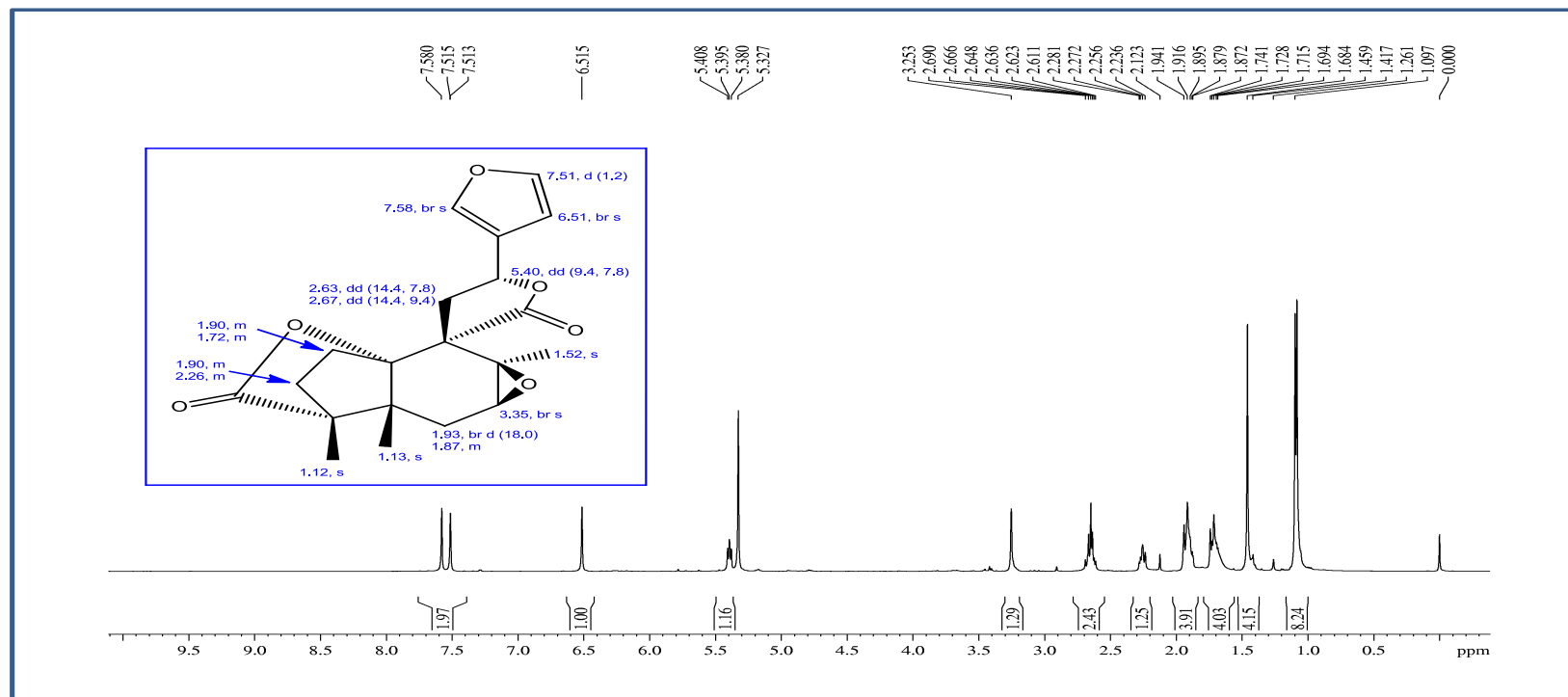
Appendix 6E: HSQC (400 MHz) spectrum of 8-acetoxyisochiliolide lactone (**86**), CD₂Cl₂.



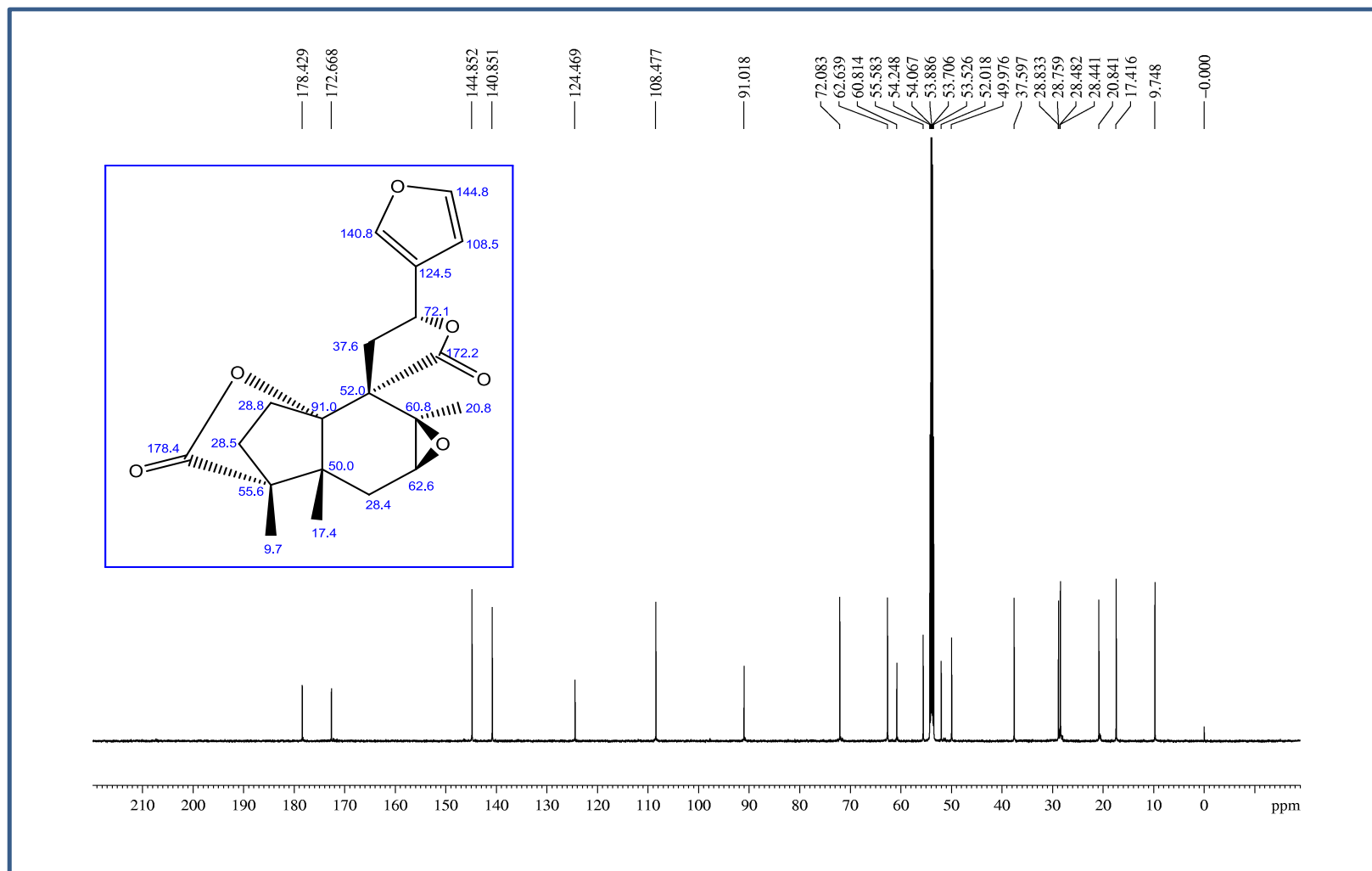
Appendix 6F: HRE-IMS (600 MHz) spectrum of 8 acetoxyisochiliolide lactone (**86**), CD₂Cl₂.



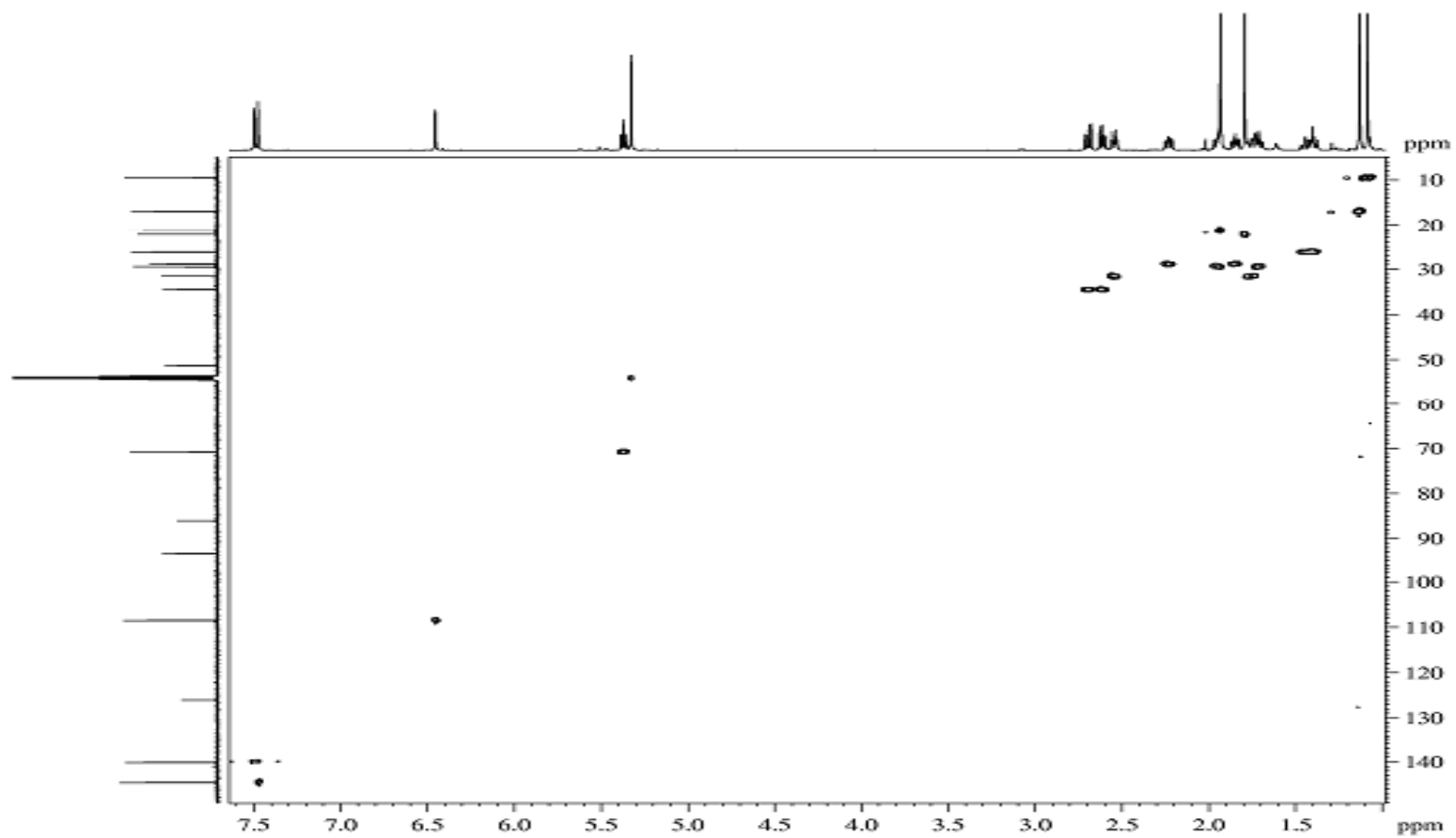
Appendix7A: $^1\text{H-NMR}$ (600 MHz) spectrum of 7,8-epoxyisocholiolide lactone (**87**), CD_2Cl_2 .



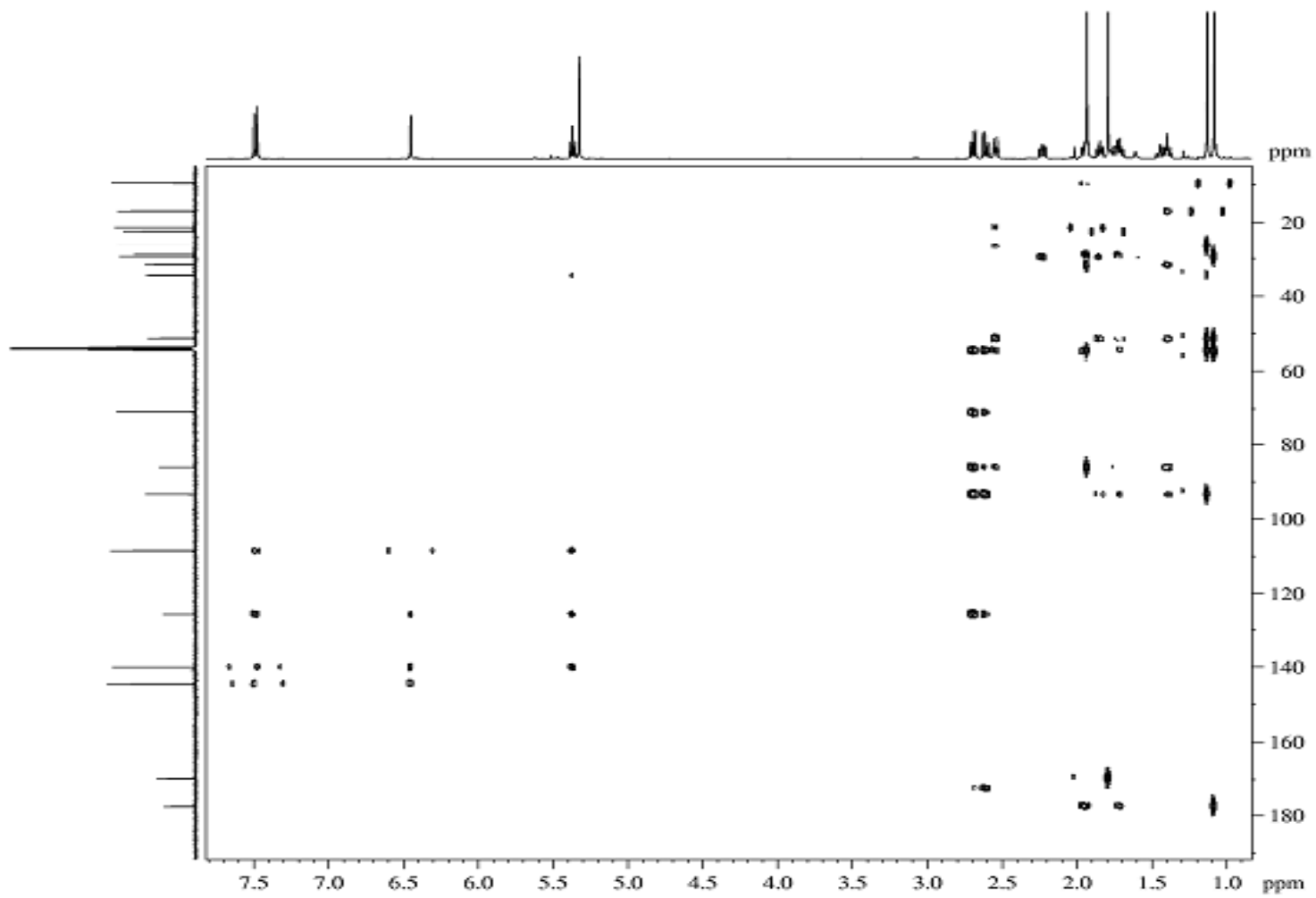
Appendix 7B: ^{13}C -NMR (150 MHz) spectrum of 7,8-epoxyisocholiolide lactone (**87**), CD_2Cl_2 .



Appendix 7C: HMBC (400 MHz) spectrum of 7,8-epoxyisocholiolide lactone (**87**), CD₂Cl₂.



Appendix 7D: HMBC (400 MHz) spectrum of 7,8-epoxyisocholiolide lactone (**87**), CD₂Cl₂.



Appendix 7E: EI-MS spectrum of 7,8- epoxyisochiliolide lactone (87)

