

**INHIBITION OF  $\alpha$ -AMYLASE AND  $\alpha$ -GLUCOSIDASE BY ALKALOIDS AND  
PHENOLICS FROM STEM BARK OF *Zanthoxylum chalybeum* and *Zanthoxylum gillettii***

**BY  
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THE DEGREE OF MASTER OF SCIENCE IN CHEMISTRY**

**SCHOOL OF PHYSICAL AND BIOLOGICAL SCIENCES**

**MASENO UNIVERSITY**

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**DECLARATION**

This thesis is my original work and has never been previously submitted for examination for a degree in Maseno University or any other institution.

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## **DEDICATION**

This thesis is dedicated to my family, and to all that supported me throughout the study period.

## ABSTRACT

Diabetes prevalence is on the rise with conventional drugs either being unaffordable, unavailable or having undesirable side effects. The plants *Zanthoxylum chalybeum* (ZC) and *Zanthoxylum giletii* (ZG) are reported for management of diabetes by local communities. Root bark and stem bark extract from two plants has shown *in vivo* anti-diabetic activity against alloxan streptozotocin-induced diabetic rats. Two benzophenanthridine alkaloids and a phenolic were established to inhibit the  $\alpha$ -amylase and  $\alpha$ -glucosidase activities. Based on the molecular variations of benzophenanthridine alkaloids and phenolics, the inhibitory effects of only three compounds were not predictive enough for the trends of inhibition against  $\alpha$ -amylase and  $\alpha$ -glucosidase. Molecular structural requirement including enzymatic inhibitory modes of alkaloids and phenolics from *Zanthoxylum* plants was also unknown. Further research to establish the structural characteristics exhibiting the enzymatic inhibition of these compounds is thus necessary to provide insight into their potential in hyperglycemic management. Therefore, the study intends to determine the structures of these compounds, their relative  $\alpha$ -amylase and  $\alpha$ -glucosidase inhibitory activities, and kinetic modes of inhibition of the active molecule. The stem bark of ZC was collected from Homa Hills, Kenya and stem bark of ZG from Kakamega forest, Kenya. Both the samples were air dried, ground into fine powder followed by acid-base alkaloid extraction comprising of 25% v/v of aq.NH<sub>4</sub>OH: ethyl acetate-petroleum ether: 2% v/v H<sub>2</sub>SO<sub>4</sub>; chloroform fractionation into alkaloid and non-alkaloids extracts. The fractions were subjected to chromatographic separations then characterization of the pure isolates using mass spectrometry, NMR, and UV-vis spectroscopic techniques. Eleven compounds were isolated: six alkaloids {Norchelerythrine (**9**), Phenanthridine A (**42**), chalybemide A (**39**), chalybemide B (**40**), chalybemide C (**41**) and skimmianine (**21**); two phenolics {2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) and sesamine(**46**)} from *Z. chalybeum*. Whereas *Z. giletii* yielded six compounds: four alkaloids {**9**, **42**, Zanthocapsine (**43**), Zanthoamide D (**45**); and two phenolics {Zanthocapsol (**44**) and sesamine (**46**)}. Ten compounds showed inhibitory activities against  $\alpha$ -amylase and  $\alpha$ -glucosidase in the range of IC<sub>50</sub> = 43.22-49.36  $\mu$ M which had no significant difference ( $P > 0.05$ ) relative to acarbose (IC<sub>50</sub> = 42.67, 44.88) following *in vitro* bioassay by Worthington Enzyme assay protocols except sesamine (IC<sub>50</sub> = 54.67, 54.77) which showed lower activity. The kinetic analysis based on Lineweaver-Burk plots {1/velocity vs 1/[Substrate] at constant [Inhibitor] (sample compounds)} established modes of inhibition and on each kinetic mode, Dixon plots {1/velocity vs [Inhibitor] at constant [Substrate]} were used to establish the respective enzymatic dissociation constants (K<sub>i</sub>). Compound **10** showed competitive inhibition with K<sub>i</sub> of 5.54 and 17.21  $\mu$ M as acarbose (K<sub>i</sub> = 6.14 and 22.40  $\mu$ M) against  $\alpha$ -amylase and  $\alpha$ -glucosidase, respectively. Alkaloids **9**, **21**, **42**, and **43** showed mixed modes of inhibitions (K<sub>i</sub> ranges 2.74-3.10  $\mu$ M and 4.73-9.17  $\mu$ M) in  $\alpha$ -amylase and  $\alpha$ -glucosidase enzymes, respectively. Compounds with amides functionality **39**, **40**, **41**, **45** showed non-competitive inhibition (K<sub>i</sub> = 11.05-26.69 and 17.56-44.58) against  $\alpha$ -amylase and  $\alpha$ -glucosidase, respectively, except compound **40** which showed uncompetitive inhibition (K<sub>i</sub> = 44.58  $\mu$ M) against  $\alpha$ -glucosidase same as **44** (K<sub>i</sub> = 26.28 and 20.62  $\mu$ M). Compounds showing competitive inhibition similarly exhibited stronger binding (lower K<sub>i</sub> values) to the enzymes followed by mixed inhibitors and noncompetitive modes while uncompetitive inhibitors showed the highest K<sub>i</sub>. Compounds **10**, **9**, **21**, **42** and **43** exhibited good activity against  $\alpha$ -amylase and  $\alpha$ -glucosidase thus were considered as lead compounds for management of hyperglycemia. The study offers guidance for future investigations into drug discovery using these kinds of molecular templates.

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## LIST OF ABBREVIATIONS AND ACRONYMS

<b><math>^{13}\text{C}</math> NMR</b>	-	Carbon-13 Nuclear Magnetic Resonance
<b><math>^1\text{H}</math> NMR</b>	-	Proton Nuclear Magnetic Resonance
<b>ANOVA</b>	-	Analysis of Variance
<b>br</b>	-	broad
<b>CC</b>	-	Column Chromatography
<b><math>\text{CDCl}_3</math></b>	-	Deuterated trichloromethane
<b>COSY</b>	-	Correlation Spectroscopy
<b>DMSO</b>	-	Dimethylsulfoxide
<b>d</b>	-	doublet
<b>DBP</b>	-	Dihydrobenzo[c]phenanthridine
<b>DCM</b>	-	Dichloromethane
<b>dd</b>	-	doublet of doublet
<b>DEPT</b>	-	Distortionless Enhancement Polarization Transfer
<b>DM</b>	-	Diabetes Mellitus
<b>DNA</b>	-	Deoxyribonucleic Acid
<b>DNS</b>	-	Dinitrosalicylic acid
<b>EIMS</b>	-	Electron Ionization Mass Spectroscopy
<b>ES</b>	-	Enzyme-substrate
<b>ESI</b>	-	Enzyme-Substrate-Inhibitor
<b>ESIMS</b>	-	Electron spray Ionization Mass Spectrometry
<b>EtOAc</b>	-	Ethyl acetate
<b>GC/MS</b>	-	Gas Chromatography/Mass Spectrometry
<b>GH</b>	-	Glycoside Hydrolase
<b>HBP</b>	-	Hexahydrobenzo[c]phenanthridine
<b>HIV</b>	-	Human Immunodeficiency Virus
<b>HMBC</b>	-	Heteronuclear Multiple Bond Correlation
<b>HMQC</b>	-	Heteronuclear Multiple Quantum Correlation
<b>HPLC</b>	-	High Performance Liquid Chromatography
<b>HREIMS</b>	-	High Electron Ionization Mass Spectroscopy
<b>HSQC</b>	-	Heteronuclear Single Quantum Correlation

<b>Hz</b>	-	Hertz
<b>IC<sub>50</sub></b>	-	Inhibitory Concentration of 50%
<b>J</b>	-	Coupling constant
<b>K<sub>i</sub></b>	-	Dissociation constant
<b>[M]<sup>+</sup></b>	-	Molecular ion
<b>m</b>	-	multiplet
<b>MeOH</b>	-	Methanol
<b>mg</b>	-	milligram
<b>MHz</b>	-	Mega Hertz
<b>mL</b>	-	Milliliter
<b>mM</b>	-	millimole
<b>m.p</b>	-	melting point
<b>MS</b>	-	Mass spectrometry
<b><i>m/z</i></b>	-	Mass to charge ratio
<b>NMR</b>	-	Nuclear Magnetic Resonance
<b>NOE</b>	-	Nuclear Overhauser Effect
<b>NOESY</b>	-	Nuclear Overhauser Effect Spectroscopy
<b>ppm</b>	-	parts per million
<b>PTLC</b>	-	Preparative Thin Layer Chromatography
<b>QBP</b>	-	Quaternary benzo[c]phenanthridine
<b>R<sub>f</sub></b>	-	Retention factor
<b>s</b>	-	singlet
<b>SAR</b>	-	Structure-Activity Relations
<b>t</b>	-	triplet
<b>TLC</b>	-	Thin Layer chromatography
<b>TMS</b>	-	Tetramethylsilane
<b>UV-Vis</b>	-	Ultraviolet-visible
<b>WHO</b>	-	World Health Organization
<b>ZC</b>	-	<i>Zanthoxylum chalybeum</i>
<b>ZG</b>	-	<i>Zanthoxylum gillettii</i>
<b>δ</b>	-	Chemical shift in pp

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# CHAPTER ONE

## INTRODUCTION

### 1.1 Background Information

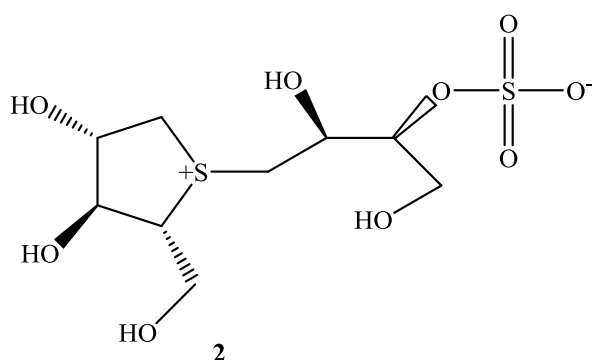
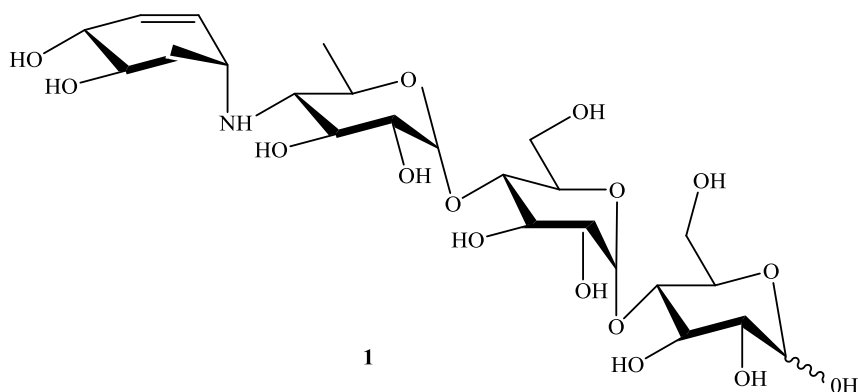
Diabetes mellitus, a metabolic disease resulting from irregularities in insulin secretion and activity, is characterized by chronic hyperglycemia (WHO, 1999). This occurs when the pancreas is unable to produce enough insulin (Ünal *et al.*, 2012; Gardner and Shoback, 2011). Insulin receptor sensitivity or insufficiency is a major factor in type 1 and type 2 diabetes as well as gestational diabetes. Type 1 diabetes is brought about by the pancreas' incapacity to produce enough insulin, whereas gestational diabetes develops during pregnancy. Type 2 diabetes is a complex metabolic disorder marked by prolonged hyperglycemia (Fernandez-Mejia, 2013) and defective insulin action (Lin and Sun, 2010), often linked to overproduction of free radicals (Johansen *et al.*, 2005). Despite being less common than Type 1, Type 2 diabetes remains a significant challenge and requires more research attention.

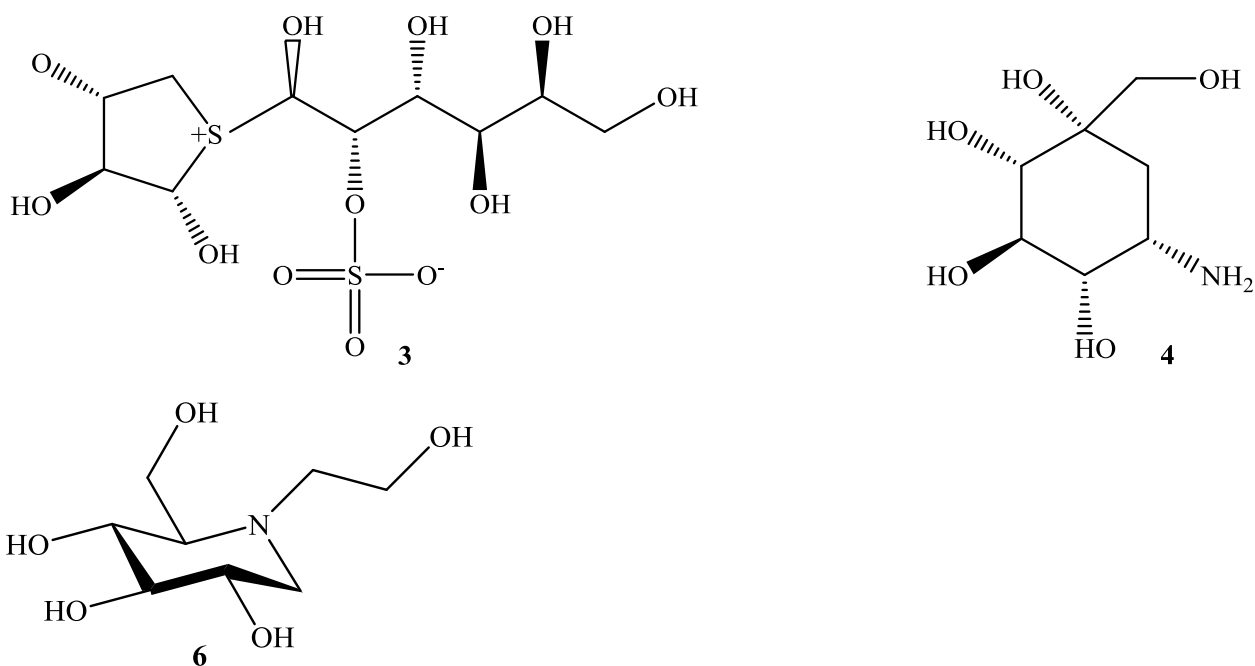
By the year 2016, 422 million people had been diagnosed with diabetes (WHO, 2016). In addition 80% of deaths from the disease in low- and middle-income nations had occurred by 2006 (Mathers and Loncar, 2006). Factors contributing to this increase include a shift towards processed foods, urbanization, and a decline in physical activities, as well as inadequate management techniques and faulty diagnosis (WHO, 2016). Type 1 diabetes is normally managed by insulin injection therapy (Gough and Narendan, 2010), while type 2 diabetes is managed by oral anti-diabetic medications like sulphonylureas, biguanides, thiazolidinediones, and  $\alpha$ -glucosidase inhibitors (Krentz and Bailey, 2005). However, many reports of unsuccessful all-round therapy suggest the need for further research on more effective treatments.

Diabetes management involves managing starch digestion and absorption rates using starch digestive enzyme inhibitors moderate postprandial hyperglycemia (Ye *et al.*, 2008). These enzymes including Glycosidase Hydrolases (GH), break down glycosidic bonds in starch, producing simple sugars (Lombard *et al.*, 2013). The key enzymes include  $\alpha$ -amylase which break down internal  $\alpha$ -1,4-glycosidic bonds (Rajagopalan and Krishnan, 2008; Gupta *et al.*, 2011), while brush border hydrolases (sucrase-amylase, maltase-gluco-amylase,  $\alpha$ -glucosidase and amylase) digest dietary carbohydrates in the small intestine (Hooton *et al.*, 2015). Modulating these enzymes may predispose the body to positive pathophysiological conditions, potentially aiding in diabetes type 2 management. The search for natural modulators of these enzymes is thus envisaged as safe avenue for discovery alternative anti-diabetic options (Gupta *et al.*, 2011).

Shai *et al.* (2010) suggested that employing natural  $\alpha$ -amylase inhibitors derived from food and medicinal plants is a better option for controlling hyperglycemia by decreasing post-prandial serum glucose levels. Notably, a number of  $\alpha$ -amylase inhibitors, including nojirimycin, valioline, voglibose, and acarbose (Niwa *et al.*, 1970) as well as Miglitol (**6**), salacinol (**2**) and kotalanol (**3**) (Yoshikawa *et al.*, 1998), which were isolated either from higher plants or lower plants then later synthesized. However, many of the synthetic analogues have shown certain limitations including being non-specific, displaying serious side effects and failure to manage hyperglycemia successfully.

Grant (2003), highlighted the limitations and high costs of prescription drugs, particularly insulin, in managing diabetes. Furthermore, with prolonged management of diabetes mellitus with insulin has specific disadvantages, including brain atrophy, insulin resistance, anorexia nervosa, fatty liver tissues and (Verspohl, 2012). It is therefore necessary to look at alternative therapy and management approaches in light of these restrictions. Natural sources like plants offer an alternative management resources, but their applications are limited due to insufficient coherent scientific information to support the usage.





Traditional medicine suggests that various plants have anti-diabetic properties, benefiting patients with diabetic retinopathy, insulin-dependent and non-insulin-dependent diabetes as well as diabetic peripheral neuropathy (Trojan-Rodrigues *et al.*, 2012). Among these plants are *Zanthoxylum chalybeum* and *Zanthoxylum gilletii* (Agwaya *et al.*, 2016; Kokwaro, 2009). *Z. chalybeum* has anti-hyperglycemic effects *in vivo*, while *Z. gilletii* manages various ailments such as malaria, tuberculosis, including viral and cancerous (Kokwaro, 2009). Both plants have anticancer, anti-leukemia, antimicrobial (Nakanishi *et al.*, 1998), anti-viral (Tan *et al.*, 1991), and anti-diabetic properties (Mbaze *et al.*, 2007). The plants were separately noted to be used in management diabetic cardiomyopathy (Agwaya *et al.*, 2016). However, possible chemical leads displaying these different beneficial effects remain uncertain.

The phytochemical information on *Z. gilletii* and *Z. chalybeum* and their anti-diabetic activities are inconsistent, with only 3- $\beta$ -acetoxy-16- $\beta$ -hydroxybetulinic acid showing mild  $\alpha$ -glucosidase inhibitory effects. Triterpenoids isolated from *Z. chalybeum* displayed mild inhibitory activities against  $\alpha$ -amylase. This previous study pointed out *in vitro* inhibitory effects against  $\alpha$ -glucosidase and  $\alpha$ -amylase activities of two benzophenanthridine alkaloids, while lupeol, 3 $\alpha$ , 20-hydroxy-3-oxo-28-lupanoic acid, 20-dihydroxy-28-lupanoic acid and 3 $\alpha$ ,20,28-trihydroxylupane registered low activity (Wasiali, 2019).

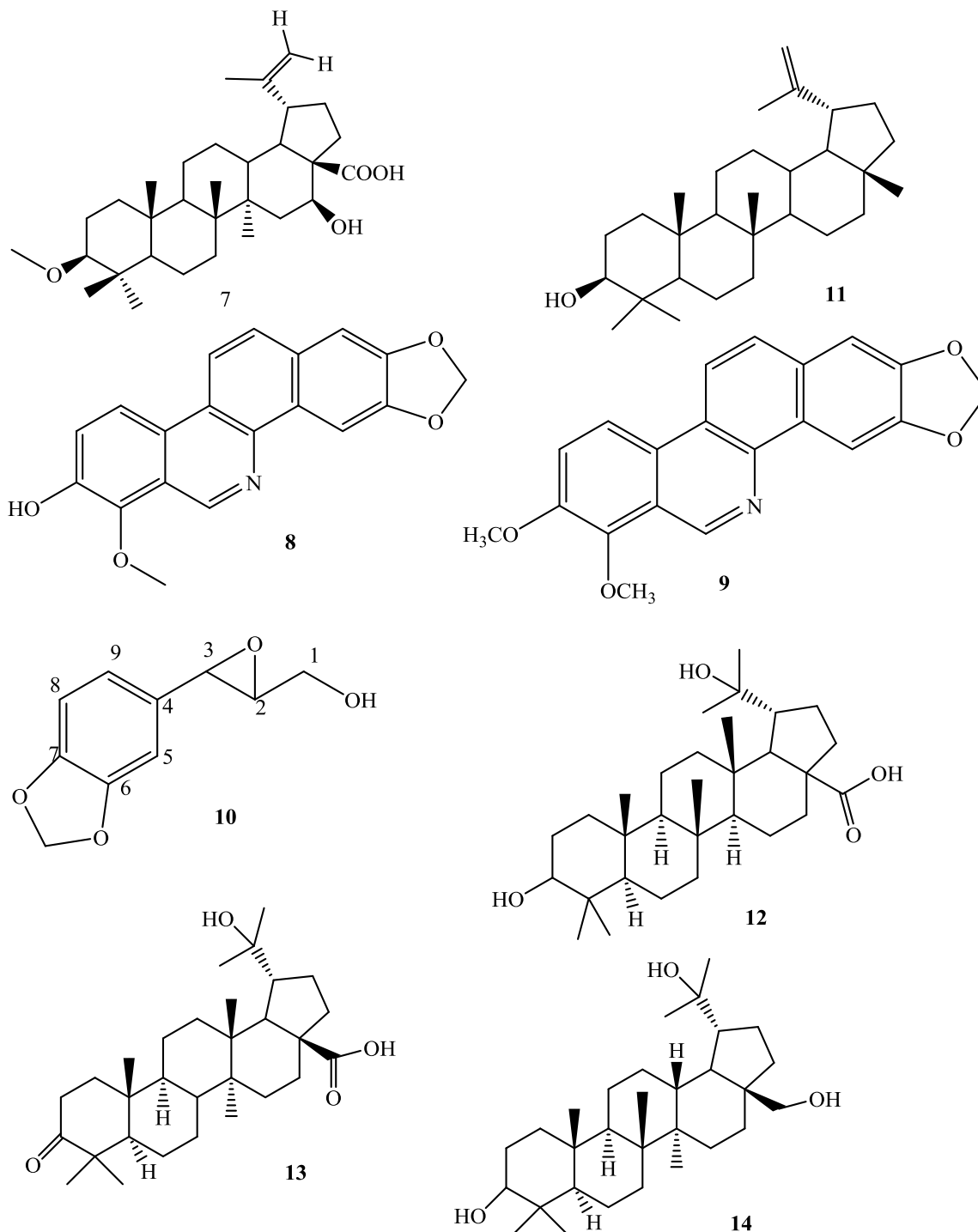
Phytochemical information linking the two plants to management of diabetic conditions are inconsistent, for instance *Z. gilletii* is known for furoquinoline alkaloids, cinnamic acid amides and benzophenanthridine alkaloids (Islam *et al.*, 2001) which have not been evaluated for anti-

hyperglycemic potential. Except for 3- $\beta$ -acetoxy-16- $\beta$ -hydroxybetulinic acid (**7**), a triterpenoid from the *Z. gilletii* stem bark that showed mild  $\alpha$ -glucosidase inhibitory effects (Mbaze *et al*, 2007). Conversely, triterpenoids isolated from *Z. chalybeum* showed mild activities against  $\alpha$ -amylase inhibitory effects relative to benzophenanthridine alkaloids against  $\alpha$ -amylase inhibitory effects (Wasiali, 2019). Two benzophenanthridine alkaloids {6-hydroxy-N-methyl decarine (**8**) and norchelerythrine (**9**)} and a phenolic {2,3-epoxy-6,7-methylenedioxyconiferyl (**10**)} showed potent *in vitro* inhibitory effects against  $\alpha$ -glucosidase and  $\alpha$ -amylase activities, while lupeol, 3 $\alpha$ ,20-dihydroxy-28-lupanoic acid, 20-hydroxy-3-oxo-28-lupanoic acid, and 3 $\alpha$ ,20,28-trihydroxylupane registered low activity (Wasiali, 2019). This means that the most active compounds were not actually the triterpenoids in *Z. gilletii*, but since triterpenoids are stable and are easy to isolate, consequently the only pure isolate that was tested but in the crude some other compounds could have been active as well but might have not been isolated.

The anti-hyperglycemic potentials of *Zanthoxylum* has been attributed to the existence of benzophenanthridine alkaloids together with additive effects of phenolics compounds (Wasiali, 2019). However, the genus is known to elaborate different classes of benzophenanthridine alkaloids with marked structural variation that normally exhibit variable biological activities (Tavares *et al*, 2014). Generally, benzophenanthridine alkaloids are classified into three classes: dihydrobenzo[c]phenanthridine-type (DBP), quaternary benzo[c]phenanthridine-type (QBP), and hexahydrobenzo[c]phenanthridine-type (HBP) (Krane *et al*, 1984). The two alkaloids (**8** and **9**) reported as active against  $\alpha$ -glucosidase and  $\alpha$ -amylase structurally belong to the DBP although the other two (HBP and QBP) forms are also noted with scaffold necessary for specific biological activities (Tavares *et al*, 2014). Since all these benzophenanthridine alkaloids types and their various derivatives have been isolated from the genus *Zanthoxylum*, it implies existences of interesting molecular chromophores worth exploring. The activity of only one type cannot qualify that all the benzophenanthridine alkaloids and phenolic from *Zanthoxylum* plants are active against  $\alpha$ -glucosidase and  $\alpha$ -amylase hydrolytic effects on carbohydrates. To this extent, the varied functionalities among benzophenanthridine alkaloids and phenolic that predispose inhibition of both  $\alpha$ -amylase and  $\alpha$ -glucosidase are unknown.

Based on chemotaxonomic similarities, the benzophenanthridine alkaloids and phenolics from *Z. gilletii* and that of *Z. chalybeum* presumably should display common biological activities which vary depending on difference in molecular structures. Such an assertion, endorses structure-activity relationship study on benzophenanthridine alkaloids and phenolics from these plants, which then require targeted systematic extraction and separation procedures for alkaloids and phenolics.

Besides, establishing the possible activity trends among these benzophenanthridine alkaloids and phenolics together with their possible inhibitory kinetics parameters would determine the precise mode of inhibition as either uncompetitive, noncompetitive, mixed inhibition or competitive inhibition, are unknown.



## 1.2 Statement of the Problem

The plants, *Zanthoxylum chalybeum* and *Zanthoxylum gillettii* have been identified as medicinal plants with potential for managing diabetes including hyperglycemia although there is scanty pharmacological evidence to validate the claims. Reported data indicate two benzophenanthridine alkaloids and a phenolic from *Zanthoxylum chalybeum* as active against  $\alpha$ -amylase activity despite the different classes of these compounds elaborated by the genus. Based on the varied classes of benzophenanthridine alkaloids and phenolics, inhibitory effects of only one class cannot qualify that all benzophenanthridine alkaloids from *Zanthoxylum* genus are active as  $\alpha$ -glucosidase and  $\alpha$ -amylase inhibitors hydrolytic effects on carbohydrates. Data that can be used for predicting the activities of these compounds is lacking. In other words, the varied functionalities among benzophenanthridine alkaloids and phenolic that predispose inhibitory effects on both  $\alpha$ -glucosidase and  $\alpha$ -amylase are unknown. Besides, the possible enzymatic inhibitory activities trends, the inhibitory kinetics modes have not been established among these compounds.

## 1.3 Objectives

### 1.3.1 General Objective

To determine molecular structures of alkaloids and phenolics from the stem barks of *Zanthoxylum chalybeum* and *Zanthoxylum gillettii* and their kinetic modes of inhibition towards  $\alpha$ -amylase and  $\alpha$ -glucosidase activities.

### 1.3.2 Specific Objectives

- i. To isolate and elucidate structures of compounds from the stem bark of *Zanthoxylum chalybeum* and *Zanthoxylum gillettii* using chromatographic and spectroscopic techniques.
- ii. To ascertain the relative inhibitory activity of the isolated compounds from objective (i) against  $\alpha$ -amylase and  $\alpha$ -glucosidase activities on carbohydrate substrate.

- iii. To determine the modes of inhibition of the active isolated compounds from the objective (ii) against  $\alpha$ -amylase and  $\alpha$ -glucosidase activities on carbohydrates substrate.

#### **1.4 Null Hypothesis**

Benzophenanthridine alkaloids and phenolics from *Zanthoxylum chalybeum* and *Zanthoxylum gilletii* do not significantly inhibit activities  $\alpha$ -amylase and  $\alpha$ -glucosidase relative to a standard drug (acarbose).

##### **1.4.1 Research Questions**

What are the molecular structural variations of alkaloids and phenolic compounds present in the stem bark of *Z. chalybeum* and *Z. gilletii* which are responsible for amylase and glucosidase inhibition?

What are the modes/mechanism of inhibition of  $\alpha$ -amylase and  $\alpha$ -glucosidase enzymes by alkaloids and phenolics from *Z. chalybeum* and *Z. gilletii* stem bark?

#### **1.5 Justification of the Study**

Diabetes mellitus is a global epidemic for which there are no effective treatments or their side effects are either unfavorable. Because most people in underdeveloped countries find herbal medications to be acceptable, economical, and easily available, they are growing in popularity. Establishing an efficient supplementary and alternative diabetes management strategy is necessary for the impoverished communities in developing nations, as the current anti-hyperglycemic medicines are not very effective and have undesirable side effects. Herbal remedies, of which *Zanthoxylum spp.* are a plentiful source, may provide such safe and efficient substitutes. Based on the few reports inclining the metabolites from these plants especially alkaloids and phenolics which vary depending on difference in molecular structural features. Molecular variation normally predisposes varied biological activities, which require predictive structure-activity relationship (SAR) study which can be used for development of active drug



components. Moreover, establishment of the possible activity trends among these alkaloids and phenolics together with their possible kinetics modes of inhibition would provide necessary information relevant for the development of the urgently need alternative medicinal remedies from natural resource.

## CHAPTER TWO

### LITERATURE REVIEW

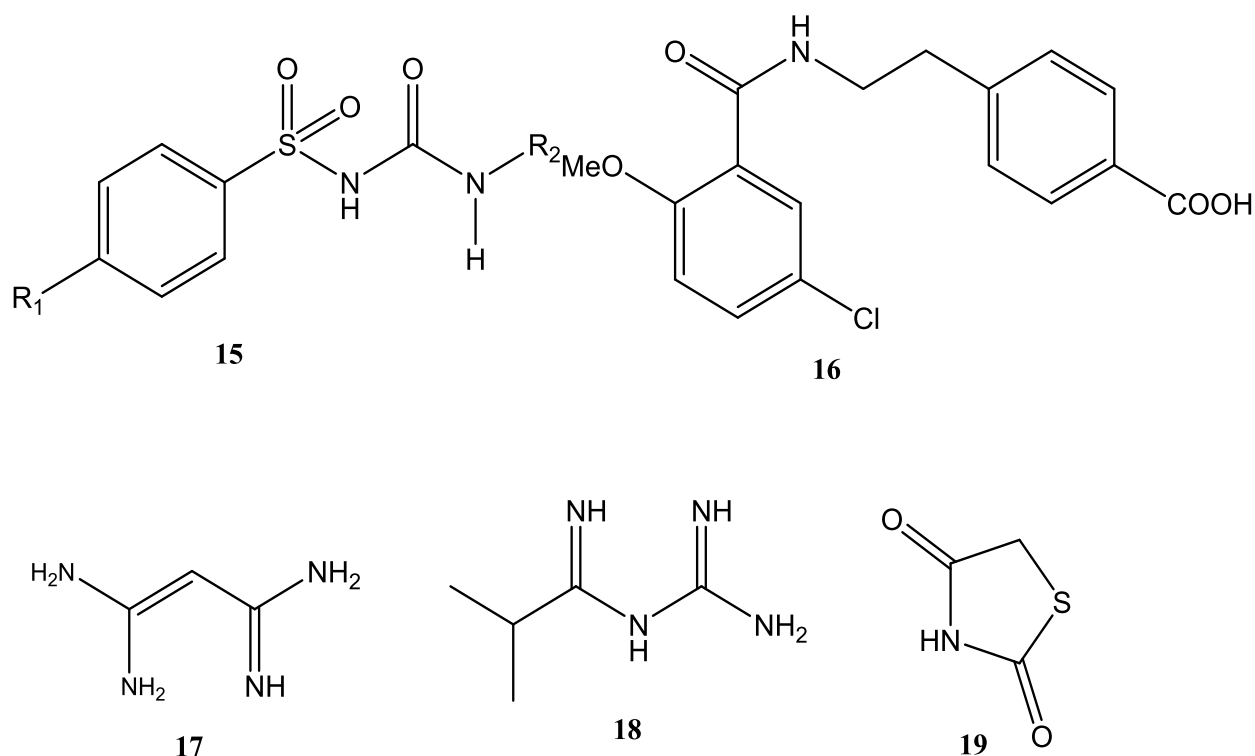
#### 2.1 Diabetes Mellitus

Mammals must digest complex carbohydrates, starch, for provision of energy the body needs (Quezada *et al.*, 2007). During starch metabolism, complex pathophysiological conditions can interfere with starch hydrolysis, leading to diabetes mellitus. This condition is associated with hyperglycemia, or elevated blood glucose levels, is a complication of this chronic illness (Gerich *et al.*, 2006; Zeyana *et al.*, 2013). In 2016, over 360 million people worldwide were diagnosed with diabetes due to unhealthy lifestyles, obesity, and ageing (WHO, 2016). Prolonged hyperglycemia can cause organ failure including kidneys, heart, blood vessels, nerves, eyes, and kidneys (Gerich, 2006).

In 2014, 8.5% of people who were 18 years of age or older had diabetes. In 2019, diabetes-related mortality accounted for 1.5 million deaths, with 48% of these deaths occurring in those under the age of 70. Diabetes was responsible for 20% of deaths from cardiovascular disease and 460,000 deaths from renal disease, according to the (WHO, 2016). Between 2000 and 2019, there was a 22% global drop in non-communicable diseases, however diabetes raised age-standardized death rates by 3%, leading to a 13% increase in mortality in lower-middle income nations (WHO, 2020).

Type 1, type 2, and gestational diabetes are three types of diabetes. Type 1 causes insufficient insulin production due to pancreatic  $\beta$ -cell destruction occasioned by genetic susceptibility (Eckel *et al.*, 2011), while type 2 results from malfunctions and insulin resistance (Lin and Sun, 2010). Gestational diabetes occurs during pregnancy due to hormonal changes (Eckel *et al.*, 2011). Treatments include insulin from outside sources to manage type 1 diabetes whereas healthy diet, exercise, and administration hypoglycemic agents are applied for type 2 (Krentz and Bailey, 2005). To date, finding a safe, efficacious drug to combat all conditions simultaneously without side effects is a challenge to be addressed.

A part from insulin administration other oral anti-hyperglycemia therapeutic options available include sulphonylureas (**15**) and meglitinides (**16**) drugs which stimulate pancreatic insulin secretion. Biguanides (**17**) and metformin (**18**) medication reduce hepatic glucose production and thiazolidinediones (**19**) improve insulin action. While Miglitol (**6**) and acarbose (**1**) delay digestion and absorption of glucose (Luna and Feinglos, 2001). The latter option has not received much attention due to several options and its mild effectiveness, although it may prove to be safe in terms of preventive measures.



## 2.2 Carbohydrate metabolism

Carbohydrates is key dietary component for energy provision for animals is broken down by enzymes at the small intestine. (Haque and Tan, 2017). Enzymatic hydrolysis of glycosidic bonds in disaccharides and polysaccharides is crucial for metabolism to simple sugars. Management of the rate of carbohydrate digestion to avoid too much or less glucose in the bloodstream has thus been regarded as an option in management of diabetes mellitus (Ünal *et al*, 2012).

### 2.2.1 Glycoside Hydrolases

Glycoside hydrolases are enzymes that break down the glycosidic bond between carbohydrates or carbohydrate and non-carbohydrate components. Glycoside hydrolases have been categorized using a similarity in sequence, which has resulted in the identification of more than a thousand distinct enzyme families that are selective to various substrates (Henrissat and Davies, 1995).

#### 2.2.2 $\alpha$ -Amylase

$\alpha$ -amylases are found in saliva and pancreatic secretions. They are responsible for breaking down the internal  $\alpha$ -1,4-glycosidic linkages in starch (Fig. 1(a)). Amylases convert starch into low-molecular-weight products. These low-molecular products are in form of maltose, short oligosaccharides, and limit dextrins, they are branch points of amylopectin (Rajagopalan and Krishnan, 2008). The main function of  $\alpha$ -Amylase is to reduce workload in the small intestines and

prevent oligosaccharide accumulation (Horton and Tomasik, 2015; Wickham *et al*, 2015). However, high amylase activities may lead to excess oligosaccharides, requiring control to avoid excess glucose generation.

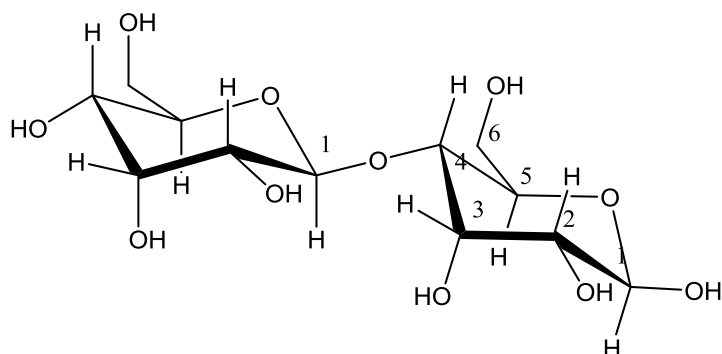


Fig.1 (a):  $\alpha$ -1, 4-glycosidic linkage

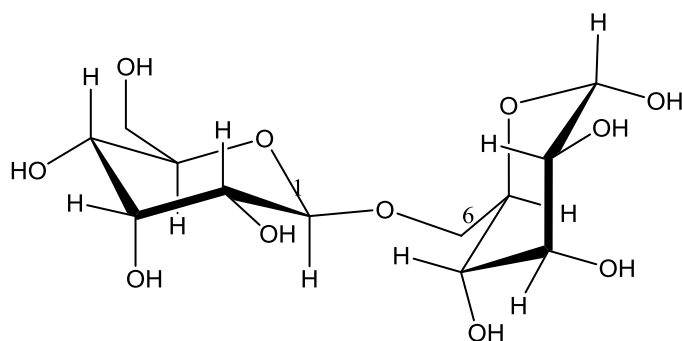


Fig 1(b)  $\alpha$ -1, 6-glycosidic

### Figure 1: Structure of Glycoside

#### 2.2.3 Brush-border hydrolases

The final stage of digestion of dietary carbohydrates is performed by brush border hydrolase enzymes on the face of small intestinal enterocytes (Vaughan, 2002). Brush border enzymes, tethered to the plasma membrane. They vary in density and distribution depending on animals age and small intestine segments or may vary due to food type the animals was fed on. After digestion monosaccharides are absorbed into epithelial cells (Ross *et al.*, 2016).

The brush border membrane contains two enzymes, sucrase-isomaltase and maltase-glucoamylase, (together are referred to as  $\alpha$ -glucosidase). They perform the final stages of carbohydrate digestion (Simpson *et al*, 2015). Maltase-glyco-amylase has multiple active sites, converting maltose or maltotriose residues into  $\alpha$ -D-glucose and releasing  $\beta$ -D-glucose. Similarly, sucrase-isomaltase has

two active sites with distinct specificities, catalyzing the breakdown of sucrose into fructose and glucose (Mhlongo, 2014). Controlling these enzymes may require general  $\alpha$ -glucosidases inhibitors to prevent excessive glucose release into the bloodstream.

#### 2.2.4 Enzyme Inhibition Modes

Enzyme structure studies can aid in understanding inhibitory mechanisms, although structural data may be unavailable, consequently limiting determination of enzyme-inhibitor binding affinity. Nevertheless, kinetic investigation using Michaelis-Menten (equation 1), has often been used to establish the inhibition modes of enzymes by different inhibitors (Kumar, 1985),

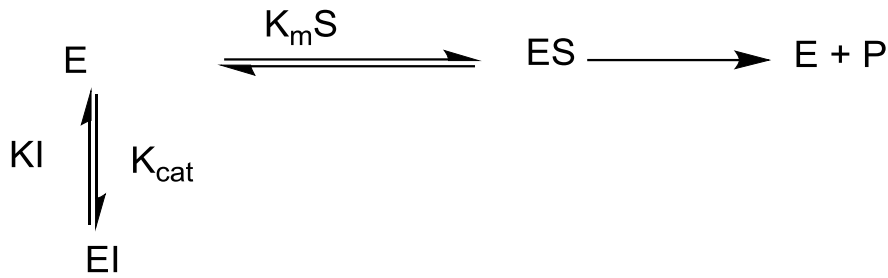
$$V_0 = \frac{V_{\max} [S]}{K_m + [S]} \quad \text{Equation 1}$$

where enzymatic activities is described in terms of; Michaelis constant,  $K_m$  and maximum velocity,  $V_{\max}$  (Michaelis, 1913). Equation 1 can be modified to the double reciprocal rate equation known as Lineweaver – Burk equation (equation 2) in order to obtain linear plots for enzyme inhibition reactions (Tong and Jingyan, 1999). From this equation,  $K_m$  and  $V_{\max}$  can be obtained.

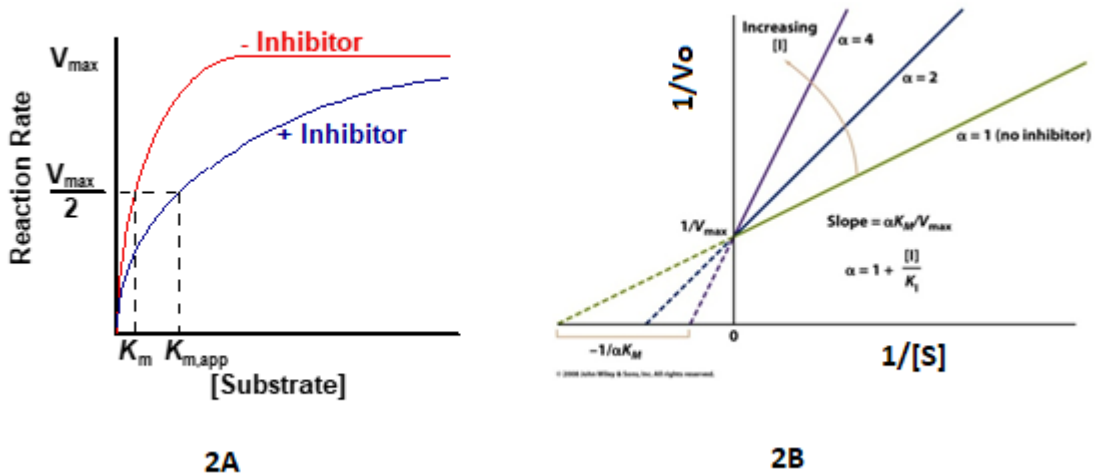
$$\frac{1}{v} = \frac{K_m}{V_{\max} \left(1 + \frac{[I]}{K_i}\right)} \cdot \frac{1}{[S]} + \frac{1}{V_{\max}} \quad \text{or} \quad \frac{1}{v} = \frac{K'_m}{V_{\max}} \cdot \frac{1}{[S]} + \frac{1}{V_{\max}} \quad \text{Equation 2}$$

An inhibitor may bind to an enzyme's active site (isosterically) and then fail to release the enzyme, preventing the formation of an enzyme-substrate complex and irreversibly inhibiting the conversion of the substrate into the product (Whitely, 2000). Conversely, inhibitors may also isosterically bind with an enzyme or allosterically bind but release the enzyme (reversible inhibition) (Punekar, 2018).

Reversible inhibitions can be classified as competitive, noncompetitive, or uncompetitive (Brandt *et al.*, 1987). In the case of competitive inhibition, the inhibitor binds to the enzyme's active site to form the enzyme-inhibitor complex (scheme 1), which reduces the enzyme's affinity for the substrate and is indicated by an increase in the Michaelis constant ( $V_{\max}$ ,  $K_m$  apparent) (Fig 2(a)). When the substrate concentration increases, the excess substrate overcomes the inhibitor and the reaction proceeds to acquire the maximum rate of reaction ( $V_{\max}$ ). Therefore, in competitive inhibition,  $V_{\max}$  is the same as that of the control while the turn over number ( $K_{\text{cat}}$ ) remains constant (Cheng and Prusoff, 1973) as illustrated in figure 2(b).



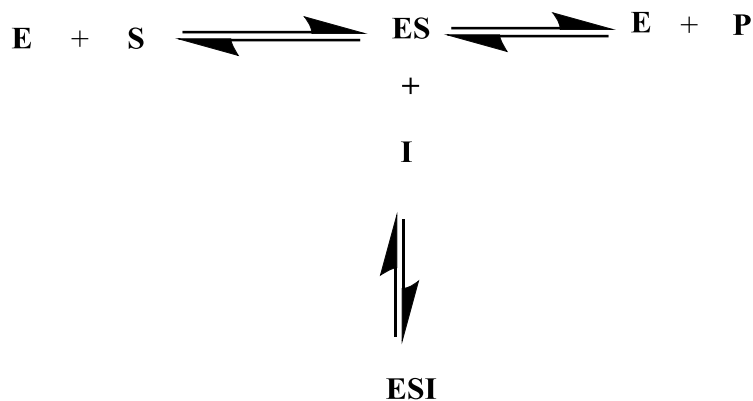
Scheme 1



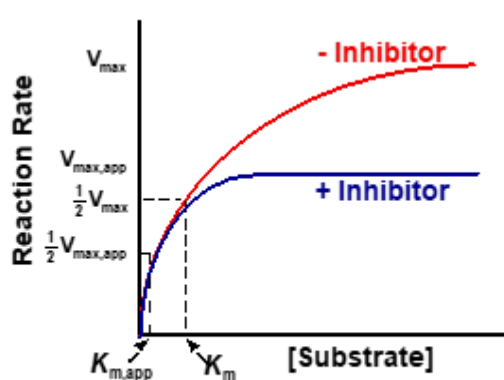
**Figure 2: Competitive Inhibition curves: 2A) Michaelis-Menten plot reaction velocity vs substrate concentrations; 2B) Lineweaver-Burk plot of 1/initial velocity vs 1/[Substrate].**

Inhibitor binds with both the free enzyme and the enzyme-substrate complex, resulting in noncompetitive inhibition. A tertiary complex (ESI) is created when the inhibitor attaches to the enzyme-substrate complex as well as when it binds to the free enzyme. This prevents the enzyme from converting the substrate into the product (scheme 2). The increase in substrate concentration does not overcome inhibition. The  $V_{\text{max}}$  decreases due to decrease in the number of functional enzymes since there is decrease in the number of active sites whereas  $K_m$  remains constant (Fig 3(a)). The turn over number is lowered because as the enzyme binds with the inhibitor, the inhibitor changes the shape of the enzyme making it non complementary to the substrate (Fig 3(b)) (Brandt *et al*, 1987).

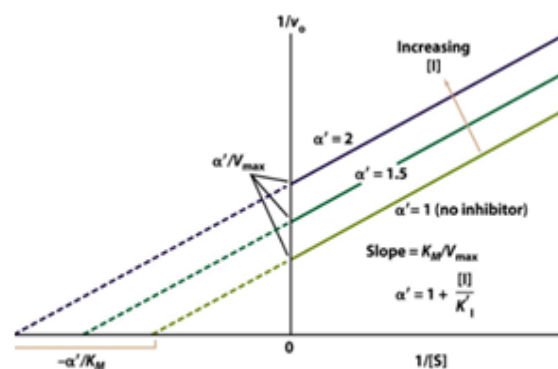




Scheme 3



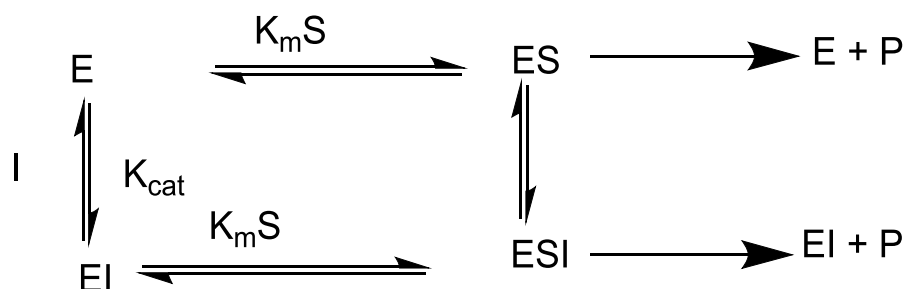
(4A)



(4B)

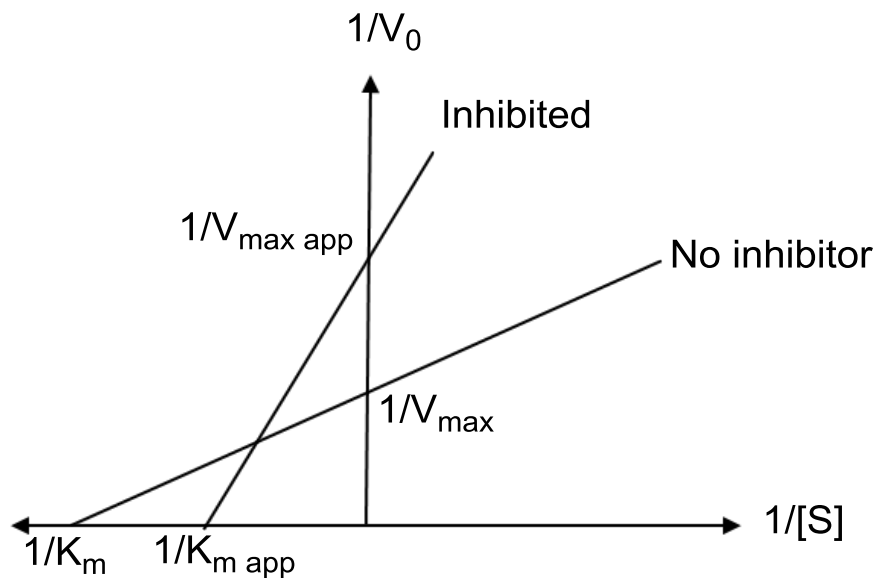
**Figure 4: Uncompetitive Inhibition curves: (4A) Michaelis-Menten plots of reaction velocity vs substrate concentration; (4B) Lineweaver-Burk plot of reciprocal of initial velocities vs reciprocal of substrate concentration.**

Enzyme-inhibitor (EI) and enzyme-substrate-inhibitor (ESI) complexes are formed when an inhibitor binds to both enzyme (E) and enzyme-substrate (ES) complexes. These complexes dissociate in distinct ways because the EI and ESI constants have different dissociation constants. Both  $V_{\max}$  and  $K_m$  are lowered (Figure 4).



Scheme 4





**Figure 5: Lineweaver-Burk plot for Mixed Inhibition**

### 2.3 Starch- Digesting Enzymes inhibitors

Enzyme-inhibitor (EI) and enzyme-substrate-inhibitor (ESI) complexes form when an inhibitor binds to both, obstructing the small intestine's process of converting disaccharide to monosaccharide. Such inhibitors thus slow glucose entry and can be used for management of postprandial hyperglycemia (Shimabukuro *et al*, 2006).

#### 2.3.1 Acarbose and Miglitol

Acarbose (**1**) is an anti-hyperglycemic drug used to manage type-2 diabetes mellitus which until recently was believed to be effective, and safe and for treatment of type 2 diabetes (Zhang, 2013). Nevertheless, it has been noted as ineffective in patients on a relatively high carbohydrate diet (Zhu *et al*, 2011). Moreover, side effects associated with poor absorption leading to abdominal discomfort from the gastrointestinal tract have been reported on patients who use the drug (Putter *et al*, 19).

Conversely, miglitol (**6**) is an oral medication that lowers blood sugar levels by preventing the body from converting complex carbohydrates into glucose (Guntram, 2013) The amount of non-monosaccharide carbohydrates in an individual's diet determines how much miglitol (**6**) and other structurally related iminosugars inhibit  $\alpha$ -amylase enzymes by slowing the digestion of carbs and decreasing postprandial hyperglycemia (Rebecca, 2013). Unlike acarbose (**1**), miglitol (**6**) is well

absorbed with a 100% systemic bioavailability. It is not metabolized and is eliminated by the kidneys in its unaltered form because of its strong structural similarity to glucose (Joubert *et al.*, 1990). As mentioned earlier, these two medications along with other plant-based inhibitors are used for treatment of postprandial hyperglycemia, as they have been demonstrated to have a strong inhibition activity against  $\alpha$ -glucosidase and a little inhibitory impact on the activity of  $\alpha$ -amylase (Afonne, 2002).

In recent times, herbal medicines have gain relevance in the management of diabetes mellitus since they are less costly and have less adverse effects than synthetic anti-hyperglycemic medications (Mukherjee *et al.*, 2006). However, the specificity of the inhibition modes remains unknown since many herbal plants that have been cited to act against diabetes mellitus and not much has been documented such inhibitors.

#### **2.4 Compounds from *Zanthoxylum* Plants and their biological activities**

There have been reports of notable anti-hyperglycemic effects from a number of plants in the genus *Zanthoxylum* (Aloke *et al.*, 2012). For example, root bark, stem bark and leaves of *Z. zanthoxyloides* is used medicinally for treating diabetes mellitus (Wagman, 2001).

In addition, stem bark of *Z. armatum* stem bark has been reported to have anti-hyperglycemic potential by lowering blood glucose levels in rats with diabetes caused by streptozotocin. This effect was related to the presence of alkaloids and phenolic compounds (Karki *et al.*, 2014). When *Z. chalybeum* stem bark extract was given to streptozotocin-induced diabetic rats, blood glucose levels significantly decreased. This study finding partially supported the plant's traditional use in the treatment of diabetes (Kimani, 2015). The genus *Zanthoxylum* is a rich source of alkaloids, lignans, steroids, coumarins, benzenoids, terpenes, coumarines, flavonoids, fatty acids and amides (Fang *et al.*, 2014). Alkaloids and coumarins are the primary elements of the genus, according to a review by Negi *et al.* (2011).

A number of species in the genus *Zanthoxylum* have been identified through phytochemical characterization as containing alkaloids, including chelerythrine (**20**), skimmianine (**21**) and arnottianamide (**22**), nitidine (**23**), N-methyl flindersine (**24**) (Krane *et al.*, 1984), zanthomurolanines (**25**) and dihydrochelerythrine (**26**) (Matu, 2011), rutaecarpine (**27**) and lignans (**28**) (Sheen *et al.*, 1994). *Z. nitidine* produced nitidine (**23**), which was isolated (Gakunju *et al.*, 1995). Seven compounds were isolated from *Zanthoxylum holtzianum* during a phytochemical investigation for antimicrobial principles. Of these, three were benzophenanthridine alkaloids [dihydrochelerythrine (**26**), 6-acetyldihydrochelerythrine (**27**) and 8-oxochelerythrine

(**28**)], along with a phenol canthine-6-one (**29**) and fatty acids, flavanones, and amides. Strong anti-diabetic effects were reported for flavonoids, chelerythrine (**20**), berberine (**30**), and phenol canthine-6-one (**29**) that were extracted from the methanol extract of *Z. zanthoxyloides* root bark (Odebisi and Sofowora, 1979; Tsuchiya, 1996), supporting the activity described on *Z. chalybeum*. Similarly, skimmianine (**21**) isolated from seed extract of *Z. chalybeum* showed both anti-hyperglycemic and anti-lipidemic properties (Olila *et al*, 2001). Muganga *et al.*, (2014) in a bio-guided methanol root bark extract of *Z. chalybeum* fractionation identified nitidine (**23**) as the main antimalarial compound alongside chelerythrine (**20**) and methyl canadine (**31**). Adia *et al*, (2016) reported the isolation of fagaramide (**32**) from Uganda's *Z. chalybeum* stem bark that was associated with management of diabetes. Anza *et al*, (2014) reported that *Z. chalybeum* root extract (CH<sub>2</sub>Cl<sub>2</sub>/CH<sub>3</sub>OH 1:1) underwent phytochemical screening, which identified the presence of anthraquinones, alkaloids, flavonoids, terpenoids, and tannins, but no pure molecule was ever extracted. A coniferyl alcohol derivative, 2,3-epoxy-6,7-methylenedioxy coniferyl (**10**) and a known alkaloid, dihydrochelerythrine (**26**), which had been previously isolated from *Z. nitidum*, were obtained from the extract by column chromatographic separation (Yang *et al*, 2014; Krane *et al*, 1984).

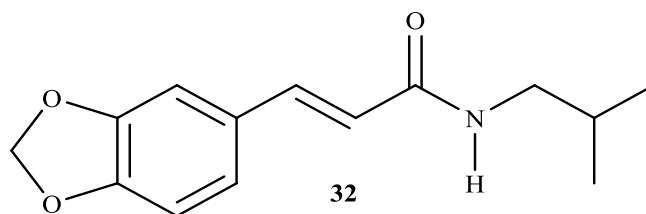
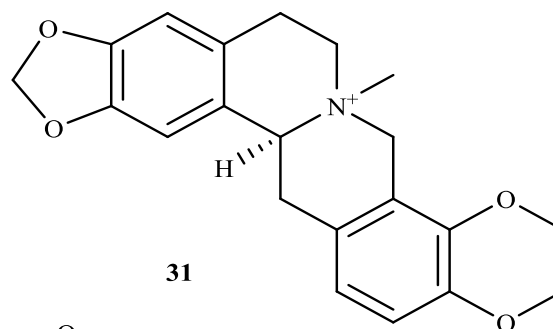
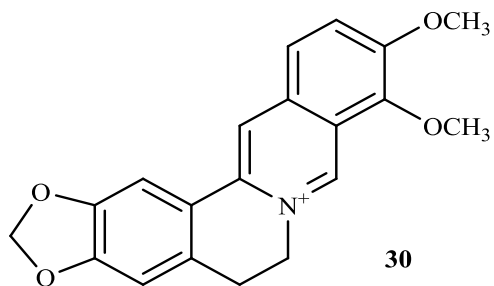
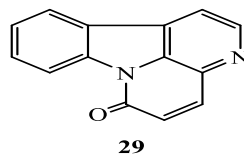
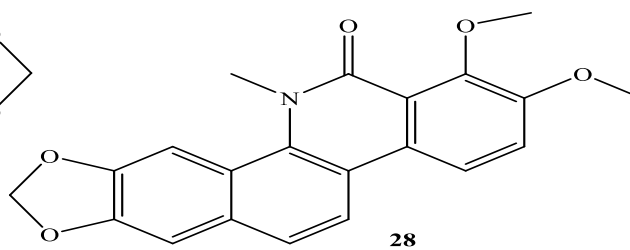
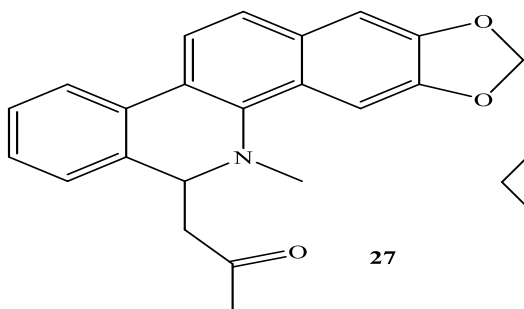
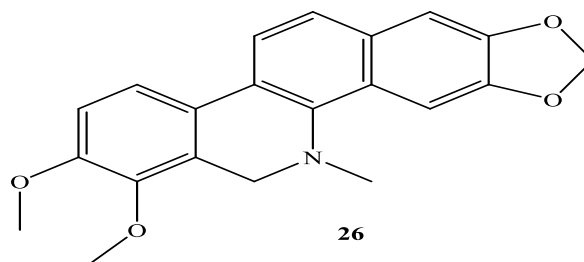
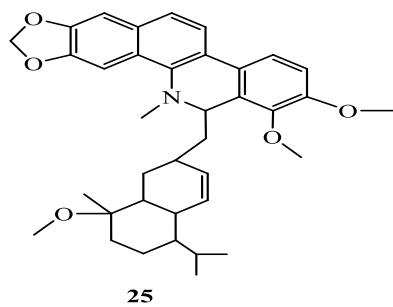
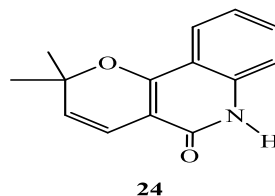
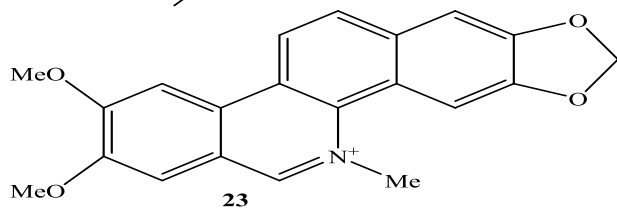
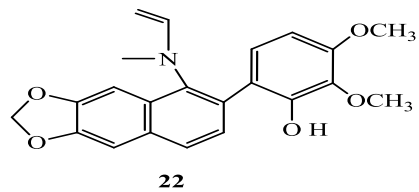
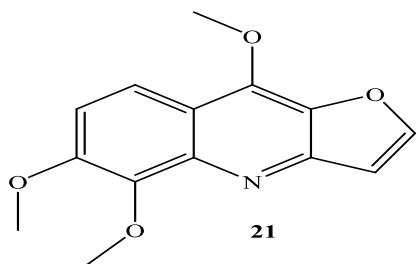
The aqueous root extract of *Z. chalybeum* showed good anti-hyperglycemic and anti-hyperlipidemic capabilities in an inquiry on the impact of the plant on diabetes-induced cardiac dysfunction in alloxan-induced type-1 diabetic rat. The study concluded that the plant possesses wide therapeutic benefits including anti-diabetic and anti-oxidant activities (Agwaya *et al*, 2016). A single phenolic 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) and two benzophenanthridines, 6-hydroxy-N-methyl decarine (**8**) and norchelerythrine (**9**) were isolated from the stem bark of *Z. chalybeum* and shown to exhibit antidiabetic activity by inhibiting carbohydrate hydrolytic activities (Wasiali, 2019). Compounds **8**, **9** and **10** had comparable inhibitory activity ( $p > 0.05$ ) against both  $\alpha$ -amylase and  $\alpha$ -glucosidase (IC<sub>50</sub> = 64.54, 58.91 and 54.81; and 66.77, 62.53 and 58.03  $\mu$ M) in comparison to the positive control acarbose (IC<sub>50</sub> = 54.66 and 51.54  $\mu$ M) (Wasiali, 2019).

In contrast, the triterpenoids that were extracted from the same extract and identified as lupeol (**11**), 3 $\alpha$ ,20-dihydroxy-28-lupanoic acid (**12**), 20-hydroxy-3-oxo-28-lupanoic acid (**13**) and 3 $\alpha$ ,20,28-trihydroxylupane (**14**) had minimal inhibitory effects on the two enzymes when compared to the same control (Wasiali, 2019). The work of Wasiali 2019, thus attributed the potential of *Z. chalybeum* to benzophenanthridine and possible phenolics, however for comprehensive structure

activity validation of active compounds the number of compounds used in the study could not provide a clear picture.

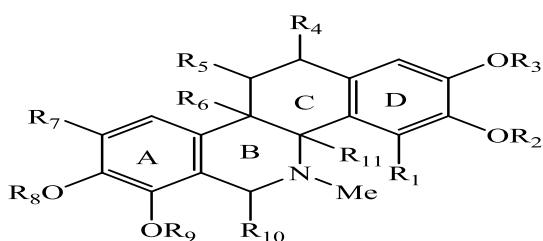
With extremely similar morphological characteristics to *Z. chalybeum*, and *Z. gillettii* is an indigenous deciduous tree that can reach heights of 10 to 35 meters. Even though it is found in Africa, it is also widely spread in nations like Zimbabwe, Zambia, Kenya, Sudan, Angola, and Malawi. Commonly reported traditional applications include treatment of stomachache by chewing the bark; against urinogenital infections, rheumatism and other pain related ailments (Kokwaro, 2009). In addition to its claimed anti-leukemic (Nakanishi et al., 1998), antibacterial, anti-inflammatory, analgesic, and anti-HIV effects, the plant was also found to possess strong anticancer properties (Tan et al., 1991). In identification of secondary metabolites, *Z. gillettii* has so far been able to establish benzophenanthridine alkaloids such as nitidine (**23**), dihydrochelerythrine (**26**) and chelerythrine (**20**) as well as furoquinoline alkaloids such as skimmianine (**21**), cinnamic acid amide, and fagaramide (**32**).

*Z. gillettii* serves as a better reservoir for alkaloids and phenolics some of which could be active compounds. However, these compounds have not been evaluated for anti-hyperglycemic potential. With the exception of pentacyclic triterpene acetate, 3 $\beta$ -acetoxy-16 $\beta$ -hydroxybetulinic acid (**7**), which was described as having modest  $\alpha$ -glucosidase inhibitory activities. (Mbaze et al., 2007).

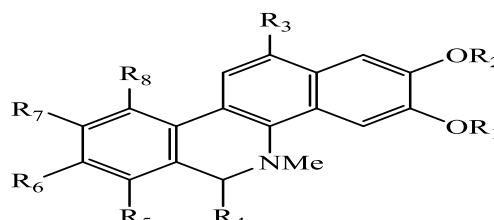


### 2.4.1 Structure Activity Relationship of Benzophenanthridine Alkaloids

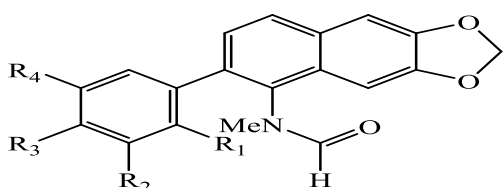
Benzophenanthridine alkaloids belong to the benzyl isoquinoline alkaloid family and are primarily found in the Rutaceae and Papaveraceae families (Krane *et al.*, 1984). More than 100 have been documented using various separation and identification procedures. They are divided into three types namely quaternary benzo[c]phenanthridine (QBP) type, dihydrobenzo[c]phenanthridine (DBP) type, and hexahydrobenzo[c]phenanthridine (HBP) type (Krane *et al.*, 1984). HBP type consists of two aromatic rings, two cycloalkanes and *N*-Me group: DBP type consist of three aromatic rings, one cycloalkane and *N*-Me group with an open ring B as *N*-acetyl or methoxy group. Furthermore, Dihydrobenzo[c]phenanthridine alkaloids can be categorized further depending on the linkage between C-7 and C-8 and a polymerized type at C-8. QBP type takes the shape of a quaternary ammonium salt. (Figure 6).



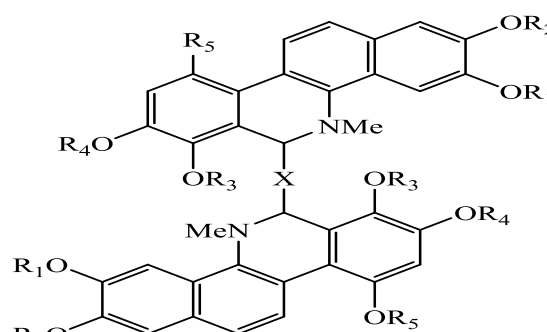
Hexahydrobenzo[c]phenanthridine type



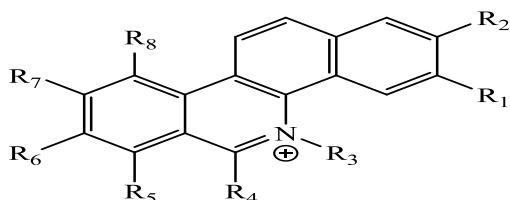
Dihydrobenzo[c]phenanthridine type



Dihydrobenzo[c]phenanthridine type I



Dihydrobenzo[c]phenanthridine type II



Quaternary benzo[c]phenanthridine

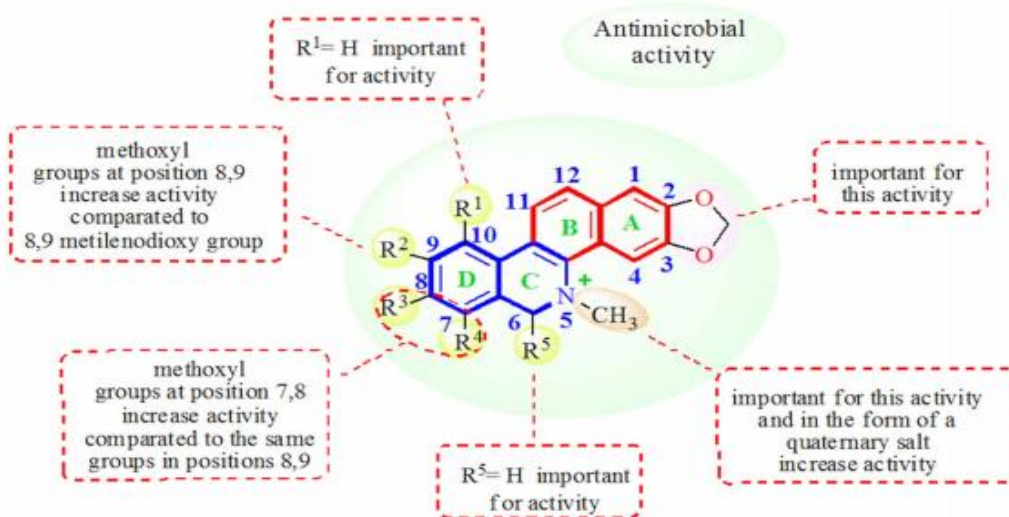
**Figure 6: The structures of different types of Benzophenanthridine alkaloids**

The class DBP includes decarine (8), norchelerythrine (9), dihydrochelerythrine (26), avicine (33), dihydroavicine (34) among others whereas the QBP include nitidine (23), faragonine (35) and quaternary benzophenanthridine include faragamide (32) which are ubiquitous in the *Zanthoxylum* spp.

Benzophenanthridine alkaloids isolated from *Zanthoxylum* spp have been shown with as anti-inflammatory, anti-tumor, and anti-microbial qualities (Chen *et al*, 2005) and it has been demonstrated through modeling studies that the majority of them are protein kinase C inhibitors that preserve erythrocyte deformability (Yang *et al*, 2008) Some have been suggested to produces more reactive oxygen species, which can then exhibit bacteriolytic effects (Tane *et al.*, 2005). Both antibacterial and antifungal activities have been predicted to be dependent on a B/C ring fusion and the existence of an N-methyl group (Ma *et al*, 1999).

Higher antimicrobial activities has been reported for chelerythrine (20) and avicine (33), dihydrochelerythrine (26) and dihydroavicine (34). The corresponding hydrogenated alkaloids, suggest that the kind and location of substitution on ring D affects this kind of activity (Tavares *et al*, 2014). Comparably, it has been observed that molecules with a methylenedioxy group, like avicine (34) are more active than those having two methoxyl groups at positions C-8 and C-9, like nitidine (23) (Tavares *et al*, 2014).

Furthermore, the methoxyl groups' position is a relevant structural feature for antimicrobial activities of these alkaloids, since ortho coupled methoxyl groups at C-7 and C-8 as observed in chelerythrine (20) tend to enhances the antimicrobial activities better than dimethoxyl groups at C-8 and C-9 as in nitidine (23) (Tavares *et al*, 2014). The study by Tavares *et al*, (2014) (Figure 7) concluded the following moieties as important pharmacophores in benzophenanthridine alkaloids.



**Figure 7: The structure-activity connection for the benzophenanthridine alkaloids' antibacterial activity (Tavares *et al.*, 2014)**

There have been reports that quaternary benzo[*c*]phenanthridine, sanguinarine (35) and nitidine (23), block HIV-1 reverse transcriptase by thus considering benzophenanthridine alkaloids as potential anti-HIV compounds (Tan *et al.*, 1991). Even non QBP molecules such as decarine (36) have been proved to have inhibitory potential against HIV-1 replication in H9 lymphocytes (Cheng *et al.*, 2005) thus further validating activities of benzophenanthridine alkaloids (Cheng *et al.*, 2005).

Based on the structure-activity relationship (SAR), which anticipated a C=N between positions 7 and 8 as a significant functionality, seven benzophenanthridine alkaloids with varying degrees of analgesic and anti-inflammatory properties were extracted from *Zanthoxylum nitidum* (Roxb.) DC (Hu *et al.*, 2006). Conversely, replacement on C-8 reduced the anti-inflammatory activity, but the 9-OH and 10-OMe groups had greater anti-inflammatory activity (Hu *et al.*, 2006).

According to research by Jursky and Balioya (2011), the two most potent compounds, chelerythrine (20) and sanguinarine (37), were found to selectively inhibit the glycine transport proteins GlyT 1 rather than GlyT, hence suppressing pain. Studies on anticancer agents similarly predicted C=N bond of benzophenanthridine alkaloids following a possible reaction between



C=N group on the B ring and -SH nucleophilic sites on the protein residues consequently exhibiting actions against proliferation and angiogenesis (Wolf and Knipling, 1993).

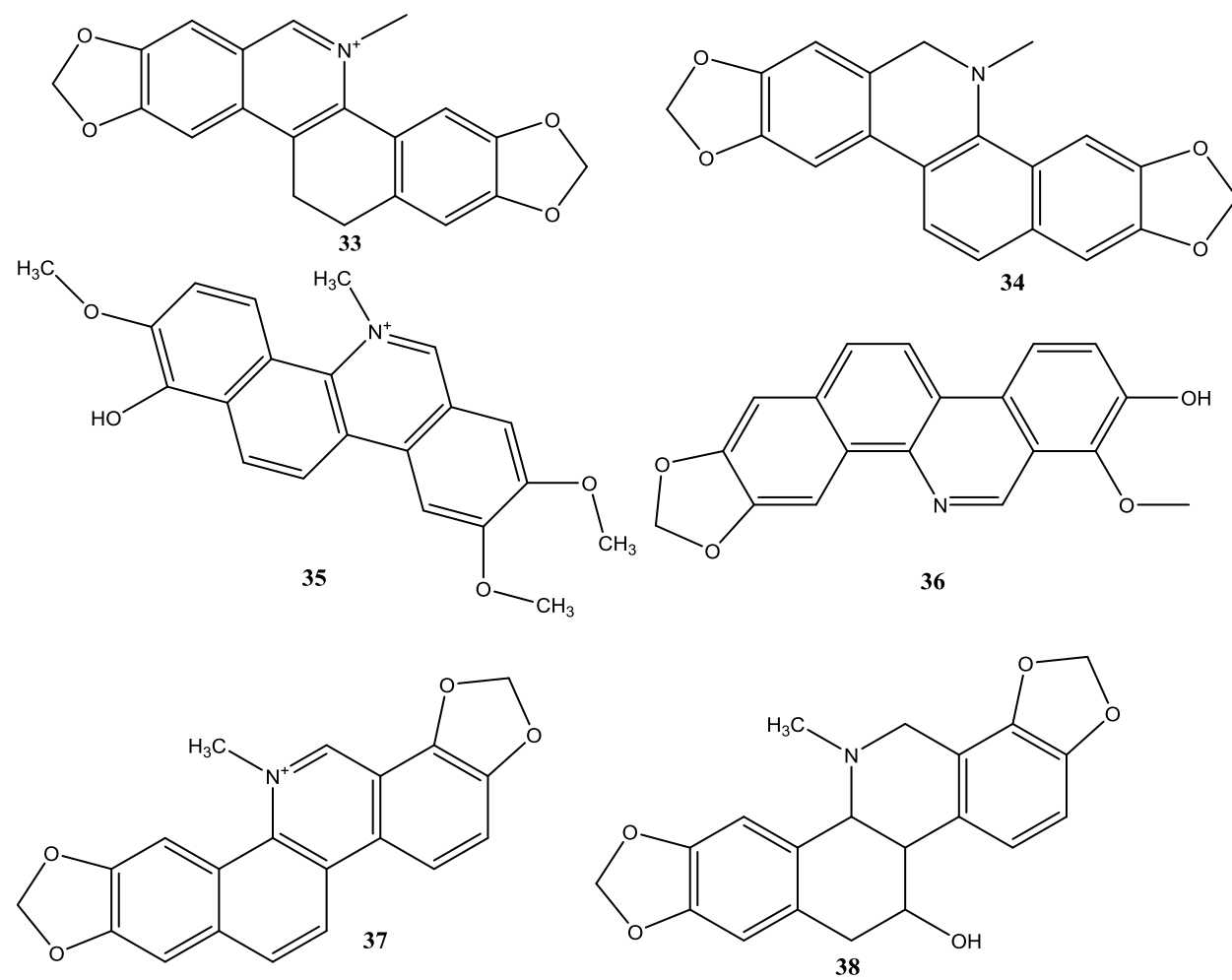
Furthermore, the chemicals have the potential to impact DNA Topoisomerase I and II enzymatic activities through insertion into DNA molecules, hence impeding the rapid growth of tumor cells (Philchenkov et al., 2008). These mechanisms were reported based on study using chelidonine (**38**), sanguinarine (**37**) and chelerythrine (**20**) which are known for anti-microtubular effects (Wolf and Knipling, 1993). However, chelidonine (**38**) relies more on spatial steric hindrance brought on by the non-planar structures of the B and C rings, whereas sanguinarine (**37**) and chelerythrine (**20**) have similarly analogous mechanisms of action that depend on the C=N group on the B ring (Wolf and Knipling, 1993).

Structure-activity relationships on a set of 19 alkaloid molecules including benzophenanthridine alkaloids using ischemia-hypoxia H9C2 cells discovered that hexahydrobenzo[c]phenanthridine and dihydrobenzo[c]phenanthridine showed better activity more than proberberine and aporphine (Yang *et al.*, 2014). The activities were noted to be reducing upon acetylation 6-OH or substitution of 8-H with OH, methyl group hydroxylation at positions C-2, 3, and 11, as well as the lack of N-Me (Yang *et al.*, 2014).

Benzophenanthridine alkaloids have also been noted for very good antioxidant activities against DPPH and ABTS better than known positive antioxidants. For instance, Proberberine is more potent clearance free radical than the alkaloids due to the phenolic hydroxyl group's location on the HBP skeleton (Han *et al.*, 2016). Benzophenanthridine alkaloids are consequently acknowledged to exhibit promising possibilities for the creation of novel pharmacological substances.

Nevertheless, because only a small number of benzophenanthridine alkaloids have been isolated, research into these compounds is still ongoing. New analysis methods must be developed in order

to find more compounds, and their activities must be carefully screened in order to find novel treatments for a range of illnesses.



## CHAPTER THREE

### MATERIALS AND METHODS

#### 3.1 Materials

##### 3.1.1 General Instrumentation

The stem bark was ground into a powder using an electric pulverizer (Kika, M20), and the powdered materials samples were weighed using a top-loading analytical balance (Shimadzu, UX). Samples were concentrated using a rotary evaporator (EYELA, N-1000) that evaporated the liquids at low pressure. Gallen Kamp was used to record melting points with uncorrected capillary tubes. Kieselgel F<sub>254</sub> aluminum silica gel plates were utilized in thin layer chromatography, and column chromatography was performed over silica gel (Merck 60G, 70-230 mesh) and sephadex LH-20 (Kobian, Kenya and Sigma Chemical Company, St. Louis, USA).

The TLC plates were viewed under UV lamp then heated using heat gun for two minutes. Detection and visualization were done by UV-Visible light spectrophotometer (molecular device Sunnyvale, USA) at the instrument room, Department of Chemistry, Maseno University. Bruker or Varian-Mercury spectrometers were used to record the 1 D (<sup>1</sup>H-NMR (500 MHz) and <sup>13</sup>C-NMR (125 MHz)) and 2 D (COSY, NOESY, HSQC, and HMBC) using DMSO, CDCl<sub>3</sub>, CD<sub>2</sub>Cl<sub>2</sub>, or CD<sub>3</sub>OD as solvents with tetramethylsilane (TMS) serving as the internal standard. Chemical shifts were determined in ppm in  $\delta$  values and coupling constants (*J*) in Hz compared to deuterated solvents and TMS. ESIMS spectra were obtained using a Finnigan MAT mass spectrometer with a direct input at 70 eV on SSq 710.

Using a SpectraMax GeminiXS Spectrofluorometer (Molecular Devices, Sunnyvale, CA), the  $\alpha$ -amylase enzymatic activity (fluorescence) was measured.  $\alpha$ -glucosidase enzyme activity (absorbance) was determined using a SpectraMax 190 Spectrophotometer (Molecular Devices, Sunnyvale, CA).

##### 3.1.2 Chemicals and Reagents

The organic solvents and reagents used in the study were obtained from Sigma-Aldrich Chemical Company, St. Louis, USA, and Kobian, Kenya. These included ammonia solution, n-hexane, dichloromethane, ethyl acetate, glacial acetic acid, p-anisaldehyde, and Dragendorff's crystals. Organic solvents were distilled, Dimethylsulfoxide (DMSO) was analar grade from Kobian, Kenya, and  $\alpha$ -glucosidase, bovine serum, p-nitrophenyl- $\alpha$ , D-glucopyranoside, and acarbose were purchased from Sigma-Aldrich Chemical Company.

### 3.1.3 Chromatographic conditions

TLC was performed on pre-coated aluminum silica gel plates (Merck 60, F254). Chromatographic zones on TLC plates were spotted by spraying *p*-anisaldehyde-sulphuric acid, iodine vapor, or ammonia vapor, and burned at 100 °C. For alkaloid tests, Dragendorff's reagent was sprayed on TLC plates. The *p*-anisaldehyde-sulphuric acid visualizing stain was prepared from a mixture of 0.5 ml of *p*-anisaldehyde, 85 ml of ethanol, 5 ml concentrated sulphuric acid, and 10 ml glacial acetic acid (Krishnaswamy, 2003). Column chromatography was carried out using silica gel (Merck 60G, 70-230 mesh) in glass columns measuring 3.0 × 60 cm and 2.0 × 50 cm. Solvents including n-hexane, dichloromethane (DCM), ethyl acetate (EtOAc), and 95% methanol were used in the study.

### 3.1.4 Plant Materials Collection and Preparation

Plants were identified from its natural habitat by assistance of a plant taxonomist and the samples collected were conserved at University Herbarium after authentication. The stem bark of *Zanthoxylum chalybeum* was sampled from Homa Hills, (0°, 23'S, 34°30'E) and *Zanthoxylum gillettii* stem bark was gathered in Kakamega Forest, (0°17'30''N, 34°51'22''E) in January, 2019. Samples were identified and authenticated by a taxonomist at Maseno University, Department of Botany where a voucher specimen (KYC/MSN/2019) for *Z. chalybeum* sample and (KYG/MSN/2019) for *Z. gillettii* were deposited at Maseno University herbarium. The plants materials were sliced into small pieces then allowed to dry by air under the shade. ZC stem bark dried after 21 days and ZG after 25 days. The dried materials were then pulverized differently into fine powder using an electric grinder at Maseno University Research Laboratory.

## 3.2 Methods

### 3.2.1 Extraction and isolation of compounds from *Z. chalybeum*

Using an electric pulverizer, the *Z. chalybeum* stem bark was ground into a fine powder after being sliced into small pieces and allowed to dry by air for 21 days in a shade. The 0.8 kg of powdered stem bark was extracted using 4 × 1.5 L of 95% aqueous methanol, and filtered through Whatman No. 1 using a Buchner filter funnel. Using a rotary evaporator, the consecutive filtrates were combined and concentrated in vacuum chamber to produce (45 g) of 5.625% crude extract at a temperature of 45°C.

Total alkaloid extraction: total alkaloid was isolated 28 g of the crude extracts were suspended in 200 mL of 0.1 M H<sub>2</sub>SO<sub>4</sub> in ethyl acetate and agitated for 30 minutes. The extracts were then

combined and the non-polar compounds were extracted three times using 300 mL of n-hexane in a separating funnel.

A 0.1 M ammonia hydroxide solution was used to neutralize the acidic polar fractions. 200 mL of dichloromethane ( $\text{CH}_2\text{Cl}_2$ ) was then used to extract the material further until a negative Dragendorff's test result was obtained. Anhydrous sodium sulphate was used to dry the organic fraction ( $\text{CH}_2\text{Cl}_2$  extracts), and then further concentration in a vacuum produced 10 g (1.25%). The alkaloid extract (9 g) was subjected to column chromatography on a normal phase silica gel 60 column using gradient polarity (v/v) of  $\text{CH}_2\text{Cl}_2$ :  $\text{CH}_3\text{OH}$  10:0, 9:1, 8:2, 7:3, 3:2 1:1, (500 ml in each case) and finally 100% MeOH.

Following the use of Dragendorff's reagent spray on TLC plates, 34 fractions generated by this procedure were recombined into 10 fractions, which were then vacuum-concentrated using a rotary evaporator. Following their elution with 1%  $\text{CH}_3\text{OH}$  in  $\text{CH}_2\text{Cl}_2$ , fractions 2 (35 mg) and 3 (72 mg) were combined and further separated using preparative thin-layer chromatography (PTLC) developed with 8%  $\text{CH}_3\text{OH}$  in  $\text{CH}_2\text{Cl}_2$ . Two UV-active compounds, compound **(9)** and compound **(43)**, were obtained by scrapping the bands associated with each isolated component, dissolving it in DCM, and filtering the mixture.

Based on their TLC profiles, fractions 4 and 5 were eluted with  $\text{CH}_2\text{Cl}_2$ :  $\text{CH}_3\text{OH}$  8:2 v/v were combined and then subjected to a column (60 cm  $\times$  3.0 cm,  $\text{SiO}_2$ ) chromatography eluting with  $\text{CH}_2\text{Cl}_2$ :  $\text{CH}_3\text{OH}$  gradient polarity of 9:1, 8:2, and 7:3 yielding 8 fractions (30mL). These fractions were then recombined into three fractions.

Compounds **21** (30 mg), **9** (50 mg), and **42** (80 mg) were produced through recrystallization using  $\text{CH}_2\text{Cl}_2$ : n-hexane from sub-fractions one, two, and three of Fractions 4 and 5. Fractions 6 (410 mg) and 7 (296 mg) were eluted with 7:3 v/v  $\text{CH}_2\text{Cl}_2$ : $\text{CH}_3\text{OH}$  mixtures. Based on their TLC profiles, these fractions were combined and subjected to column (60 cm  $\times$  4.0 cm,  $\text{SiO}_2$ ) chromatography. Twelve fractions (25 ml each) were obtained, which were then combined into four fractions in accordance with their TLC profiles.

Sub-fraction one of fraction 6 and 7 yielded crystalline compound **39** (86 mg) from  $\text{CH}_2\text{Cl}_2$ : n-hexane mixture. Third portion of fraction combined 6 and 7 yielded compound **40** (77 mg) as white crystalline solids from  $\text{CH}_2\text{Cl}_2$ : n-hexane mixture. Lastly, fractions 8, 9 and 10 (158, 214 and 176 mg, respectively) eluted with 1:1 v/v  $\text{CH}_2\text{Cl}_2$ :  $\text{CH}_3\text{OH}$  were combined and re-chromatographed on CC (60 cm and 4.0 cm,  $\text{SiO}_2$ ), gradient elution with  $\text{CH}_2\text{Cl}_2$ :  $\text{CH}_3\text{OH}$  (7:3, 1:1 and 3:7 v/v). Four fractions were obtained, from which the second portion yielded compound **41** (30 mg) after

crystallization with CH<sub>2</sub>Cl<sub>2</sub>:CH<sub>3</sub>OH. Third sub-fraction yielded compound **41** (45 mg) and yellow solids of compound **42** (61 mg), following fractional re-crystallizations in CH<sub>2</sub>Cl<sub>2</sub>: *n*-hexane.

Alkaloid-free extracts: The aqueous non-alkaloid fraction was concentrated *in vacuo* to dryness and afforded 10.9 g (1.36% of crude extracts) which was then suspended in 200 mL distilled water followed by sequential solvent extractions beginning with 3 × 300 mL *n*-hexane, followed by 3 × 300 mL ethyl acetate, and finally 3 × 300 mL *n*-butanol (C<sub>4</sub>H<sub>9</sub>OH) using a separatory funnel. The combined decanted organic layer extracts, were evaporated in *vacuo* to yield 2.05 g (0.13%), 5.01 g (0.63%) and 3.1 g (0.39%), respectively.

Dry *n*-hexane non-alkaloid extract (2 g) was loaded on silica gel in column (60 × 3 cm, SiO<sub>2</sub>) and eluted stepwise with *n*-hexane adjusting the polarity with ethyl acetate as (1:0, 99:1, 95:5, 9:1, 4:1, 3:1, 2:1). Out of the 35 eluent fraction (25 mL each), five fractions were obtained after pooling the fractions with similar TLC profiles and monitored using *p*-anisaldehyde-sulphuric acid reagent. 80 mg of Compound **10** was obtained as white crystals from the third fraction after crystallizing it in methanol. Further purification of the fifth fraction using silica gel column (120 × 60 mm, SiO<sub>2</sub>) eluting with *n*-hexane: ethyl acetate (9:1, 4:1) gave more of the white crystals of 25 mg with single spot on TLC which turned purple – blue after spraying the developed plate with *p*-anisaldehyde-sulphuric acid, characteristics of a terpenoid. A portion of *n*-C<sub>4</sub>H<sub>9</sub>OH fraction was (1.4 g) also chromatographed on reverse (C18 reverse phase SiO<sub>2</sub>, 40–63 μm) flash chromatography, eluting with H<sub>2</sub>O-CH<sub>3</sub>OH (9:1); CH<sub>3</sub>OH; CH<sub>3</sub>OH - CH<sub>2</sub>Cl<sub>2</sub> and finally CH<sub>2</sub>Cl<sub>2</sub>. A total of 30 fractions (each 50 ml) which were further combined to 3 fractions according to their TLC profiles were obtained. The first fraction yielded a white amorphous solid that was confirmed to be sesamine (**46**, 103 mg), the second fraction yielded few mixtures of compound (**46**) and compound (**44**) that crystallized in CH<sub>2</sub>Cl<sub>2</sub>; *n*-hexane as white crystalline solid (62 mg).

### 3.2.2 Extraction and Isolation of compounds from *Z. gilletii*

*Z. gilletii* stem bark were chopped into small pieces separately, air-dried at room temperature under shade for 25 days and pulverized into fine powder. The powdered stem bark material (1.05 kg) were extracted with 95% aqueous methanol (3 × 2.5 L) for 72 hours and filtered. The filtrates were combined and concentrated in *vacuo* using rotary evaporator to yield 53.74 g (5.21 %).

Total alkaloid extraction: The crude extracts suspended in 200 mL of 2% 0.1 M H<sub>2</sub>SO<sub>4</sub> in dichloromethane under agitation for 40 min, replicated four times and the extracts were combined, followed by three times extraction of the non-polar compounds using 300 mL

*n*-hexane using a separatory funnel. The acidic polar fractions was neutralized using 0.25 M ammonia hydroxide solution, followed by further extraction using 200 mL dichloromethane (CH<sub>2</sub>Cl<sub>2</sub>), until a negative Dragendorff's test was observed. The organic fraction (CH<sub>2</sub>Cl<sub>2</sub> extracts) was dried with anhydrous sodium sulphate, followed by further concentration *in vacuo* to yield 15.5 g (1.48 % of the crude extract).

The alkaloid crude extract (12.4 g) was subjected to a silica gel CC (8 × 120 cm, 1.7 kg SiO<sub>2</sub>) and eluted *n*-hexane:CH<sub>2</sub>Cl<sub>2</sub> mixture (1 : 0 to 9 : 1, 10% gradient, 500mL of each eluent; 9:1, 4:1, 7:3 to 3 : 2, 10% gradient, 500 mL of each eluent; 1:0 to 0 :1, 10% gradient, 500 mL of each eluent) and CH<sub>2</sub>Cl<sub>2</sub>-MeOH (1 : 0 to 9 : 1, 10% gradient, 500mL of each eluent) to yield 40 crude fractions which were combined to 6 fractions (D1–D6). TLC profiles of the five fractions then monitored with *p*-anisaldehyde-sulphuric acid reagent. The first three fractions D1, D2 and D3 showed one spot each using the UV lamp. The single spots were confirmed by heating the TLC plates using Heat gun at 100°C. The fractions were then vaporized *in vacuo* and produced two compounds; Compounds **9**, yellow solid, (40 mg), and **43**, (35 mg) were obtained from fractions D1, D2, and D3 respectively. 2.4g of fraction D4 subjected to silica gel CC (5 × 80 cm, 160 g SiO<sub>2</sub>) using mixtures of *n*-hexane-CH<sub>2</sub>Cl<sub>2</sub> (1:0 to 7: 3, 10% gradient, 500mL of each eluent; 7:3 to 0:1, 10% gradient, 500mL of each eluent) yielding compound **9** (110 mg) as yellow solid. Further Fraction D5 (1.7 g) was eluted over silica gel CC (3 × 30 cm, 85 g SiO<sub>2</sub>) with *n*-hexane-CH<sub>2</sub>Cl<sub>2</sub> (1:4 to 0:1, 10% gradient, 500 mL of each eluent) and CH<sub>2</sub>Cl<sub>2</sub>-MeOH (1:0 to 9:1, 10 gradients, 0.2 L of each eluent) to afford white crystals of compounds **43** (45.5 mg) and yellow crystals of compound **42** (56.8 mg).

Alkaloid-free extracts: Aqueous non-alkaloid portion was concentrated *in vacuo* to dryness obtaining 13.5 g (1.29 % of crude extracts) then suspended in 200 mL distilled water. Sequential solvent extractions was done beginning with 3 × 300 mL *n*-hexane, then, 3 × 300 mL ethyl acetate, lastly, 3 × 300 mL *n*-butanol (C<sub>4</sub>H<sub>9</sub>OH) in a separatory funnel. The combined decanted organic layer extracts, were evaporated *in vacuo* to yield 2.23 g (0.21%), 6.38 g (0.61%) and 4.1 g, respectively.

*n*-Hexane extract (2.2 g) subjected on silica gel in column (60 x 3 cm, SiO<sub>2</sub>) and eluted with *n*-hexane-EtOAc (99:1, 95:5, 9:1, 4:1, 7:3, 3:2, 2:1 and 1:1) 200ml each eluent resulting into 40 fractions (50 ml each) the fractions were combined according their similar TLC profiles monitored using *p*-anisaldehyde-sulphuric acid reagent giving 3 fractions. The first fraction gave white precipitate which produce a blue sport on TLC under UV lamp *n*-hexane-EtOAc (9:1) which dried

yielding white amorphous solid ( $R_f = 0.36$ ) confirmed to be compound **12**, (56.8 mg) using TLC profile in comparison to authentic sample. The second fraction showed two spots on TLC and was dried *in vacuo* using a rotatory removed evaporator and re-crystallized using 60 %  $\text{CH}_2\text{Cl}_2$  in *n*-hexane to produce a white amorphous solid (**46**, 31.2 mg). The mother liquor was recrystallized from 80 %  $\text{CH}_2\text{Cl}_2$  in *n*-hexane, filtered and dried yielding, compound **44** (14.25 mg).

Dry EtOAc extract was eluted using *n*-hexane - EtOAc (99:1, 9:1, 4:1, 3:1, and 1:1) on silica gel CC giving twenty fractions. The fractions were then pooled and combined according to their similar TLC profiles to produce five fractions (E1 (1-7), E2 (8-19), E3 (20-26), E4 (27-36) and E5 (37-40). E1 showed a single blue spot on TLC. The portion was dried producing white amorphous solid (63.7 mg) producing a purple-blue spot when sprayed with *p*-anisaldehyde-sulphuric acid mixture and heated at  $110^\circ\text{C}$ , characteristic of a terpenoid. Fraction E4 was dried *in vacuo* on a rotatory evaporator and loaded on a Sephadex LH 20 column leading to isolation of a white amorphous solid of compound (**44**) (15.15 mg).

### 3.2.3 Physical and spectroscopic data pure isolates from *Z. Chalybeum* and *Z. gilletti*

**Compound (9)** : yellow crystals; UV active with blue fluorescence; melting point of  $210\text{-}212^\circ\text{C}$ ;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 9); ESIMS  $[\text{M}+\text{H}]^+$   $m/z = 334$  (calcd for  $\text{C}_{20}\text{H}_{15}\text{NO}_4$ ).

**Compound (10)**: white crystals; melting point of  $124 - 125^\circ\text{C}$ ;  $^1\text{H}$  and  $^{13}\text{C}$  NMR ( $\text{CDCl}_3$ ) data (Table 8); calcd for  $\text{C}_{10}\text{H}_{10}\text{O}_4$ .

**Compound 21**: yellow powder; UV active; melting point of  $180^\circ\text{C}$ ;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 13; ESIMS:  $[\text{M}+1]^+$   $m/z$  260 (calcd. For  $\text{C}_{14}\text{H}_{13}\text{NO}_4$ ).

**Compound 39**: white powder; UV ( $\text{CHCl}_3$ )  $\lambda_{\text{max}}$  ( $\log \epsilon$ ): 305 (0.2), 265 (0.47) nm; IR (KBr)  $\nu_{\text{max}}$ : 3457, 3316, 2943, 1625, 1551, 1505, 1466, 1257  $\text{cm}^{-1}$ ;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 1); HREIMS:  $m/z$  287.1317  $[\text{M}]^+$  (calcd. for  $\text{C}_{17}\text{H}_{21}\text{NO}_3$ , 287.1312).

**Compound 40**: white crystalline; UV ( $\text{CHCl}_3$ )  $\lambda_{\text{max}}$  ( $\log \epsilon$ ): 284 (0.76), 218 (0.48), 204 (0.24) nm; IR (KBr)  $\nu_{\text{max}}$ : 3424, 1736, 1649, 1560, 1431, 1398, 1255, 1202, 1070  $\text{cm}^{-1}$ ;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 2); HREIMS:  $m/z$  219.1103  $[\text{M}]^+$  (calcd. for  $\text{C}_{13}\text{H}_{17}\text{NO}_2$ , 219.1105).

**Compound 41**: white crystalline; UV ( $\text{CHCl}_3$ )  $\lambda_{\text{max}}$  ( $\log \epsilon$ ): 284 (0.76), 218 (0.48), 204 (0.24) nm; IR (KBr)  $\nu_{\text{max}}$ : 3424, 1736, 1649, 1560, 1431, 1398, 1255, 1202, 1070  $\text{cm}^{-1}$ ;



$^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 3); HREIMS:  $m/z$  285.11373  $[\text{M}]^+$  (calcd. for  $\text{C}_{17}\text{H}_{19}\text{NO}_3$ , 285.1365).

**Compound 42:** yellow crystals; melting point range 199-200°C;  $^1\text{H}$  NMR ((DMSO- $d_6$ )) and  $^{13}\text{C}$  NMR (DMSO- $d_6$ ) (Table 4); Rf value 0.64 ( $\text{CH}_2\text{Cl}_2$ : MeOH (4:1)); ESIMS:  $[\text{M}+\text{H}]^+$   $m/z$  349.1057 (calcd for  $\text{C}_{21}\text{H}_{18}\text{NO}_4^+$ ).

**Compound 43:** white crystals (DCM:MeOH extract); yield (45.5mg); Rf of 0.54 (40% DCM in hexane); Mp of 141-142 °C;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 5); HREIMS:  $m/z$  497.1839  $[\text{M}]^+$  (calcd. for  $\text{C}_{30}\text{H}_{27}\text{NO}_6$ , 497.1838).

**Compound 44 :** white solid (n-hexane: EtOAc extract); yield (15.15 mg) Rf of 0.47 (80% EtOAc in n-hexane); Mp of 191-192°C;  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 6); HREIMS:  $m/z$  324.0996  $[\text{M}]^+$  (calcd. for  $\text{C}_{19}\text{H}_{16}\text{O}_5$ , 324.0998).

**Compound 45:** white crystal; Rf of 0.57 (80% DCM in MeOH);  $^1\text{H}$  and  $^{13}\text{C}$  NMR data (Table 7); HREIMS:  $m/z$  268.1322 (calcd. for  $\text{C}_{12}\text{H}_{23}\text{NO}_4$ , 268.1324).

**Compound 46:** white solid; UV active; melting point of 121-122°C; HREIMS;  $m/z$  354.35 cald for  $\text{C}_{20}\text{H}_{18}\text{O}_6$ .

### 3.2.4 $\alpha$ -Amylase inhibition assay

The anti-hyperglycemic activities of the pure isolated compounds were assayed against human salivary  $\alpha$ -amylase using assay as described in the Worthington Enzyme Manual (Worthington, 1993; Kwon *et al*, 2006). Pre-incubation was carried out for 10 minutes at 25°C in 500  $\mu\text{L}$  of 0.02M sodium phosphate buffer (pH 6.9 with 0.006 M NaCl) containing 0.5 mg/mL of  $\alpha$ -amylase. 500  $\mu\text{L}$  of a 1% starch solution in 0.02M sodium phosphate buffer (pH 6.9 with 0.006M NaCl) was added to each tube containing pure isolates (0.05mM, 0.1mM, 0.2mM, 0.5mM, 1.0mM) at timed intervals. 1.0 mL of dinitrosalicylic (DNS) acid color reagent was used to stop the reaction. A boiling water bath was used to incubate the test tubes for five minutes, after which they were allowed to cool to room temperature. The reaction mixture was diluted by adding 5 to 15 mL of distilled water. Fluorescence measured using a SpectraMax GeminiXS Spectrofluorometer (Molecular Devices, Sunnyvale, CA) with excitation and emission wavelengths of 540 nm. The absorbance readings were compared with the controls that contain buffer instead of sample extract. The percentage  $\alpha$ -amylase inhibitory activity was calculated using the equation;

$$\%inhibition = \frac{Absorbance (control) - Absorbance (Extract)}{Absorbance (Control)} \times 100$$

The percentage inhibitions were expressed as probit units then regressed against concentrations of the samples to obtain the 50% inhibitory concentration of each sample (IC<sub>50</sub>).

### 3.2.5 $\alpha$ -Glucosidase inhibition assay

The anti-hyperglycemic activity of the pure isolated compounds were assayed against  $\alpha$ -glucosidase (Omosa *et al*, 2019) pure compounds prepared as described in  $\alpha$ -amylase assay above. The test compound and 2 mU of  $\alpha$ -glucosidase was diluted to 97  $\mu$ L in 0.1 M potassium phosphate buffer (pH 6.5) and pre-incubated in 96-well plates at 37 °C for 15 min. The reaction was initiated by adding 3  $\mu$ L of 3 mM *p*-nitrophenyl- $\alpha$ -D-glucopyranoside (*p*-NPG) as substrate and incubated for an additional 15 min at 37 °C, followed by addition of 100  $\mu$ L of 1 M Na<sub>2</sub>CO<sub>3</sub> to stop the reaction. All test compounds were prepared in 5% DMSO in distilled water. The final concentrations of the compounds were between 5 and 200  $\mu$ M while the final concentration of  $\alpha$ -Glucosidase was 20 mU/mL. The enzyme activity was then determined by measuring the release of *p*-nitrophenol from the *p*-NPG substrate and the reaction monitored by change of absorbance at 410 nm using a SpectraMax 190 Spectrophotometer.

The  $\alpha$ -Glucosidase inhibition rate was calculated relative to control as follows:

$$\%inhibition = \frac{Absorbance (control) - Absorbance (Extract)}{Absorbance (Control)} \times 100$$

### 3.2.6 Kinetic Analyses

Mode of inhibition of compounds against porcine pancreas  $\alpha$ -amylase and yeast  $\alpha$ -glucosidase activity were measured with increasing concentrations of *p*NPG (0.25, 0.5, 1, 2 and 5 mM) as a substrate, in the absence and presence of the pure compounds and acarbose at 0, 0.5, 1, 2.5, 10 and 20 mM. The mode of inhibition was determined by the Lineweaver-Burk plots, followed by secondary plots (Dixon plots) depending on the established mode of inhibition. The inhibition constants (K<sub>i</sub>) were calculated using the following equation (Segel, 1993):

$$v = V_{max} \times S / K_m (1 + [I] / K_i) + S (1 + [I] / \alpha K_i)$$

where  $v$  is the initial velocity in the absence and presence of inhibitor;  $S$  and  $I$  are the concentration of substrate and inhibitor, respectively;  $V_{\max}$  is the maximum velocity;  $K_m$  is the Michaelis-Menten constant;  $K_i$  is the competitive inhibition constant and  $\alpha K_i$  is the uncompetitive inhibition constant.

### **3.3 Statistical Analysis**

Lineweaver-Burk and Dixon secondary linear regression analysis used for establishing the inhibitory constants were developed on MS Excel 2019. Variance analysis was performed using Tukey HSD/Tukey Kramer post-analysis (Segel, 1993).

## CHAPTER FOUR

### RESULTS AND DISCUSSION

#### 4.1 Structure Elucidations

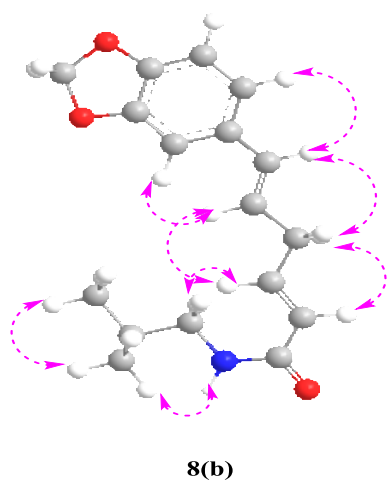
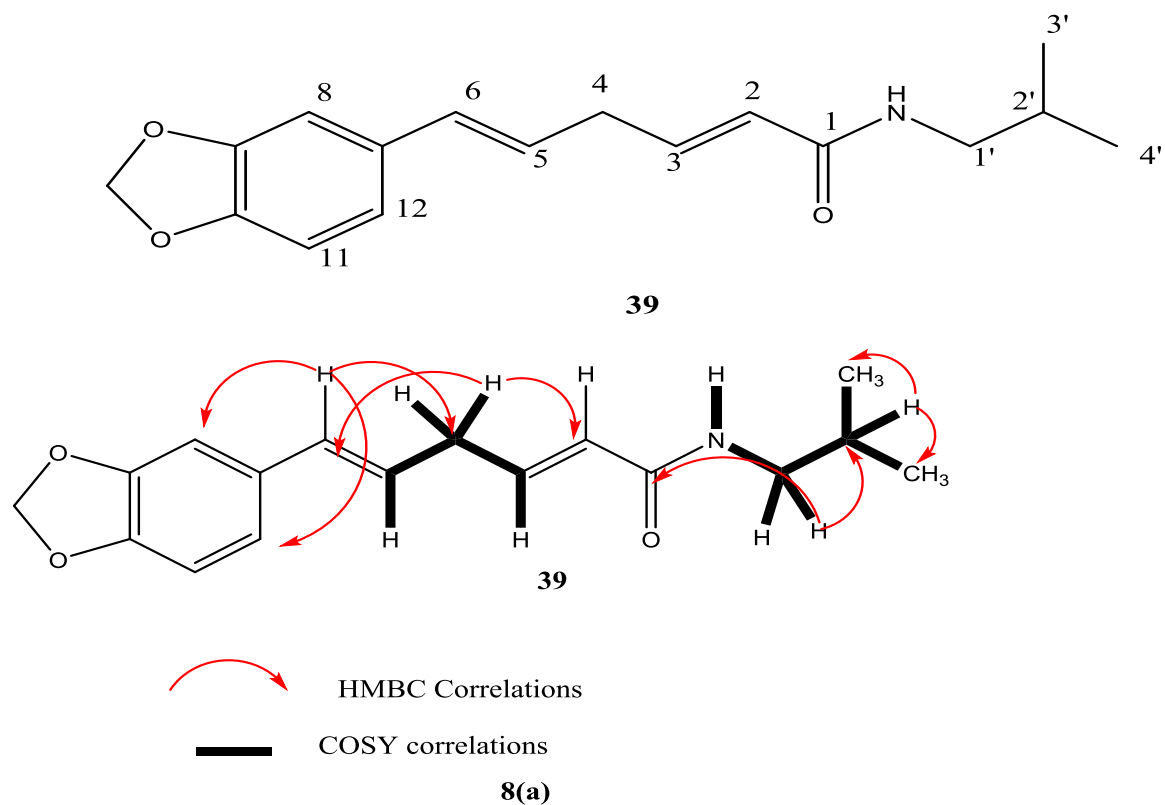
##### 4.1.1 Chalybemide A {6-benzo[1,3]dioxol-5-yl-hexa-2,5-dienoic acid isobutylamide} (39)

Compound 39 was isolated as white powder with molecular mass determined by HREIMS,  $m/z = 287.1317[M]^+$  (calcd for  $C_{17}H_{21}NO_3$  287.1312) giving its formula as  $C_{17}H_{21}NO_3$  indicating eight degrees of unsaturation. The IR spectrum showed an amide functionality, aromatic and/or alkene based on  $3457\text{ cm}^{-1}$ ,  $1505\text{-}1447\text{ cm}^{-1}$  absorptions, respectively.

From  $^1\text{H}$  and  $^{13}\text{C}$  spectra (Table 1), characteristic signal for a benzo[1,3]dioxol unit was predicted based on the protons  $\delta_{\text{H}} 5.92$  (br, s, 2H,  $\text{OCH}_2\text{O}$ ) attached to  $\delta_{\text{C}} 99.39$ ,  $\delta_{\text{H}} 6.89$  (d,  $J = 1.2$  Hz, H-8) attached to  $\delta_{\text{C}} 107.87$ ,  $6.73$  (d,  $J = 8.0$  Hz, H-11) attached to  $\delta_{\text{C}} 109.14$  and  $\delta_{\text{H}} 6.75$  (dd,  $J = 8.0, 1.2$  Hz, H-12) attached to  $120.15$  and  $\delta_{\text{C}} 132.30, 147.87$  and  $\delta_{\text{C}} 148.56$  carbon signals closely related to the data reported by Li et al., (2019). The HMBC correlations supported the benzo[1,3]dioxol as observed from  $\delta_{\text{H}} 5.92$  ( $\text{OCH}_2\text{O}$ ) displaying contours to  $\delta_{\text{C}} 147.9$  (C-9),  $148.6$  (C-10); while proton at  $\delta_{\text{H}} 6.87$  (H-8) correlating to  $\delta_{\text{C}} 148.6$  (C-10) and  $\delta_{\text{C}} 120.2$  (C-12); proton at  $\delta_{\text{H}} 6.73$  (H-11) correlating to  $\delta_{\text{C}} 132.5$  (C-7) and  $\delta_{\text{C}} 147.9$  (C-9) and proton at  $\delta_{\text{H}} 6.83$  (H-12) to  $\delta_{\text{C}} 107.8$  (C-8) and  $\delta_{\text{C}} 148.6$  (Figure 8a).

Another characteristic signal indicated the sets of *trans* oriented olefinic protons at  $\delta_{\text{H}} 5.74$  (d,  $J = 15.2$  Hz, H-2),  $6.05$  (dd,  $J = 15.2, 7.0$  Hz, H-),  $\delta_{\text{H}} 6.77$  (d,  $J = 15.2$  Hz, H-6) attached to two pairs of olefinic carbon signals at  $\delta_{\text{C}} 123.8, 146.8, 129.9$  and  $129.5$  as supported by DEPT and HSQC (Table 1).  $^1\text{H}$ - $^1\text{H}$  COSY showed correlation between  $\delta_{\text{H}} 2.39$  (m, 2H) with two of the olefinic protons;  $\delta_{\text{H}} 5.74$  (d,  $J = 15.2$ , H-2) and  $\delta_{\text{H}} 6.24$  (dd,  $J = 15.2, 7.0$ , H-5) together with HMBC correlations to carbon signals at  $\delta_{\text{C}} 123.8$  and  $\delta_{\text{C}} 129.5$  indicated a methylene link between the  $\text{C}=\text{C}$  showing the presence of penta-1,3-diene moiety. The  $^1\text{H}$  NMR further displayed a signal at  $\delta_{\text{H}} 3.13$  (t,  $J = 6.0$  Hz, 2H) attached to  $\delta_{\text{C}} 49.5$  (HSQC) which showed COSY coupling with N-H and another proton at  $\delta_{\text{H}} 1.78$  (m, 1H) attached to  $\delta_{\text{C}} 29.3$ , which coupled  $\delta_{\text{H}} 0.98$  (d,  $J = 6.0$  Hz, 2- $\text{CH}_3$ ) indicating the presence of isobutylamide moiety (Mansoor *et al*, 2013). Furthermore, HMBC correlations between  $\delta_{\text{H}} 3.13$  ( $\text{CH}_2$  to  $\delta_{\text{C}} 169$  ( $^3J, \text{C}=\text{O}$ ) and  $\delta_{\text{C}} 29.3$  ( $^2J, \text{C}-2'$ );  $^2J$  correlations between the methane proton  $\delta_{\text{H}} 1.78$  and methylene carbon ( $\delta_{\text{C}} 49.5$ ) and methyl carbons ( $\delta_{\text{C}} 19.4$ ) (Figure 8A) confirmed  $(\text{C}=\text{O})\text{NHCH}_2\text{CH}(\text{CH}_3)_2$  moiety, a common structural features of *Zanthoxylum* amide alkaloids. These NMR spectra established a

typical amide alkaloid except the presence of penta-1,3-diene moiety, from which one of the alkene proton at  $\delta_{\text{H}}$  6.77 (H-6) and  $\delta_{\text{C}}$  129.2 showed a mutual correlation to benzo[1,3]dioxol signal at  $\delta_{\text{H}}$  6.24 and  $\delta_{\text{C}}$  107.9 (C-8), while another olefinic proton  $\delta_{\text{H}}$  5.74 (H-2) correlation to the carbonyl carbon ( $\delta_{\text{C}}$  169.3, C-1) and the methylene carbon ( $\delta_{\text{C}}$  35.3) (Figure 8a) confirming penta-1,3-diene as the moiety joining the isobutylamide and the benzo[1,3]dioxol. These spectral evidences characterized compound **39** as 6-benzo[1,3]dioxol-5-yl-hexa-2,5-dienoic acid isobutylamide obtained as new natural product from *Z. chalybeum* and named as chalybemide A.



**Figure 8:** (a) is HMBC, COSY significant correlations and 8(b) is ROESY correlations of compound **39**

**Table 1: <sup>1</sup>H (600MHz) and <sup>13</sup>C NMR (150 MHz) (CDCl<sub>3</sub>) spectral data for Chalybemide A**

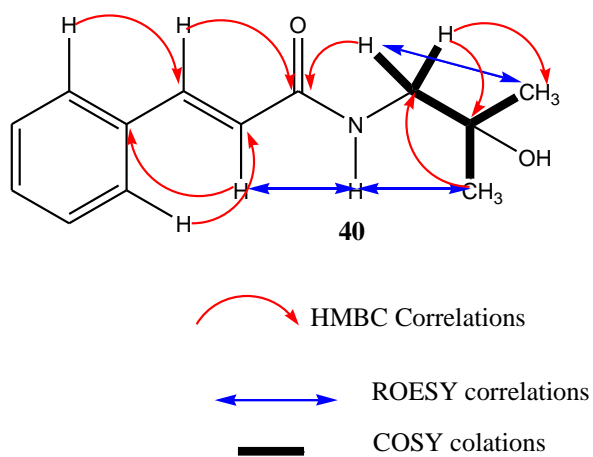
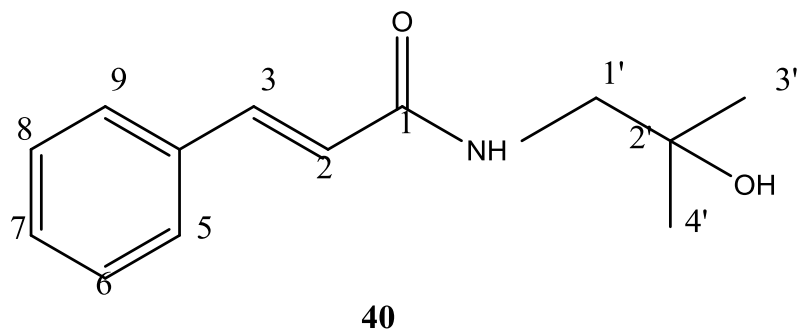
C- Position	<sup>1</sup> H, multi	<sup>13</sup> C	HMBC	COSY
1		169.3(C=O)		
2	5.74, d,( <i>J</i> = 15.2)	123.8 (CH)		
3	6.05 dd ( <i>J</i> = 15.2, 7)	146.8 (CH)		H <sub>3</sub> -H <sub>4</sub>
4	2.39 m	35.3 (CH <sub>2</sub> )		H <sub>4</sub> -H <sub>5</sub>
5	6.24 dd ( <i>J</i> = 15.2, 7)	129.2 (CH)		
6	6.77 d ( <i>J</i> = 15.2)	129.5 (CH)		
7		132.5 (C)		
8	6.89 d ( <i>J</i> = 1.2)	107.9 (CH)	C-10,12	
9		147.9 (C-O)		
10		148.6 (C-O)		
11	6.73 d ( <i>J</i> = 1.2)	109.1 (CH)	C-7,9	
12	6.83 dd ( <i>J</i> = 8, 1.2)	120.2 (CH)	C-8,10	
1'	3.13 t ( <i>J</i> = 15.2)	49.5 (CH <sub>2</sub> )	C-1,2'	H <sub>1</sub> '-H <sub>2</sub> '
2'	1.78 m	29.3 (CH)		H <sub>2</sub> '- H <sub>3</sub> ',H <sub>2</sub> '-H <sub>4</sub> '
3'	0.91 d ( <i>J</i> = 15.2)	19.3 (CH <sub>3</sub> )		
4'	0.94 d ( <i>J</i> = 15.2)	19.3 (CH <sub>3</sub> )		
OCH <sub>2</sub> O	5.92 br s	99.4 (CH <sub>2</sub> )	C-9,10	

**4.1.2 Chalybemide B {4-Methoxy-*N*-(2-methoxy-phenyl)-*N*-methyl-benzamide}**

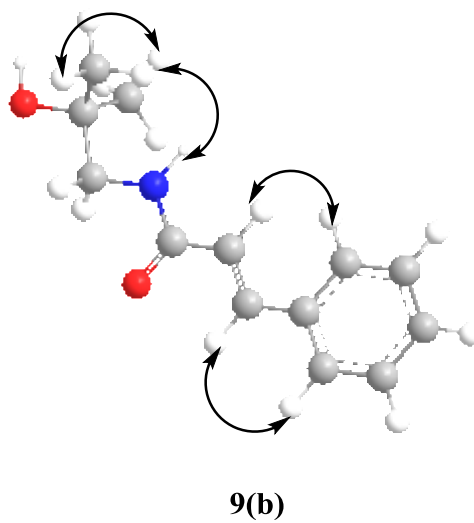
Compound **40** was obtained as white crystalline solids with molecular formula C<sub>13</sub>H<sub>15</sub>NO<sub>2</sub> deduced from its HREIMS at *m/z* 219.1103 [M]<sup>+</sup> (calculated for C<sub>13</sub>H<sub>15</sub>NO<sub>2</sub>, 219.1103)

predicted for seven degrees of unsaturation.  $^{13}\text{C}$  NMR (Table 2) supported the mass spectrometry data by exposing thirteen carbon atoms characterized by DEPT-135 spectra as seven methines, one methylene and three quaternary carbons including one hydroxylated and one carbonyl carbon.

$^1\text{H}$  NMR displayed two aromatic signals as two broad doublets integrated for two protons at  $\delta_{\text{H}}$  7.36 (d,  $J = 7.2$  Hz H-6 (2H)) and  $\delta_{\text{H}}$  7.53(d,  $J = 7.2$  Hz) and multiplet at  $\delta_{\text{H}}$  7.47 (1H) attributed to a monosubstituted aromatic ring. The presence of *trans* configured alkene moiety was noted based on the signals  $\delta_{\text{H}}$  6.64 (d,  $J = 15.5$ ) and  $\delta_{\text{H}}$  7.73 (d,  $J = 15.5$ ), which showed HMBC correlations with carbonyl carbon at  $\delta_{\text{C}}$  166.5 (C=O) and quaternary aromatic carbon at  $\delta_{\text{C}}$  134.9 (C-4) predictive of a phenylethylene skeleton attached to carbonyl carbon. Moreover, HSQC shows that proton signal at  $\delta_{\text{H}}$  3.47 (-CH<sub>2</sub>, s) attached to  $\delta_{\text{C}}$  58.8 having a COSY correlations with N-H and HMBC correlations to  $\delta_{\text{C}}$  70.1 (C-2') and to carbonyl  $\delta_{\text{C}}$  166.5 suggesting a methylene amide group. HMBC further shows two mutually coupled methyl groups ( $\delta_{\text{H}}$  1.19) to carbon at  $\delta_{\text{C}}$  70.1 (C-2') and to  $\delta_{\text{C}}$  57.1 (C-H<sub>2</sub>) (Figure 9(a) confirming the moiety as a 2-isobutylamide (Djeukeu et al, 2019). Following HMBC correlations of the ethylene protons and methylene protons to a carbonyl carbon and the ROESY correlations (Figure 9(b)) which confirmed compound **40** as *N*-(2-hydroxy-2-methylpropyl)-3-phenylacrylamide, isolated and characterized from *Zanthoxylum chalybeum* for the first time and named as Chalybemide B.



**Figure 9 (a): HMBC, ROESY and COSY for compound 40**



**Figure 9 (b): 3D and ROESY correlations for 40**



**Table 2: <sup>1</sup>H (600MHz) and <sup>13</sup>C NMR (150 MHz) (CDCl<sub>3</sub>) spectral data for Chalybemide B**

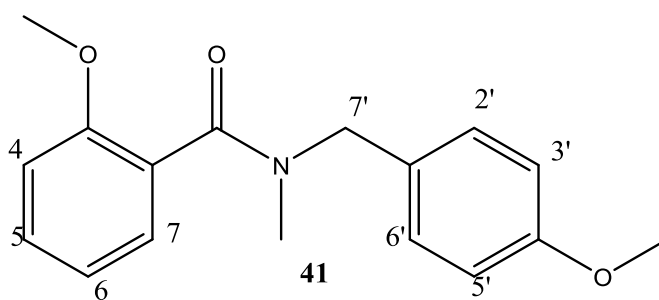
C-Position	$\delta$ H, mult (J, Hz)	$\delta$ C	HMBC	COSY
1		166.5(C=O)		
2	6.64 (d, $J = 15.5$ )	119.8(CH)	C-4	
3	7.73 (d, $J = 15.5$ )	141.6 (CH)	C-1	
4		134.9 (C)		
5	7.53 (d, $J = 7.2$ )	126.2 (CH)	C-2	
6	7.36 (d, $J = 7.2$ )	128.4 (CH)		
7	7.47 m	127.7 (CH)		
8	7.36 m	128.4 (CH)		
9	7.53 m	126.2 (CH)	C-3	
1'	3.47 br s	58.8 (CH <sub>2</sub> )	C-1, 2', 3'	H <sub>1'</sub> -H <sub>2'</sub>
2'	–	70.1 (C-O)		
3'	1.19 br s	27.2 (CH <sub>3</sub> )		H <sub>3'</sub> -H <sub>4'</sub>
4'	1.19 br s	27.2 (CH <sub>3</sub> )		

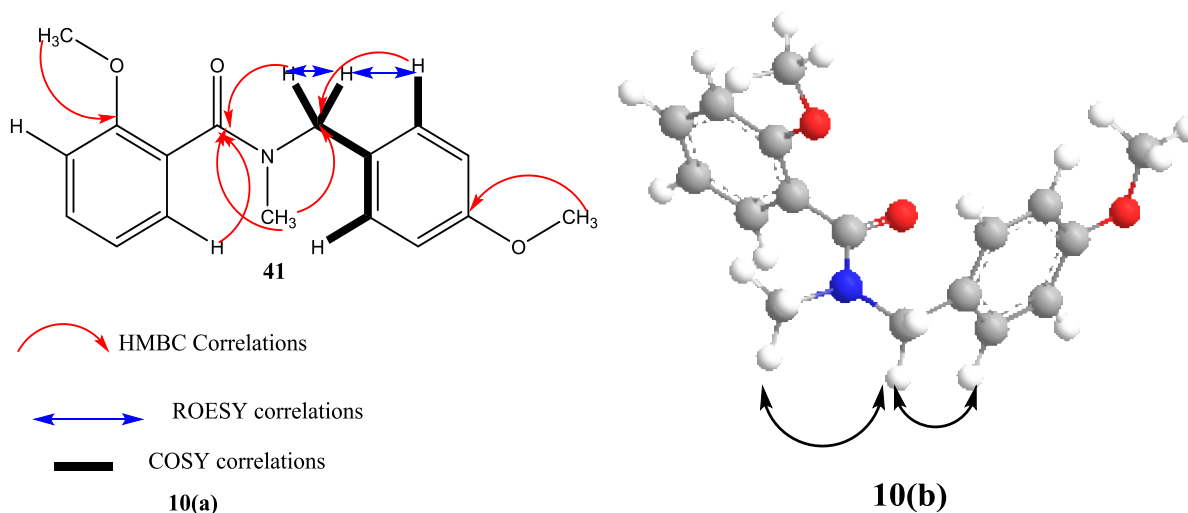
**4.1.3 Chalybemide C {*N*-(2-Hydroxy-2-methyl-propyl)-3-phenyl-acrylamide} (41)**

Compound 41 was isolated as white crystals with molecular formula of C<sub>17</sub>H<sub>19</sub>NO<sub>3</sub> determined by HREIMS at  $m/z$  285.1373[M]<sup>+</sup> (cald. for. C<sub>17</sub>H<sub>19</sub>NO<sub>3</sub>, 285.1373), indicating nine degrees of unsaturation. <sup>1</sup>H NMR spectrum displayed a pair of protons which were mutually coupled with aromatic signals at  $\delta$ H 6.86 (d,  $J = 8.5$  Hz, 2H) and  $\delta$ H 7.15 (d,  $J = 8.5$  Hz, 2H) typical of *para* disubstituted ring. Another ortho disubstituted benzene ring signaled by  $\delta$ H 6.89 (1H, d,  $J = 8.5$  Hz, H-3),  $\delta$ H 7.52 (1H, m, H-4),  $\delta$ H 7.05 (1H, t,  $J = 7.7$  Hz, H-5) and  $\delta$ H 7.86 (1H, dd,  $J = 7.7, 1.8$  Hz, H-6). <sup>13</sup>C NMR (Table 3) spectra predicted the presence of one aliphatic methylene ( $\delta$ C 57.1), an *N*-methyl ( $\delta$ C 34.9), three oxygenated carbons: two methoxyls ( $\delta$ C 54.4 and 56.6) and a carbonyl ( $\delta$ C 168.1) and twelve aromatic carbons as classified by DEPT.

The data was comparable to that of tessmamide previously isolated from *Zanthoxylum tessmanii* (Kimani, 2015) except for the presence of methylene group and absence of methylenedioxy group on the compound.

HMBC and COSY correlations (Figure 10A) confirmed the overall connectivity and the planar structure of compound 41. A proton at  $\delta_{\text{H}}$  3.73 (OCH<sub>3</sub>) showed a  $^3J$  correlations to  $\delta_{\text{C}}$  113.2 (C-3'/C-5') and  $\delta_{\text{H}}$  6.86 (H-3'/H-5') to  $\delta_{\text{C}}$  56.4 (OCH<sub>3</sub>) placing it at C-4' in ring B and the other methoxy group  $\delta_{\text{H}}$  3.72 correlated to  $\delta_{\text{C}}$  112.2 (C-1) and  $\delta_{\text{C}}$  122.2 (C-4). In addition, a proton  $\delta_{\text{H}}$  6.89 (H-3) correlated to  $\delta_{\text{C}}$  56.6 (OCH<sub>3</sub>) placing it at C-3 in ring A. Moreover, a proton for N-CH<sub>3</sub> showed a  $^3J$  correlation to a methylene carbon  $\delta_{\text{C}}$  57.1 (C-7') and to carbonyl carbon  $\delta_{\text{C}}$  168.1 (C-1) corroborated by COSY off diagonal peaks between  $\delta_{\text{H}}$  3.19 (N-CH<sub>3</sub>) and  $\delta_{\text{H}}$  3.91 (CH<sub>2</sub>) which in turn coupled with  $\delta_{\text{H}}$  7.15 (H-2'/H-6') and confirmed *N,N*-methylene-methyl amide moiety. From the data mentioned above, compound 41 was characterized as 4-methyl-*N*-(2-methoxyphenyl)-*N*-methylbenzamide, isolated and characterized for the first time from *Z. chalybeum* and named as chalybemide C





**Figure 10:**(a) HMBC, ROESY and COSY correlations on 2D structure while 10(b) is 3D ROESY correlations for compound 41

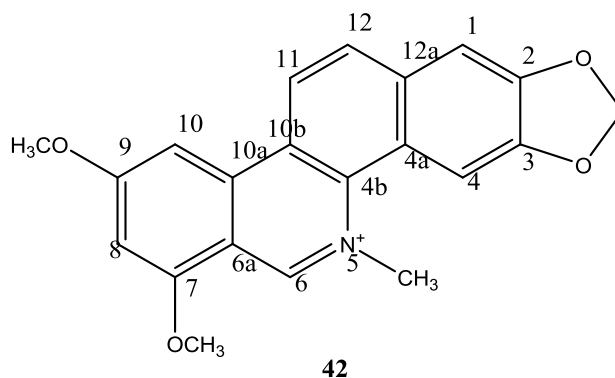
**Table 3:**  $^1\text{H}$  (600MHz) and  $^{13}\text{C}$ NMR (150 MHz) ( $\text{CDCl}_3$ ) spectral data for Chalybemide C

Position	$\delta\text{H}$ , mult ( $J$ , Hz)	$\delta\text{C}$	HMBC	COSY
1		168.1(C)		
2		122.2(C)		
3		159.8(C)		
4	6.89 dd ( $J = 8.3, 1.2$ )	112.2(CH)		
5	7.52 m	132.4(CH)		
6	7.05 t ( $J = 7.7$ )	122.5(CH)		
7	7.86 dd ( $J = 8.3, 1.8$ )	132.1(CH)	C-1	
1'		131.6(C)		
2'	7.15 d ( $J = 7.2$ )	127.1(CH)	C-7'	$\text{H}_2'-\text{H}_7'$
3'	6.86 d ( $J = 7.2$ )	113.2(C)		
4'		158.2(C-O)		
5'	6.86 d ( $J = 7.2$ )	113.2(CH)		
6'	7.15 d ( $J = 7.2$ )	127.1(CH)		$\text{H}_6'-\text{H}_7'$
7'	3.91 br s	57.1(CH <sub>2</sub> )	C-1	
5-OCH <sub>3</sub>	3.73 br s	56.4 (CH <sub>3</sub> )	C-5	
4'-OCH <sub>3</sub>	3.72 br s	56.6 (CH <sub>3</sub> )	C-4'	
NCH <sub>3</sub>	3.19 s	56.6 (CH <sub>3</sub> )	C-1,7'	

#### 4.1.4{1,3-dimethoxy-12-methyl-9H-[1,3]benzodioxolo[5,6-c]phenanthridin-12-ium}

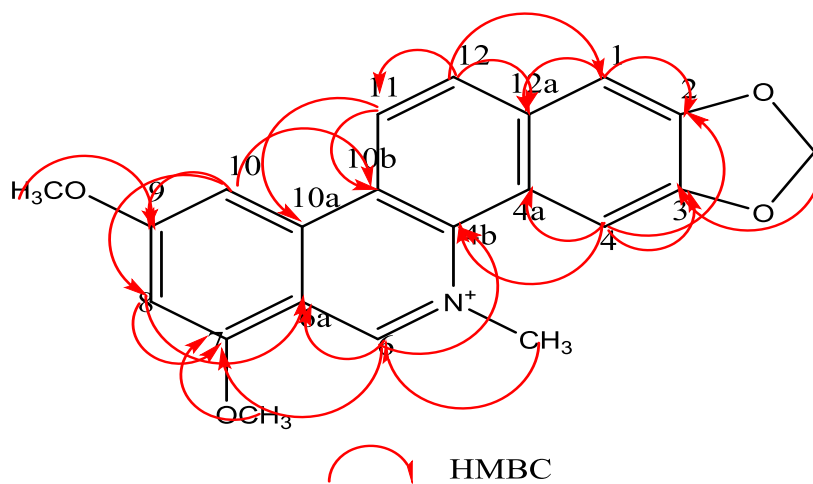
##### Phenanthridine A

Compound (**42**) was obtained as yellow needle crystals. Its ESI-MS (positive mode) displayed a molecular ion peak at  $m/z$  349.2795 corresponding to a molecular formula  $C_{21}H_{18}NO_4^+$ , ascribed to thirteen degrees of unsaturation. According to  $^1H$  NMR spectrum, an *ortho* coupled protons at  $\delta_H$  8.28 (1H, d,  $J=7.2$  Hz, H-11) and  $\delta_H$  8.28 (1H, d,  $J = 7.2$  Hz, H-12), *meta* coupled protons at  $\delta_H$  8.828 (1H, d,  $J = 2$  Hz, H-8) and  $\delta_H$  8.844 (1H, d,  $J = 2.0$  Hz, H-10), singlets at  $\delta_H$  7.781 (1H, s H-1),  $\delta_H$  8.322 (1H, s, H-4) and 10.10 (1H, s) indicated the presence of seven aromatic protons characteristics of a benzophenanthridine alkaloid skeleton. A methylene singlet at  $\delta_H$  6.357 (2H, s, 2, 3-OCH<sub>2</sub>O) characteristic of a methylenedioxy, two methoxy singlets with  $\delta_H$  4.12 (s, 7-OCH<sub>3</sub>),  $\delta_H$  3.86 (s, 9-OCH<sub>3</sub>) and a downfield signal (5.77 s, 3H) commonly associated with NCH<sub>3</sub> group.



$^{13}C$  NMR (Table 4) spectrum showed twenty one carbon signals which were classified as three oxygenated carbons (one methylenedioxy,  $\delta_C$  102.8 (OCH<sub>2</sub>O) and methoxy groups,  $\delta_C$  61.93 (9-OCH<sub>3</sub>) and  $\delta_C$  58.68 (7-OCH<sub>3</sub>), ten quaternary carbons in the aromatic region methyl amine and seven aromatic methines. The  $^1H$  and  $^{13}C$  NMR data were then reconciled by the HMQC proton-carbon attachment observed between:  $\delta_H$  7.781 (1H, s) and  $\delta_C$  106.1;  $\delta_H$  8.322 (1H, s) and  $\delta_C$  131.7;  $\delta_H$  10.10 (1H, s) and  $\delta_C$  151.3;  $\delta_H$  8.828 (1H, d,  $J=1.6$ ) and  $\delta_C$  119.3;  $\delta_H$  8.844 (1H, d,  $J=2.0$ ) and 119.5;  $\delta_H$  8.28 (1H, d,  $J=7.2$ ) and  $\delta_C$  105.4;  $\delta_H$  8.28 (1H, d,  $J=7.2$ ) and  $\delta_C$  126.3. These 1D data would closely predict four aromatic skeleton of a benzophenanthridine with N-CH<sub>3</sub>, a methylenedioxy and two methoxy groups (Miao *et al*, 2010) except for the position of the methoxy groups. The difference was noted with presence of a *meta* coupled protons at  $\delta_H$  8.83 (1H, d,  $J=1.6$ , H-8) and  $\delta_H$  8.84 (1H, d,  $J=2.0$ , H-10) instead of a pair of *ortho* coupled protons as observed with chelerythrine. This was supported by HMBC correlation between methoxy protons 4.12 to 142 C-7 and 3.89 to 145 (C-9), which were noted to have  $^2J$  correlation with (H-10) and (H-8),

respectively as shown in Figure 10 (appendix 4.4). Additional HMBC ( $^2J$  and  $^3J$ ) correlations between  $\delta_H$  5.077 (s) and these two carbon signals  $\{\delta_C$  130.8 (C-4b),  $\delta_C$  151.3 (C-6) $\}$  predicted the presence of a methyl amine with nitrogen at position 5. The spectra also confirmed the presence of two methoxy groups at carbon position 7 and 9 based on HMBC correlations between  $\delta_H$  8.828 (1H, d,  $J=1.6$ , H-8) and  $\delta_C$  58.68 (9-OCH<sub>3</sub>) and  $\delta_C$  61(7-OCH<sub>3</sub>), respectively (Figure 11B). These chemical shifts characterized chelerythrine derivative 1,3-dimethoxy-12-methyl-9H-[1,3]benzodioxolo[5,6-c]phenanthridin-12-ium named as phenanthridine A.



**Figure 11(b): Compound 42 and its HMBC**

**Table 4: <sup>1</sup>H and <sup>13</sup>CNMR (DMSO-d<sub>6</sub>) spectral data for Phenanthridine A.**

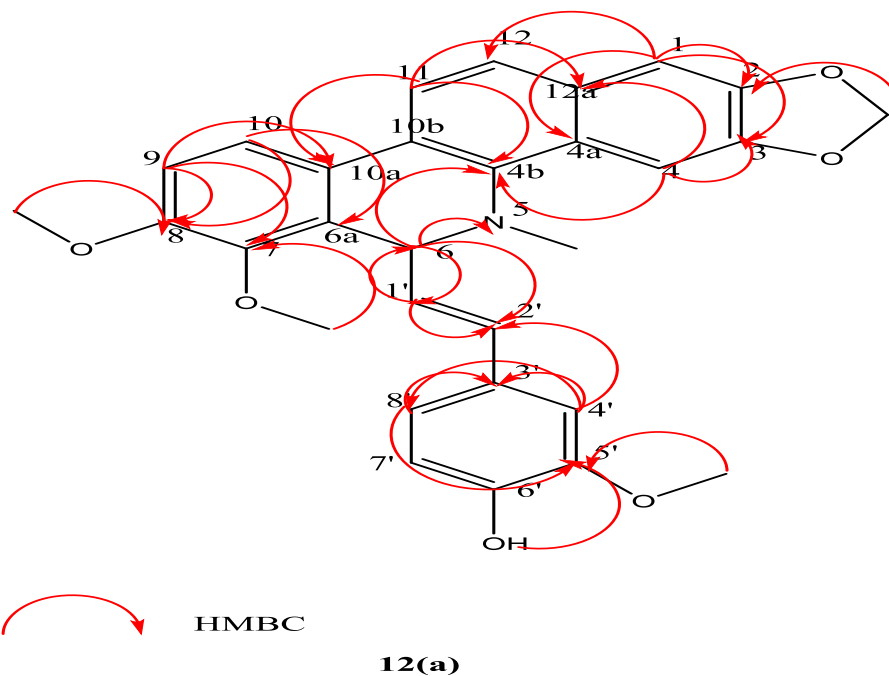
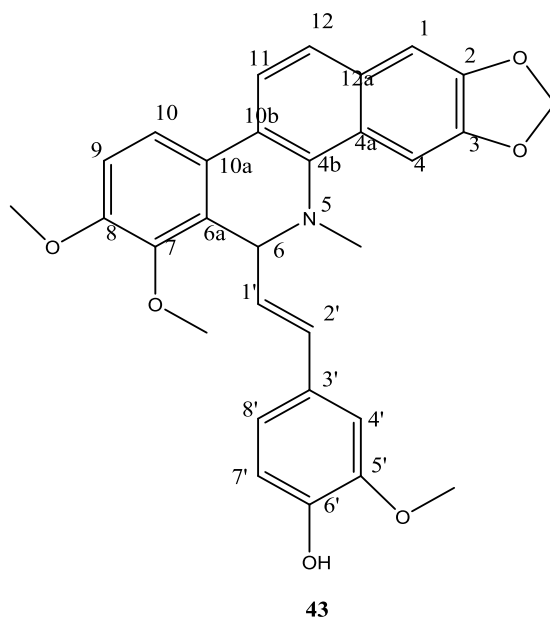
<b>C-Position</b>	<b>δ<sub>H</sub>, multi</b>	<b>δ<sub>C</sub></b>	<b>HMBC (<sup>3</sup>J)</b>
<b>1</b>	7.781 (1H, s)	106.1	C-2, C-12a, C-6a
<b>2</b>	-	147.2	-
<b>3</b>	-	147.0	-
<b>4</b>	8.322 (1H, s)	131.7	C-2, C-4a, C-6a, C-10b
<b>4a</b>	-	125.3	-
<b>4b</b>	-	130.6	-
<b>6</b>	10.10 (1H, s)	151.3	NCH <sub>3</sub> , C-2, C-4, C-12a, C-4b
<b>6a</b>	-	120.7	-
<b>7</b>	-	151.2	-
<b>8</b>	8.828 (1H, d, <i>J</i> = 1.6)	119.3	9-OCH <sub>3</sub> , 7-OCH <sub>3</sub>
<b>9</b>	-	144.3	-
<b>10a</b>	-	128.5	-
<b>10</b>	8.844 (1H, d, <i>J</i> = 2.0)	119.5	C-6, C-7, C-4, C-10a, C-12, C-12a
<b>10b</b>	-	118.8	-
<b>11</b>	8.28 (1H, d, <i>J</i> =7.2)	105.4	C-10b, C-10a, C-10b, C-4a,C-4b
<b>12a</b>	-	130.8	-
<b>12</b>	8.28 (1H, d, <i>J</i> = 7.2)	126.3	C-10a, C-10b, C-4a, C-4b
<b>2,3-OCH<sub>2</sub>O</b>	6.357 (2H, s)	102.8	C-3
<b>7-OCH<sub>3</sub></b>	4.117 (s)	61.93	C-7
<b>9-OCH<sub>3</sub></b>	3.86 (s)	58.68	C-9
<b>NCH<sub>3</sub></b>	5.077 (s)	52.24	C-4b, C-6

#### 4.1.5 Zanthocapsine (43) {(R) (1'E)-5'-methoxy-6'-hydroxy-cinnamyl-(1' → 6)-dihydrochelerythrine

Compound (43), named zanthocapsine, was isolated as a white crystal with melting point of 141-142°C. Its HREIMS indicated a molecular ion at  $m/z$  497.1839, which was in showing molecular formula  $C_{30}H_{27}NO_6$  (calcd for  $C_{30}H_{27}NO_6$ , 497.1838), with eighteen degrees of unsaturation. Analysis of the  $^1H$  NMR spectral data (Table 5) showed three sets of aromatic protons alongside other protons including: two pairs of *ortho*-coupled protons signals at  $\delta_H$  7.80 (1H, d,  $J = 8.4$  Hz, H-11) and 7.51 (1H, d,  $J = 8.4$  Hz, H-12); 7.68 (1H, d,  $J = 8.4$  Hz, H-10) and 7.13 (1H, d,  $J = 8.4$  Hz, H-9), corresponding to two 1,2,3,4-tetrasubstituted benzene rings; a second set of aromatic protons signals at  $\delta_H$  6.53 (1H, d,  $J = 8.0$  Hz, H-7'), 6.43 (1H, br d,  $J = 8.0$  Hz, H-8'), and 6.67 (1H, br s, H-4') predicted a 1,3,4-trisubstituted benzene ring supported by the COSY correlation between the *ortho* coupled protons; a third aromatic moiety was predicted by the presence of two singlets at  $\delta_H$  7.62 (1H, s, H-4) and  $\delta_H$  7.29 (1H, s, H-1), indicating a 1,2,4,5-tetrasubstituted aromatic ring.

Other protons signals noted on compound 43 included  $\delta_H$  6.13 (2H, d,  $J = 3.2$  Hz) for methylenedioxy H-atoms;  $\delta_H$  5.15 (1H, d,  $J = 4.0$  Hz, H-6) for methine adjacent to quaternary nitrogen; two aromatic methoxyl groups signals at  $\delta_H$  3.89 (3H, s) and 3.82 (3H, s) and an *N*-methyl group signal at  $\delta_H$  2.60 (3H, s). Simultaneously, evaluation of the  $^{13}C$  NMR, DEPT (Table 5) and HSQC spectra (appendix 5.4), thirty carbon signals were categorized as one *N*-methyl, one methylenedioxy, three methoxyl, twelve methines (one  $sp^3$  and eleven  $sp^2$ ), and thirteen  $sp^2$  quaternary carbons (including six oxygenated carbons atoms). These data suggested a 6-substituted dihydrochelerythrine skeleton, with exception of the signal for *trans*-oriented protons at  $\delta_H$  5.99 (1H, d,  $J = 16.0$  Hz, H-2') and 5.92 (1H, dd,  $J = 16.0, 4.0$  Hz, H-1') which showed HMQC contours to  $\delta_C$  129.72 and 127.04, respectively.

The appearance of the *trans*-oriented olefinic moiety was further confirmed to have mutual HMBC correlation with carbon signals at  $\delta_C$  127.75 and 146.26 for C-3' and C-6' on the 1,3,4-trisubstituted benzene ring which was in turn confirmed by HMBC (appendix 5.5) to hold one -OH at C-4' and  $OCH_3$  at C-5' thus pointing to the presence of a cinnamonyl unit in compound 43. Unambiguous confirmation of 5'-methoxy-6'-hydroxy-cinnamonyl moiety was based on a NOE cross-peak between the aromatic proton H-4' and a methoxyl singlet at  $\delta_H$  3.64. HMBC between the proton signal at  $\delta_H$  5.15 (6-H) to both olefinic carbons ( $\delta_C$  127.04 and 129.7) which in turn correlated to 6a (126.46) and 4b (140.06), provided the basis for asserting the 5'-methoxy-6'-hydroxy-cinnamonyl was attached at the C-6 of the dihydrochelerythrine skeleton.



**Figure 12B: Structure of compound 43 and its HMBC correlations**



**Table 5: <sup>1</sup>H and <sup>13</sup>CNMR (DMSO-d<sub>6</sub>) spectral data for Zanthocapsine (43)**

Position	<i>Spectral data for 43</i>		<i>Literature data (Luo et al., 2012)</i>		
	$\delta_H$ (J in Hz) mult	$\delta_C$	HMBC	$\delta_H$ (J in Hz) mult	$\delta_C$
1	7.29 (1H, s)	104.3	2,3,4a,12	7.29 (s)	104.3
2		147.24			147.2
3		147.91			147.9
4	7.62 (1H, s)	99.76	2,3,4b,12a	7.62 (s)	99.8
4a		126.89			126.9
4b		140.06			140.1
6	5.15 (1H, d, J = 4.0)	58.44	4b,6a,1',2', N-Me	5.15 (d, 4.0)	58.4
6a		126.46			126.5
7		145.71			145.7
8		151.97			152
9	7.13 (1H, d, J = 8.4)	111.97	7,8,10a	7.13 (d, 8.4)	112
10	7.68 (1H, d, J = 8.4)	119.14	8,6a,10b	7.68 (d, 8.4)	119.1
10a		124.21			124.2
10b		123.44			123.4
11	7.80(1H, d, J = 8.4)	119.96	4b,10a,12a	7.80 (d, 8.4)	120
12	7.51 (1H, d, J = 8.4)	123.66	1,4a,10b	7.51 (d, 8.4)	123.7
12a		130.5			130.5
1,	5.92 (1H, dd, J=16.0,4.0)	127.04	6,2',3'	5.92 (dd, 16.0, 4.0)	127
2'	5.99 (1H, d, J = 16.0)	129.72	6,1',3',4',8'	5.99 (d, 16.0)	129.7
3'		127.75			127.8
4'	6.67 (br s)	109.42	2',3',6',8	6.67 (br s)	109.4
5'		147.6			147.6
6'		146.26			146.3
7'	6.53(1H, d, J = 8.0)	115.3		6.53 (d, 8.0)	115.3
8'	6.43(1H, d, J = 8.0)	119.46	3',5'	6.43 (br d, 8.0)	119.5
OCH <sub>2</sub> O	6.13 (2H,s)	101.24	2',4',6'	6.13 (s, 3.2)	101.2
NCH <sub>3</sub>	2.60 (3H, s)	42.26	2,3	2.60 (s)	42.3
7-OCH <sub>3</sub>	3.82 (3H, s)	60.48	6,4b	3.82 (s)	60.5
8-OCH <sub>3</sub>	3.89 (3H, s)	55.7	7	3.89 (s)	55.7
5'-OCH <sub>3</sub>	3.64 (3H, s)	55.49	8	3.64 (s)	55.5
6'-OH	8.97 (br s)		5'	8.97 (br s)	

Further analysis of HMBC spectrum showed the following correlation: contours intersection 6.13 signal for the methylenedioxy protons and carbons signal at  $\delta_C$  147.24 and 147.91 for C-2 and C-

3, respectively which in turn correlated to proton signal at  $\delta_{\text{H}}$  7.62 (1H, s, H-4) and  $\delta_{\text{H}}$  7.29 (1H, s, H-1), thus confirming the 1,2,4,5-tetrasubstituted benzene ring (Figure 12B). Following the comparable spectroscopy of compound **43** and that of previously isolated phenyl substituted dihydrobenzophenanthridine alkaloid from *Zathoxylum capens* (Fotie et al., 2007) hence characterized as zanthocapsine.

#### 4.1.6 Zanthocapsol (**44**) {2-(3',4'-methylenedioxyphenyl)-5-[(*E*)-3''-hydroxy-1''-propenyl]-7-methoxybenzofuran}

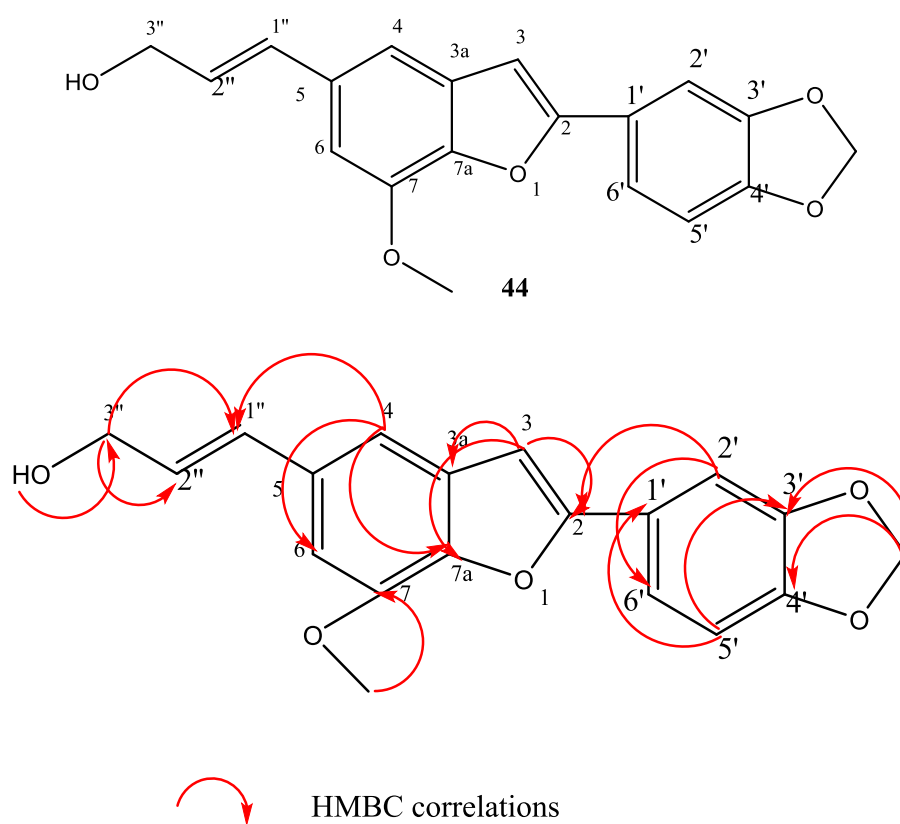
Zanthocapsol (**44**) was obtained as a pale yellow amorphous solid with the melting point 191-192°C with a molecular formula  $\text{C}_{19}\text{H}_{16}\text{O}_5$  predicted by both 1D and 2D NMR data and HREIMS at a molecular ion of  $m/z$  324.099 (calculated. for  $\text{C}_{19}\text{H}_{16}\text{O}_5$ , 324.0998) predictive of twelve degrees of unsaturation.

The  $^{13}\text{C}$  NMR spectrum of compound **44** indicated nineteen carbon signals, showing two oxygenated methylenes (one methylenedioxy,  $\delta_{\text{C}}$  101.51), one methoxyl group ( $\delta_{\text{C}}$  55.84), eight  $\text{sp}^2$  quaternary carbons (five oxygenated) and eight  $\text{sp}^2$  methines (Table 6). According to  $^1\text{H}$  NMR (DMSO- $d_6$ ) spectrum (Table 6), two broad singlets in the aromatic region at  $\delta_{\text{H}}$  7.18 (1H, br s) and 7.04 (1H, br s), signified a tetra substituted benzene ring moiety); and a tri-substituted aromatic ring moiety displaying an ABX spin systems constituted by a *meta*-coupling and *ortho* coupling ( $\delta_{\text{H}}$  7.40 (1H, br d,  $J = 8.0, 1.5\text{Hz}$ ); 7.05 (1H, d,  $J = 8.0\text{ Hz}$ ) and  $\delta_{\text{H}}$  7.44 (1H, dd,  $J = 1.5\text{Hz}$ )). Presence of a singlet at  $\delta_{\text{H}}$  7.25 (1H, s) attached to carbon signal at 101.38 with  $^3J$  HMBC correlation to aromatic carbon signal  $\delta_{\text{C}}$  111.18 holding a singlet proton ( $\delta_{\text{H}}$  7.18) and another quaternary aromatic carbon  $\delta_{\text{C}}$  142.61 indicated an ethenyl unit as one of the substituents of the tetrasubstituted benzene ring.

Moreover, the presence of a *trans*-3-hydroxy-1-propenyl group was noted based on the signals for *trans* coupled alkene protons ( $\delta_{\text{H}}$  6.61 (1H, d,  $J = 16.0\text{ Hz}$ ), 6.38 (1H, dt,  $J = 16.0, 4.8\text{ Hz}$ )) which showed HMBC correlation to an oxymethylene carbon holding two hydroxymethylene protons at  $\delta_{\text{H}}$  4.14 (2H, dd,  $J = 5.0, 4.8\text{ Hz}$ ) which in turn showed  $^3J$  correlation to alkene carbon  $\delta_{\text{C}}$  129.8. The attachment of the *trans*-3-hydroxy-1-propenyl to the tetrasubstituted aromatic rings was justified based on HMBC correlation between the proton signal at  $\delta_{\text{H}}$  6.61 and carbon signals at  $\delta_{\text{C}}$  111.18 and 104.78 (carbon atoms holding the singlet protons). The other two substituents were noted as a methoxyl group and phenyl ether linkage. The methoxy group was based on the HMBC correlation between the methoxy signals at  $\delta_{\text{H}}$  3.98 and carbon signal at  $\delta_{\text{C}}$  144.77 which in turn correlated with singlet protons at  $\delta_{\text{H}}$  6.03. Whereas the phenyl ether linkage was noted as

cyclization point with the ethene substituent to a furan group following the HMBC correlation of the alkene proton ( $\delta_{\text{H}}$  7.255) to both *O*-substituted aromatic carbon ( $\delta_{\text{C}}$  142.61) and *O*-substituted alkene carbon ( $\delta_{\text{C}}$  155.50). A benzofuran skeleton with *trans*-3-hydroxy-1-propenyl and methoxyl group was thus noted with a clear indication of substitution at carbon 2.

Considering the ABX substitution pattern of the other aromatic ring that would attach at carbon 2, a methylenedioxy group  $\{\delta_{\text{H}}$  6.101 (2H, s) $\}$  on carbon 3' and 4' automatically fitted the structure. It was thus concluded that the compound **44** named 2-(3',4'-methylenedioxyphenyl)-5-[(*E*)-3''-hydroxy-1''-propenyl]-7-methoxybenzofuran was a derivative of 2-arylbenzofuran neolignan, class of *Zanthoxylum* metabolite previously reported from *Zanthoxylum capense* (Luo *et al.*, 2013



**Figure 13: Structure of Compound 44 and its HMBC correlations**

**Table 6: <sup>1</sup>H and <sup>13</sup>C NMR (DMSO-d<sub>6</sub>) spectral data for Zanthocapsol (44)**

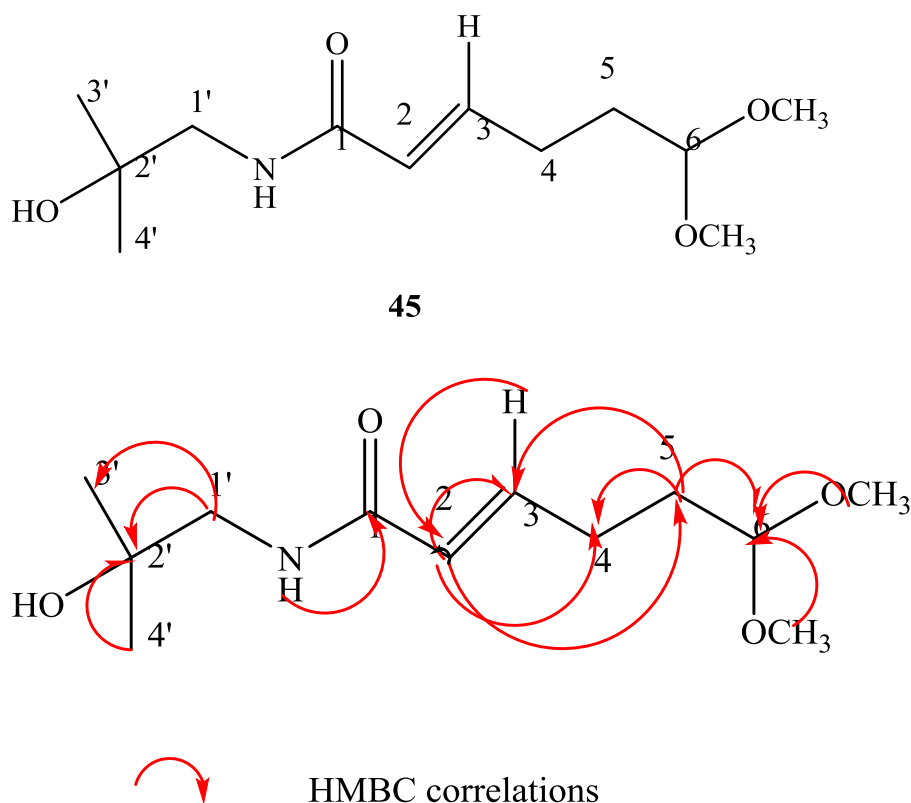
<i>Spectral data for Zanthocapsol (44)</i>				<b>Literature data (Luo X et al,2012)</b>	
<b>Position</b>	$\delta_{\text{H}}$ , multi	$\delta_{\text{C}}$	<b>HMBC</b>	$\delta_{\text{H}}$ , multi	$\delta_{\text{C}}$
<b>2</b>		155.5			155.5
<b>3</b>	7.255 (1H, s)	101.38	2,3a,7a	7.25 (s)	101.3
<b>3a</b>		130.66			130.7
<b>4</b>	7.181 (1H, br s)	111.18	3,6,7a,1''	7.18 (br s)	111.2
<b>5</b>		133.45			133.5
<b>6</b>	6.035 (1H, br s)	104.78	4,7,7a,1''	7.03 (br s)	104.8
<b>7</b>	7.442 (1H, br s)	144.72			144.7
<b>7a</b>		142.61			142.6
<b>1'</b>		123.85			123.8
<b>2'</b>	7.053 (1H, d, <i>J</i> = 8.0)	105.04	2,3',4',6'	7.44 (br s)	105
<b>3'</b>	7.403 (1H, d, <i>J</i> = 8.0)	148			148
<b>4'</b>	6.612 (1H, d, <i>J</i> = 16.0)	147.94			147.9
<b>5'</b>	6.376(1H, dt, <i>J</i> = 16.0, 4.8)	108.88	1',3',4'	7.05 (d, 8.0)	108.9
<b>6'</b>	4.149 (2H, dd, <i>J</i> = 5.0, 4.8)	118.84	2,2',4'	7.41 (br d, 8.0)	118.8
<b>1''</b>	6.101 (2H, s)	129.01	4,6,2'',3''	6.61 (d, 16.0)	129
<b>2''</b>	3.980 (3H, s)	129.81	5,1'',3''	6.39 (dt, 16.0, 4.8)	129.8
<b>3''</b>	4.884 (1H, s)	61.62	1'',2''	4.13 (dd, 5.0, 4.8)	61.6
<b>OCH<sub>2</sub>O</b>		101.51	3',4'	6.10 (s)	101.5
<b>7-OMe</b>		55.84	7	3.98 (s)	55.8
<b>3''-OH</b>			3''	4.88 (t, 5.0)	

#### 4.1.7 Zanthoamide D (45) {(*E*)-*N*-(2-Hydroxy-2-methylpropyl)-6, 6-dimethoxyhex-2-enamide}

Compound **45** was isolated as white crystals from the alkaloid fraction of the crude extract portion using hexane: DCM 1:9 solvent system for crystallization and showed a blue spot when sprayed with with *p*-anisaldehyde. The <sup>1</sup>H NMR (Table 7) showed two protons signals characteristic of alkene protons at  $\delta_{\text{H}}$  6.85 (1H, d, *J* = 15.0) and  $\delta_{\text{H}}$  6.08 (1H, d, *J* = 15.0) with *trans* orientation, three methylene protons  $\delta_{\text{H}}$  1.80 (2H, m),  $\delta_{\text{H}}$  2.24 (2H, t),  $\delta_{\text{H}}$  3.15 (2H, s), one methine proton  $\delta_{\text{H}}$

4.47 (1H, t), two methyl signals integrating to 6H ( $\delta_{\text{H}}$  1.17, 3H, s) and proton signals characteristic of methoxy groups  $\delta_{\text{H}}$  4.47 (1H, s). The  $^{13}\text{C}$  NMR (Table 7) revealed a total of twelve carbons; two quaternary carbons at  $\delta_{\text{C}}$  169.5 and 71.6, two  $\text{sp}^2$  methine carbons at  $\delta_{\text{C}}$  125.0 and 147.2, three methylene carbons at  $\delta_{\text{C}}$  33.6, 29.0 and 53.6, one methine carbon at  $\delta_{\text{C}}$  107.2, two symmetric methyl carbons at  $\delta_{\text{C}}$  28.0 and two symmetric methoxy carbons at  $\delta_{\text{C}}$  55.5. The  $^{13}\text{C}$  NMR thus, supported the  $^1\text{H}$  NMR data based on the attached proton analysis (HMQC spectrum), with exception of a broad proton signal at  $\delta_{\text{H}}$  6.96 suggested being for N-H from an amide moiety.

HMBC correlation showed  $^2J$  and  $^3J$  correlations between the methine proton signal at  $\delta_{\text{H}}$  4.47 with the methylene carbon signal at  $\delta_{\text{C}}$  33.6; methylene proton at  $\delta_{\text{H}}$  1.80 with methylene carbon at  $\delta_{\text{C}}$  29.0; methylene proton at  $\delta_{\text{H}}$  2.24 with olefinic methine carbon at  $\delta_{\text{C}}$  125.0. Moreover,  $^3J$  correlation between methine proton signal at  $\delta_{\text{H}}$  6.85 and the quaternary carbon at  $\delta_{\text{C}}$  169.4 and methine carbon at  $\delta_{\text{C}}$  125.0 revealed the connectivity between C-1 to C-6 with acetal moiety at C-6  $\text{O}=\text{C}-\text{C}=\text{C}-\text{CH}_2-\text{CH}_2-\text{C}(\text{OCH}_3)_2$ . On the other hand, methylene proton at  $\delta_{\text{H}}$  3.15 showed a correlation with carbon signal at  $\delta_{\text{C}}$  28.0 and 71.6 which confirmed connectivity from C-1' to C-3' as  $-\text{HN}-\text{CH}_2-\text{C}\{(\text{OH})(\text{CH}_3)_2\}$ . The data was confirmed by ESIMS a molecular peak at  $m/z$  268.1322, thus characterizing compound **45** as a derivative of 6,6-dimethoxy hex-2-enamide isolated for the first time from *Z. gilletii* named as Zanthoamide D



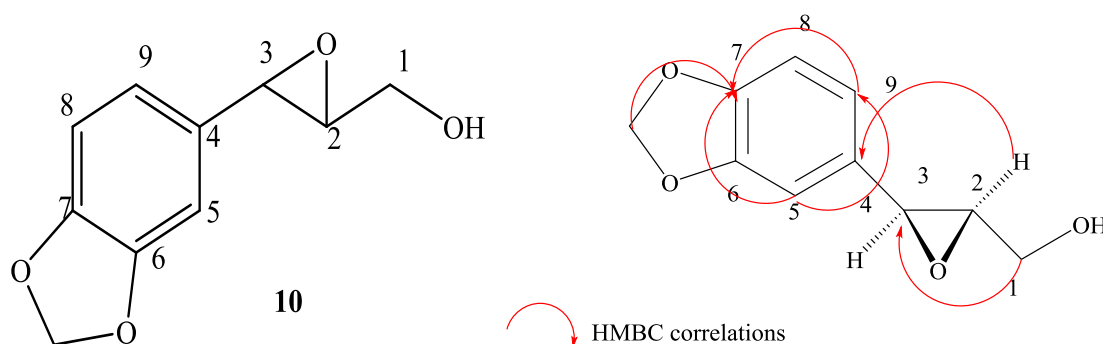
**Figure 14: Structure of compound 45 and its HMBC correlations**

**Table 7: <sup>1</sup>H and <sup>13</sup>C NMR (DMSO-d<sub>6</sub>) spectral data for Zanthoamide D**

C. position	δ <sub>H</sub> , Multiplicity	δ <sub>C</sub>	<sup>1</sup> H- <sup>1</sup> H	HMBC
<b>1</b>	-	169.5	-	-
<b>2</b>	6.08, 1H dt, (J=15.0)	125.0	6.85, 2.24	C-5, C-3, C-1
<b>3</b>	6.85, 1H, dt (J=15.0)	147.2	6.08	C-1, C-2, C-4
<b>4</b>	2.24, 2H td	29.0	1.80, 6.85	C-6, C-5
<b>5</b>	1.80, 2H m	33.6	2.24, 4.47	107.2, 29.0
<b>6</b>	4.47, 1H t	107.2	1.80	C-5
<b>1'</b>	3.15, 2H s	53.6	-	C-3', C-2'
<b>2'</b>	-	71.6	-	-
<b>3'/4'</b>	1.17, 3H s	28.08	-	C-1', C-2'
<b>2 x OCH<sub>3</sub></b>	3.25, 3H	55.5	-	-

#### 4.1.8 2,3-epoxy-6,7-methylenedioxy conferyl alcohol (10)

Compound 10 was isolated as a white crystal (melting point: 124-125°C) with a molecular formula of C<sub>10</sub>H<sub>10</sub>O<sub>4</sub> as predicted by both 1D and 2D NMR and The <sup>1</sup>H NMR spectrum (CDCl<sub>3</sub>, Table 8) corresponded to that of 2,3-epoxy-6,7-methylenedioxy conferyl alcohol isolated from *Z. chalybeum* by Wasiali (2019).



**Figure 15: Compound 10 and its HMBC**

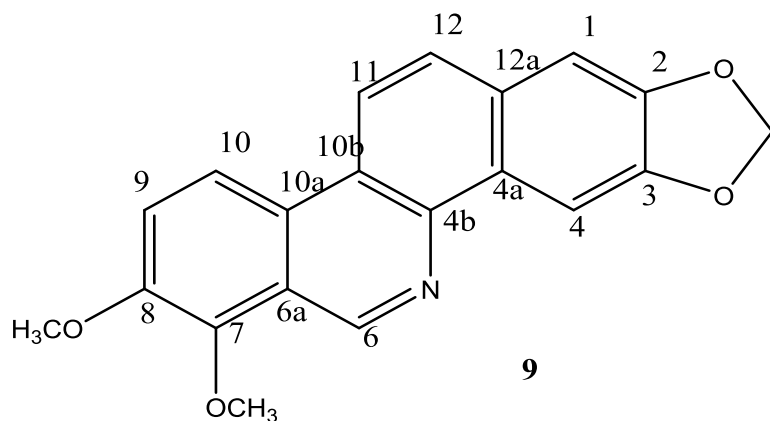
**Table 8: <sup>1</sup>H and <sup>13</sup>C NMR (CDCl<sub>3</sub>) spectral data for 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol**

C-Position	$\delta_C$	$\delta_H$ , Multiplicity	COSY ( <sup>1</sup> H- <sup>1</sup> H)	HMBC	Literature data (Anza <i>et al.</i> , 2014)	
					$\delta_H$ , Multiplicity	$\delta_C$
1	71.7	3.89 (1H, d, <i>J</i> = 16.0, 7.2) 4.25 (1H, dd, <i>J</i> = 16.0, 13.0)	H <sub>1,1'</sub> -H <sub>2</sub>	C-2,3	4.25 dd(1H, d, <i>J</i> = 16.0, 7.2) 3.89 dd(1H, dd, <i>J</i> = 16.0, 13.0)	71.7
2	54.3	3.05(m)	H <sub>2</sub> -H <sub>1,3</sub>	C-1, 3, 4	3.10 m	54.3
3	85.7	4.73(1H, d, <i>J</i> = 4.4)	H <sub>3</sub> - H <sub>2</sub>	C-1,2,4,5,9	4.74 d(1H, d, <i>J</i> = 4.4)	85.9
4	135.0	-	-	-	-	135.1
5	119.3	6.82 (1H, dd, <i>J</i> = 0.8,1.2)	-	C-3,4,6,9	6.82 dd(1H, dd, <i>J</i> = 0.8,1.2)	119.4
6	147.8	-	-	-	-	147.9
7	147.1	-	-	-	-	147.1
8	108.1	6.86 (1H, dd, <i>J</i> = 8.0,0.8)	H <sub>8</sub> -H <sub>9</sub>	C-4,6,7	6.86 dd	108.2
9	106.5	6.80 (1H, <i>J</i> = 8.0, 1.2)	H <sub>9</sub> -H <sub>8</sub>	C-3,4,5,7	6.80 dd(1H, <i>J</i> = 8.0, 1.2)	106.5
OCH <sub>2</sub> O	101.3	5.95 (2H, s)	-	C-6,7	6.02 1H,s	101.1

#### 4.1.9 1,2-dimethoxy-[1,3]benzodioxolo[5,6-c]benzophenanthridine (Norchelerythrine) (9)

Compound **9** was isolated as yellow crystals, UV active with blue fluorescence with melting point of 210-212°C. Its molecular mass determined by ESIMS [M+H]<sup>+</sup>, *m/z* = 334 giving molecular formula as C<sub>20</sub>H<sub>15</sub>NO<sub>4</sub> indicating thirteen degrees of unsaturation. <sup>1</sup>H NMR spectrum showed two pairs of *ortho* coupled proton signals at  $\delta_H$  8.20 (d, *J* = 8.7),  $\delta_H$  8.59 (d, *J* = 8.7) and  $\delta_H$  8.62 (d, *J* = 9.7),  $\delta_H$  8.65 (d, *J* = 9.7) supported by COSY spectrum together with a deshielded proton peak predictive of an imine proton (HC=N) at  $\delta_H$  9.95 (Table 9).

<sup>13</sup>C spectrum revealed the presence of 17 sp<sup>2</sup> carbons, ten quaternary carbon atoms including four oxygenated carbons; methylenedioxy group at  $\delta_C$  104.5 and two methoxy groups at  $\delta_C$  57.7 and  $\delta_C$  63.0, typical of benzophenanthridine structure. These data and the literature were similar to that of norchelerythrine previously isolated from roots of *Z. gillettii* (Wafula, 2014) and from roots of *Z. chalybeum* (Wasiali, 2019).



**Table 9: <sup>1</sup>H and <sup>13</sup>CNMR (CDCl<sub>3</sub>) spectral data for Norchelerythrine (9)**

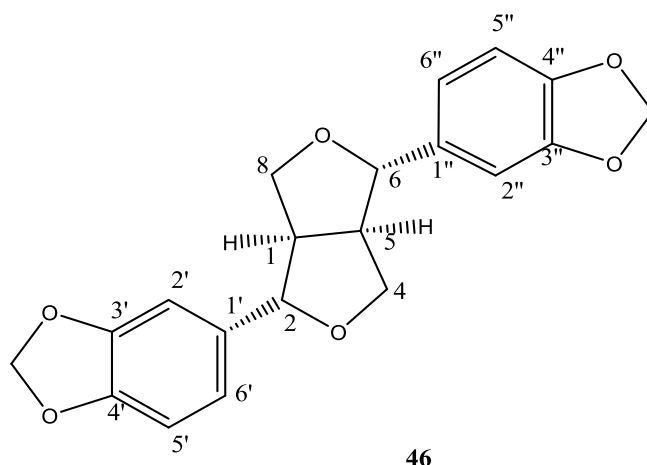
C-Position	Compound 9			Literature data (Maliwat <i>et al</i> , 1997)	
	$\delta_H$ , Multiplicity	$\delta_C$	HMBC	$\delta_H$ , Multiplicity	$\delta_C$
1	7.52 (1H, s)	107.2	C- 3, C-12	7.52 (1H, s)	104.4
2		152.2			149.4
3		151.9			148.5
4	8.18 (1H, s)	105.2	C-2, C- 4b	8.72(1H, s)	102.0
4a		133.5			129.7
4b		134.3			129.7
6	9.95 (1H, s)	151.9	C-6a, C- 10a	9.75 (1H, s)	140.0
6a		127.2			128.1
7		150.9			148.3
8		147.6			146.6
9	8.65 (1H, d, $J = 9.7$ Hz)	132.7	C- 6a, 8-OCH <sub>3</sub>	8.36 (1H, d, $J=9.09$ Hz)	129.2
10	8.62 (1H, d, ( $J = 9.7$ Hz)	120.0	C- 6a, 7-OCH <sub>3</sub> , 8-OCH <sub>3</sub>	7.59 (1H, d, $J=9.09$ Hz)	120.0
10a		121.8			121.9
10b		130.1			129.7
11	8.59 (1H, d, $J = 8.7$ Hz)	119.6	C-4b, 10a, C-10b	8.35 (1H, d, $J=8.74$ Hz)	118.7
12	8.20 (1H, d, $J = 8.7$ Hz)	127.5	C-1, C-12a	7.86 (1H, d, $J=8.74$ Hz)	127.1
12a		120.0			120.0
OCH <sub>2</sub> O	5.95 (2H, s)	101.3	C-6, C-7	6.13 (2H, s)	101.3
7-OCH <sub>3</sub>	4.28 (3H, s)	150.9	C-7	4.08 (3H,s)	61.9
8-OCH <sub>3</sub>	4.13 (3H, s)	147.6	C-8	4.13 (3H, s)	56.8

#### 4.1.10 Sesamine (46)

Compound 46 was isolated as white amorphous solid which is UV active with molecular mass of 121 - 122°C. Its molecular formula is C<sub>20</sub>H<sub>18</sub>O<sub>6</sub> predicted by both 1D and 2D NMR data and HREIMS at a molecular ion of  $m/z = 354$  (calculated. for C<sub>20</sub>H<sub>18</sub>O<sub>6</sub>, 354.35) giving twelve degrees of unsaturation. Its <sup>1</sup>H NMR and <sup>13</sup>C NMR spectral and its TLC spots together physical



properties was similar to those of a lignan, sesamine which previously isolated from *Z. budrunga* by Mukhlensur (2005) and from *Z. gillettii* by Wafula (2014).



**Table 10: <sup>1</sup>H and <sup>13</sup>CNMR (CDCl<sub>3</sub>) spectral data for sesamine(46)**

C-Position	$\delta_{\text{H}}$ , Multiplicity	$\delta_{\text{C}}$	Literature data (Ali <i>et al</i> , 2007)	
			$\delta_{\text{H}}$ , Multiplicity	$\delta_{\text{C}}$
<b>1</b>	3.11 (1H, m)	55.1	3.11 (1H, m)	55.3
<b>2</b>	4.65 (1H, d, $J = 4.8$ )	87.1	4.65 (1H, d, $J = 4.8$ )	87.1
<b>4</b>	3.82 (1H, m)	72.4	3.82 (1H, m)	72.3
	4.20 (1H, m)		4.20 (1H, m)	
<b>5</b>	3.09 (1H, m)	55.1	3.09 (1H, m)	55.0
<b>6</b>	4.70 d, $J = 5.1$	87.1	4.70 d, $J = 5.1$	87.1
<b>8</b>	3.83 m	72.4	3.83 m	72.3
	4.23 m		4.23 m	
<b>1'</b>		133.5		133.4
<b>2'</b>	6.80 d, $J = 8.1$	114.2	6.80 d, $J = 8.1$	114.1
<b>3'</b>		146.4		146.4
<b>4'</b>		145.7		145.6
<b>5'</b>	6.70 d, $J = 8.1$	116.2	6.70 d, $J = 8.1$	116.2
<b>6'</b>	6.69 dd, $J = 1.8, 8.1$	118.5	6.69 dd, $J = 1.8, 8.1$	118.5
<b>1''</b>		136.2		136.1
<b>2''</b>	6.87 d, $J = 1.5$	107.3	6.87 d, $J = 1.5$	107.2
<b>3''</b>		148.7		148.7
<b>4''</b>		148.7		148.7
<b>5''</b>	6.77 d, $J = 8.1$	108.7	6.77 d, $J = 8.1$	108.7
<b>6''</b>	6.84 dd, $J = 8.1, 1.5$	120.1	6.84 dd, $J = 8.1, 1.5$	120.1
<b>OCH<sub>2</sub>O</b>	5.91 (2H, s)	101.8	5.91 (2H, s)	102.0
<b>OCH<sub>2</sub>O</b>	5.90 (2H, s)	102.0	5.90 (2H, s)	102.0

#### 4.1.11 Skimmianine (21)

Compound 21 was isolated as yellow coloured powder with the melting point of 179-180°C with molecular ion peak at  $m/z$  260  $[M+H]^+$  corresponding  $C_{14}H_{13}NO_4$  supported by  $^1H$  and  $^{13}C$  NMR data (Table 11). The data was compared to the literature and marched exactly with that of skimmianine as was isolated from *Glycosmis pentaphylla* by Natarajan *et al*, 2020.

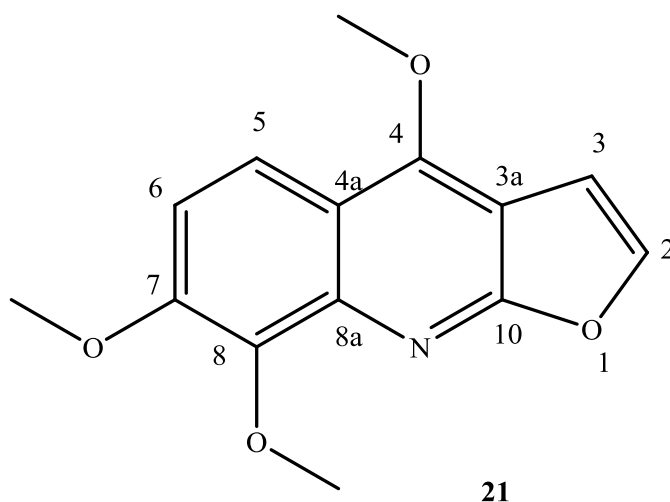


Table 11:  $^1H$  and  $^{13}C$ NMR (CDCl<sub>3</sub>) spectral data for Skimmianine (21)

C-Position	$\delta_H$ , Multiplicity	$\delta_C$	Literature data (Ali et al, 2007)	
	$\delta_H$ , Multiplicity	$\delta_C$	$\delta_H$ , Multiplicity	$\delta_C$
2	7.54, d, J=7.84	142.8	7.56, d, (J=8.0)	142.8
3	7.01, d, J=7.84	101.9	7.02, d, (J=8.0)	101.9
3a		104.3		104.4
4		152.0		152.1
4a		114.8		114.6
5	7.96, d (J= 8.0)	117.9	7.99, d, (J=8.0)	117.8
6	7.23, d J= 8.0)	112.2	7.23, d, (J=8.0)	112.2
7		157.0		157.0
8		164.2		164.4
8a		114.4		114.8
10		142.1		142.1
4-OCH <sub>3</sub>	4.03, s	56.7	4.03, s	56.7
7-OCH <sub>3</sub>	4.11,s	58.8	4.11, s	58.8
8-OCH <sub>3</sub>	4.41,s	61.5	4.41, s	61.4

#### 4.2. $\alpha$ -Amylase and $\alpha$ -Glucosidase inhibition of pure isolates from *Z. chalybeum* and *Z. gilletti*

The dose dependent  $\alpha$ -amylase and  $\alpha$ -glucosidase inhibitory activities of the isolated

compounds were processed on long probit analysis (regression analysis) to establish IC<sub>50</sub> values (Table 12). There was no significant difference in the activities ( $P > 0.05$ ) between the eight alkaloids compared to activity of acarbose (standard drug) with IC<sub>50</sub> = 42.67  $\mu$ M against  $\alpha$ -amylase. Two phenolic compounds; 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) and zanthocapsol (**44**) showed no significant difference ( $p > 0.05$ ) in the activity (IC<sub>50</sub> of 48.34 and 48.21 respectively) with acarbose while sesamine (**46**) displayed significantly low activity (IC<sub>50</sub> = 54.67  $\mu$ M) against  $\alpha$ -amylase enzyme relative to acarbose.

**Table 12: In vitro IC<sub>50</sub> values of compounds from *Z. chalybeum* and *Z. gillettii* against  $\alpha$ -amylase and  $\alpha$ -glucosidase inhibition compared to standard control inhibitor (acarbose)**

<i>Compounds</i>	$\alpha$ -amylase	$\alpha$ -glucosidase
	( $\mu$ M)	( $\mu$ M)
<i>Acarbose</i>	42.67	44.88
<i>Chaylbemide A (39)</i>	45.76	43.54
<i>Chaylbemide B (40)</i>	43.22	44.32
<i>Chaylbemide C (41)</i>	46.76	47.43
<i>Zanthoamide D (45)</i>	47.36	47.23
<i>Skimmianine (21)</i>	47.72	47.63
<i>Norchelerythrine (9)</i>	46.49	48.7
<i>Zanthocapsine (43)</i>	48.36	47.27
<i>Phenanthridine A (42)</i>	48.91	49.20
<i>Zanthocapsol(44)</i>	48.21	47.11
<i>2,3-Epoxy-6,7-methylenedioxy coniferyl (10)</i>	48.34	47.45
<i>Sesamine (46)</i>	54.67	54.77
<b>LSD (<math>P &lt; 0.05</math>)</b>	8.11	8.6
<b>CV%</b>	40	34

IC<sub>50</sub> values, calculated from regression using five different concentrations (10.0, 20.0,

50.0, 100.0, and 150.0  $\mu$ M) are expressed as mean of replicated experiments with percentage coefficient of variation (CV %) indicated, LSD ( $p < 0.05$ ): Least Significant

difference considered significant at  $p \leq 0.05$ .

The isolates showed a similar activity against  $\alpha$ -glucosidase enzyme as was (Table 12) indicating that all the isolated alkaloids and two phenolics (zanthocapsol (**44**) and 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**)) from the stem barks of the two plants had no significant difference in the inhibitory activity ( $p > 0.05$ ) compared to control acarbose but sesamine showed significantly ( $p < 0.05$ ) different compared to the control which implied that sesamine among phenolics displayed moderately lower activity against  $\alpha$ -glucosidase activities. The results from the assay deduced that inhibitory activity of the isolated alkaloids and slightly to the phenolic compounds from the plant *Z. chalybeum* and *Z. gillettii* thus agreeing with the previously established anti-hyperglycemic activities of *Z. chalybeum* stem and root extracts (Agwaya, 2016). Hydrogen bonding interaction between inhibitor and the enzyme had previously been cited as a possible inhibition mode involving active sites of  $\alpha$ -amylase and  $\alpha$ -glucosidase and the hydroxyl groups of compounds with phenolic entities (Bahadoran *et al*, 2013).

#### **4.3 In-vitro modes of $\alpha$ -amylase and $\alpha$ -glucosidase inhibition of active compounds from *Z. chalybeum* and *Z. gillettii***

Different inhibitors show varied modes of inhibition against enzymes. The modes of inhibition are predicted by considering Lineweaver-Burk plot, from which kinetic parameters are obtained such as maximum velocities ( $V_{max}$ ) and  $K_m$  (Menten constant; substrate concentration at which  $V_{max}$  is  $1/2$ ) are obtained to predict the modes of inhibition.

The modes of inhibition and the enzymes-inhibitor inhibition constants were deduced using kinetic analysis based on both Dixon plots (Appendix 12.0) and Lineweaver-Burk plots (Table 13). The results revealed that compounds showed mixed inhibitory modes. These compounds were Skimmianine (**21**), Norchelerythrine (**9**), Zanthocapsine (**43**), and Phenanthridine A (1,3-dimethoxy-12-methyl-9H-[1,3]benzodioxolo[5,6-c]phenanthridin-12-ium) (**42**) showing comparable inhibitor constant,  $K_i$  of to acarbose ( $p > 0.05$ ) which are lower indicating that it's difficult to dissociate enzyme-inhibitor complex hence better inhibitor against  $\alpha$ -amylase enzyme while the other six compounds which showed significantly ( $p < 0.05$ ) low activity compared to acarbose due their higher  $K_i$  values indicating enzyme-inhibitor complex bond easily dissociate as displayed by either noncompetitive or uncompetitive modes against  $\alpha$ -amylase actions on starch.

The compounds that exhibited mixed inhibition on amylase also exhibited comparable modes on  $\alpha$ -glucosidase activities. These compounds had considerably ( $p < 0.05$ ) higher  $K_i$  values than acarbose, indicating a better potential for inhibition against glucosidase than amylase.

On the other hand, the other compounds that exhibited uncompetitive and non-competitive inhibitions against  $\alpha$ -glucosidase activities had similar  $K_i$  values ( $p < 0.05$ ) in comparison to acarbose. One compound 2,3-Epoxy-6,7-methylenedioxy coniferyl (**10**) showed a competitive mode of inhibition both on amylase and glucosidase with  $K_i$  of  $5.54 \pm 0.58$  ( $R^2 = 0.985$ ) and  $17.21 \pm 0.15$  ( $R^2 = 0.8692$ ), respectively which were statistically comparable ( $p < 0.05$ ) to that of acarbose thus indicative of the most active inhibitors from the *Z. chalybeum* and *Z. gilletti* extracts. The most potent inhibitors of  $\alpha$ -amylase and  $\alpha$ -glucosidase were reported to be compound **10** and the alkaloids **45**, **21**, **9**, and **43**, which exhibit competitive and mixed inhibitions, respectively.

Conversely, the compounds **45**, **39**, **40**, **41**, **43**, and **46** exhibited different mechanisms of inhibition and corresponding dissociation constants in relation to the two enzymes, indicating moderate inhibitory activities. Alkaloids exhibit superior inhibitory activities against  $\alpha$ -amylase and  $\alpha$ -glucosidase possible due to H-bond interactions, while phenolics' effectiveness could be due to OH groups.

**Table 13: Inhibition modes and inhibition constant (K<sub>i</sub>) values for compounds from *Z. chalybeum* and *Z. gillettii* against  $\alpha$ -amylase and  $\alpha$ -glucosidase inhibition relative to acarbose inhibitions, as determined by Lineweaver-Burk plots and Dixon Plots, respectively**

<i>Compound</i>	$\alpha$ -amylase				$\alpha$ -glucosidase			
	Inhibition Mode	K <sub>m</sub>	V <sub>max</sub>	K <sub>i</sub> (mM)	Inhibition mode	K <sub>m</sub>	V <sub>max</sub>	K <sub>i</sub> (mM)
<i>Chaylbemide A (39)</i>	Non-competitive	2.55	0.041	13.36±2.43**	Non-competitive	8.47	0.044	20.95±2.14
<i>Chaylbemide B (40)</i>	Non-competitive	5.87	0.035	11.05±0.58**	uncompetitive	6.03	0.048	44.58±1.65
<i>Chaylbemide C (41)</i>	Non-competitive	6.35	0.034	14.83±0.50**	Non-competitive	6.86	0.044	17.56±0.24
<i>Zanthoamide D (45)</i>	Non-competitive	4.28	0.036	26.69±2.13**	Non-competitive	14.18	0.047	34.73±0.79**
<i>Skimmianine (21)</i>	Mixed	5.51	0.034	2.74±0.06	Mixed	9.58	0.044	7.64±0.02**
<i>Norchelerythrine (9)</i>	Mixed	8.14	0.044	7.57±0.59	Mixed	14.3	0.028	7.68±0.04**
<i>Zanthopensine (43)</i>	Mixed	12.11	0.033	3.34±0.03	Mixed	15.286	0.042	4.73±0.10**
<i>Phenanthridine A (42)</i>	Mixed	6.14	0.035	3.10±0.20	Mixed	11.92	0.044	9.17±0.10**
<i>Zanthocapsol (44)</i>	Uncompetitive	8.61	0.033	26.28±1.47**	uncompetitive	11.89	0.044	20.62±1.94
<i>2,3-Epoxy-6,7-methylenedioxy coniferyl (10)</i>	Competitive	3.77	0.04	5.54±1.02	competitive	3.76	0.05	17.21±0.16
<i>Sesamine (11)</i>	Uncompetitive	2.5	0.043	12.53±1.957**	Uncompetitive	5.17	0.05	24.33±1.93
<i>Acarbose</i>	Competitive	7.68	0.033	6.14±0.01	Competitive	4.59	0.043	22.40±1.23

\*\* values are significantly different from the standard inhibitor (acarbose) Inhibition constant (K<sub>i</sub>) based on Tukey HSD / Tukey Kramer post-analysis of one-way analysis of variance replicated K<sub>i</sub> values. Least significant difference was considered at p < 0.05

## CHAPTER FIVE

### SUMMARY, CONCLUSION AND RECOMEDATION

The study was intended to establish the modes of inhibition of alkaloids and phenolics from *Z. chalybeum* and *Z. gillettii* against  $\alpha$ -amylase and  $\alpha$ -glucosidase and validation of the rationale behind the use of *Z. chalybeum* and *Z. gillettii* in managing diabetes mellitus.

#### 5.1 Summary

Nine pure compounds were isolated from the stem bark of *Z. chalybeum*. Among them are seven alkaloids including two benzophenanthridine alkaloids; norchelerythrine (**9**) and phenanthridine A (1,3-dimethoxy-12-methyl-9H-[1,3]benzodioxolo[5,6-c]phenanthridin-12-ium) (**42**), four amides: chalybemide A (**39**), chalybemide B (**40**), chalybemide C (**41**) and one furoquinoline alkaloid: skimmianine (**21**), two phenolics: 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) and sesamine (**46**). Six compounds were isolated from *Z. gillettii* including four alkaloids among them are three benzophenanthridine alkaloids: Norchelerythrine (**9**), phenanthridine A (**42**) (chelerythrine derivative) and Zanthocapsine (**43**) and an amide alkaloid; zanthoamide D (**45**), and two phenolics zanthocapsol (**44**) and sesamine (**46**). This totals to thirteen pure compounds from the two plants; *Z. Chalybeum* and *Z. gillettii*.

All the eight alkaloids and three phenolics isolated from the two plants were assayed against the two enzymes:  $\alpha$ -amylase and  $\alpha$ -glucosidase. All the alkaloids, norchelerythrine (**9**), phenanthridine A (**42**), chalybemide A (**39**), chalybemide B (**40**), chalybemide C (**41**) zanthoamide D (**45**), skimmianine (**21**) and zanthocapsine (**43**) and two phenolics: 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) and zanthocapsol (**44**) showed no significant difference ( $p > 0.05$ ) in the activity compared to acarbose the standard drug but sesamine showed a significant difference in the activity compared to acarbose against  $\alpha$ -amylase. The same pattern was displayed in the activity against  $\alpha$ -glucosidase where all alkaloids and phenolics assayed showed no significant difference ( $p > 0.05$ ) in the inhibition compared to acarbose but sesamine showing relatively lower activity.

2,3-epoxy-6,7-methylenedioxy coniferyl (**10**) displayed competitive inhibition mode on both  $\alpha$ -amylase and  $\alpha$ -glucosidase with  $K_i$  of  $5.54 \pm 0.58$  ( $R^2 = 0.985$ ) and  $17.21 \pm 0.15$  ( $R^2 = 0.8692$ ), respectively. These were statistically comparable ( $p < 0.05$ ) to that of acarbose indicating the most active inhibitor from the *Z. chalybeum* and *Z. gillettii* extracts. Compound **10** and the alkaloids **42**, **21**, **9**, and **43** display competitive and mixed inhibitions, respectively were noted as the most potent inhibitors of both  $\alpha$ -amylase and  $\alpha$ -glucosidase. The other compounds **45**, **39**, **40**, **41**, **43**, and **46** showed varied modes of inhibition and associated dissociation constants towards the two enzymes (Table 11) which would be categorized as moderate inhibitory activities compared to acarbose.

## 5.2 Conclusion

- i. Alkaloids and phenolics that were isolated from *Z. Chalybeum* and *Z. gillettii* were Norchelerythrine (**9**), phenanthridine A, (**42**), chalybemide A (**39**), chalybemide B (**40**), chalybemide C (**41**) and Zanthoamide D (**45**) skimmianine (**21**), 2,3-epoxy-6,7-methylenedioxy coniferyl alcohol (**10**) sesamine (**46**) and two triterpenes: zanthocapsine (**43**) and zanthocapsol (**44**) of which phenanthridine A (**42**), chalybemide A (**39**), chalybemide B (**40**), chalybemide C (**41**) are reported for the first time while zanthocapsine (**43**) and zanthocapsol (**44**) are reported from *Z. gillettii* for the first time.
- ii. All isolated compounds showed no significant difference ( $P < 0.05$ ) in their inhibitory activities against  $\alpha$ -amylase and  $\alpha$ -glucosidase compared to acarbose ( $IC_{50} = 42.67$  and  $44.88$ , respectively) except sesamine (**46**) ( $IC_{50} = 54.67$  and  $54.77$ , respectively) which showed relatively lower activity.
- iii. 2,3-Epoxy-6,7-methylenedioxy coniferyl (**10**) showed a competitive mode of inhibition against both  $\alpha$ -amylase and  $\alpha$ -glucosidase with lower  $K_i$  of  $5.54 \pm 0.58$  ( $R^2 = 0.985$ ) and  $17.21 \pm 0.15$  ( $R^2 = 0.8692$ ), respectively showing the most active inhibitor against  $\alpha$ -amylase and  $\alpha$ -glucosidase together with compounds **42**, **21**, **9**, and **43** showed mixed mode of inhibition with relatively lower  $K_i$  values showing stronger binding and better inhibition. Compounds **45**, **39**, **40**, **41**, **43**, and **46** showed both noncompetitive and uncompetitive modes of inhibition with relatively higher  $K_i$  values compared to acarbose ( $p > 0.05$ ), indicating that they easily dissociate from inhibitor – enzyme complex due weak binding hence moderately lower inhibition against  $\alpha$ -amylase and  $\alpha$ -glucosidase activities.

## 5.3 Recommendation

- i. *Z. chalybeum* and *Z. gilletti* contain bio active compounds such as amide, quinoline and benzophenanthridine alkaloids and phenolics. The plants are recommended for in vivo studies.
- ii. The relatively active compounds recommended for further computational studies in order that their templates may be used for development of anti-hyperglycemic drugs.
- iii. *Z. chalybeum* and *Z. gilletti* contain bio active compounds such as amide, quinoline and benzophenanthridine alkaloids and phenolics which can further be evaluated for potential application in diabetic drug discovery



#### **5.4 Significance of study**

The *in vitro* anti-hyperglycemic activities of the isolated compounds from the stem bark of *Z. chalybeum* and *Z. gilletii* showed the activity against  $\alpha$ -amylase and  $\alpha$ -glucosidase in controlling hyperglycemia. The plant should therefore be subjected to further analysis to check the efficiency on the use the plants and to create a variety of herbal products that can safely, less costly and efficiently manage diabetes mellitus. The active phenolics and alkaloids isolated from the two plants should be subjected to toxicological studies and to serve as templates for developing anti-hyperglycemic drugs if found safe.

#### **5.5. Suggestions for further studies**

- i. From the kinetic results obtained, it would thus be better that molecular interaction studies based on *in silico* experiments to be done to confirm the conclusive information about the compounds and their activities.
- ii. Toxicological and *in vivo* testing will be conducted to determine the effectiveness and toxicity of the separated chemicals.

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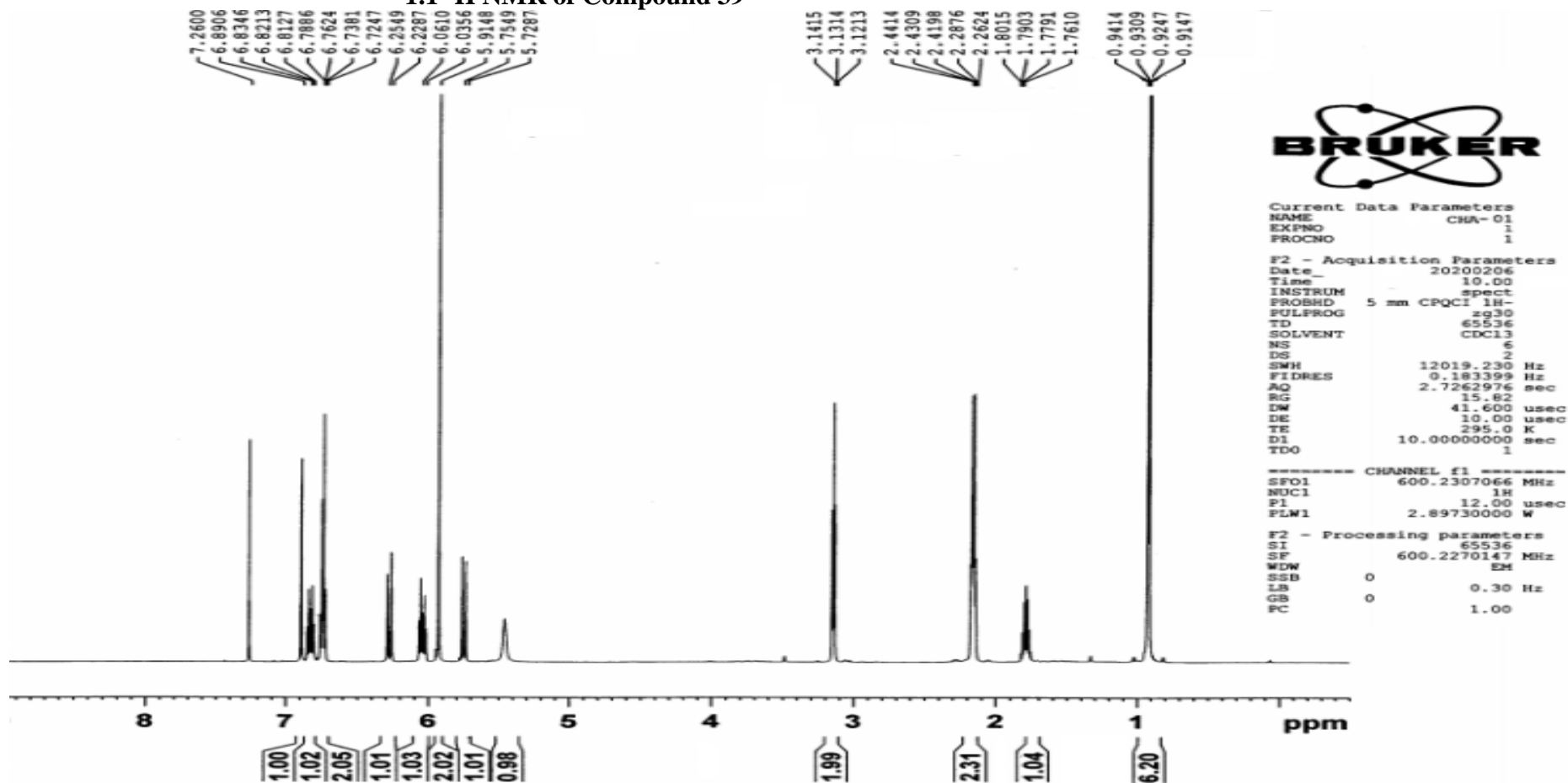
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Zhu, W., Chen, M., Shou, Q., Li, Y., & Hu, F. (2011). Biological Activities of Chinese Propolis and Brazilian Propolis on Streptozotocin-Induced Type 1 Diabetes Mellitus in Rats. *Evidence Based Complementary Alternatives for Medication*, 11, 1–8.

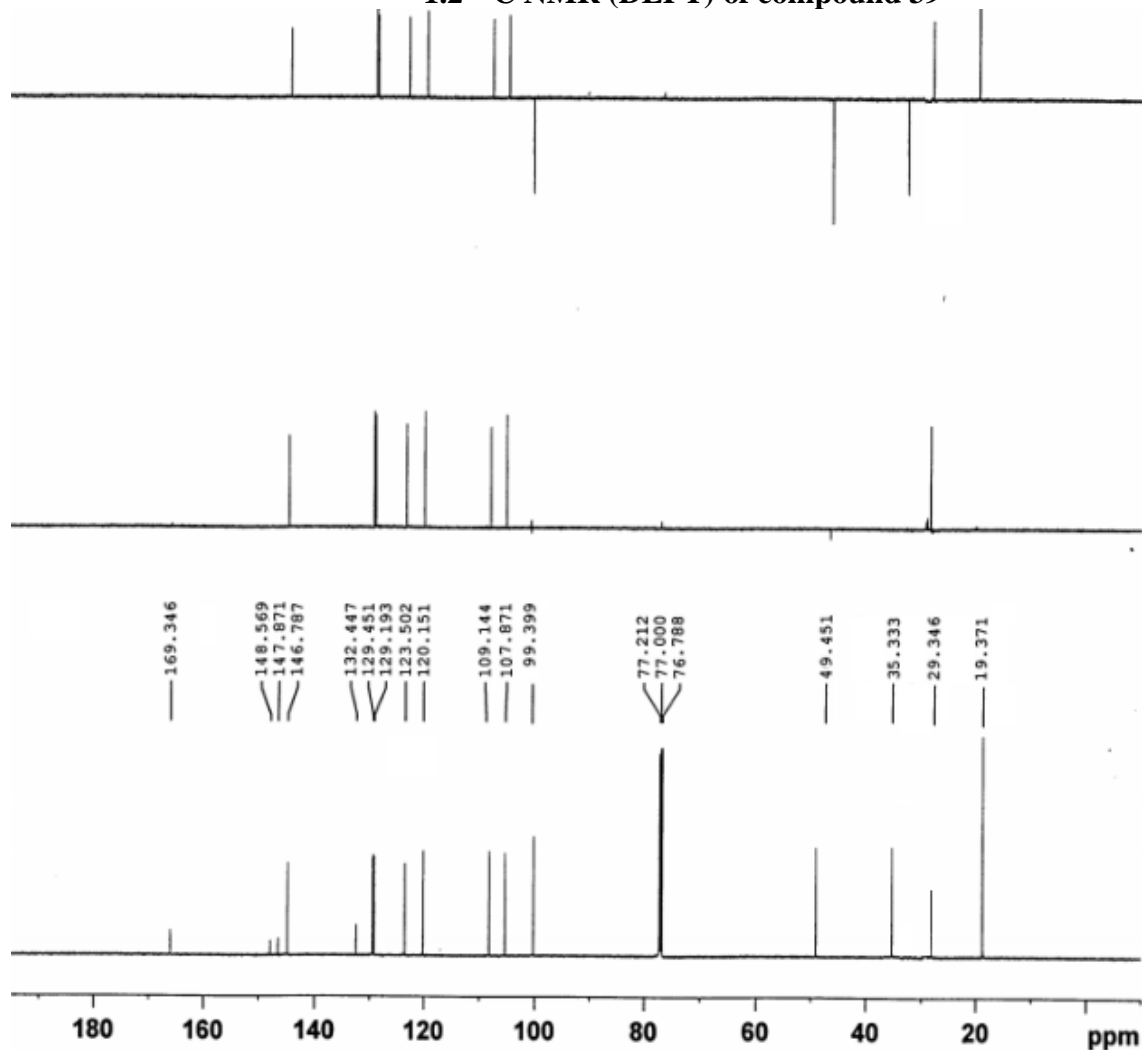
## APPENDICES

### 1. Spectra for compound 39 Chalybemide A {6-Benzo[1,3]dioxol-5-yl-hexa-2,5-dienoic acid isobutylamide}

#### 1.1 <sup>1</sup>H NMR of Compound 39



1.2 <sup>13</sup>C NMR (DEPT) of compound 39



Current Data Parameters  
 NAME CHA-01  
 EXPNO 4  
 PROCNO 1

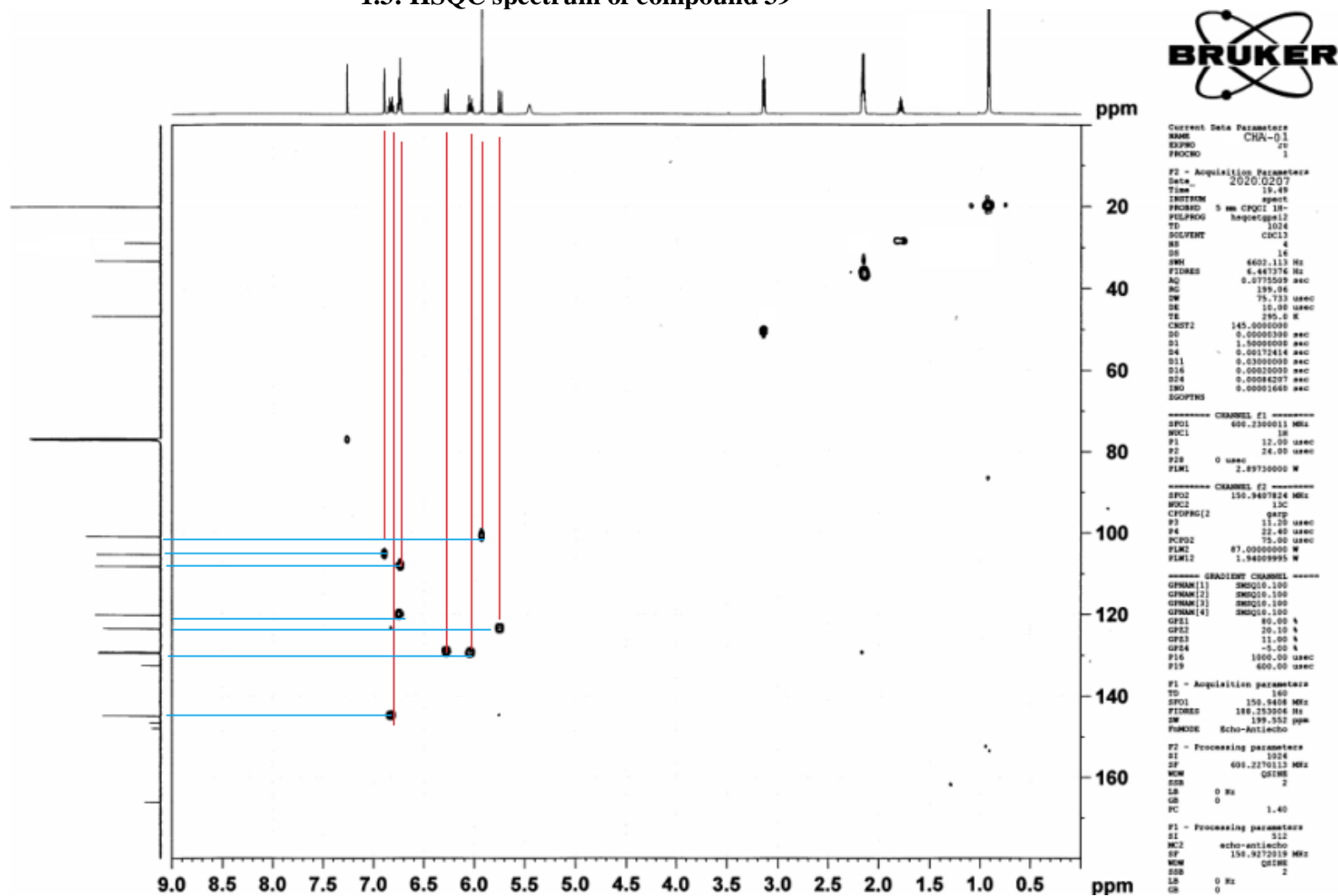
F2 - Acquisition Parameters  
 Date 2020/0207  
 Time 18.06  
 INSTRUM spect  
 PROBRD 5 mm CPQCI 1H-  
 PULPROG zgdc  
 TD 65536  
 SOLVENT CDCl3  
 NS 320  
 DS 4  
 SWH 36231.883 Hz  
 FIDRES 0.552855 Hz  
 AQ 0.9043968 sec  
 RG 199.06  
 DW 13.800 usec  
 DE 18.00 usec  
 TE 295.0 K  
 D1 2.00000000 sec  
 D11 0.03000000 sec  
 TDO 1

----- CHANNEL f1 -----  
 SFO1 150.9438010 MHz  
 NUC1 13C  
 P1 11.20 usec  
 PLW1 94.00000000 W

----- CHANNEL f2 -----  
 SFO2 600.2294009 MHz  
 NUC2 1H  
 CPDPRG2 waltz16  
 PCPD2 80.00 usec  
 PLW2 2.89730000 W  
 PLW12 0.06519000 W

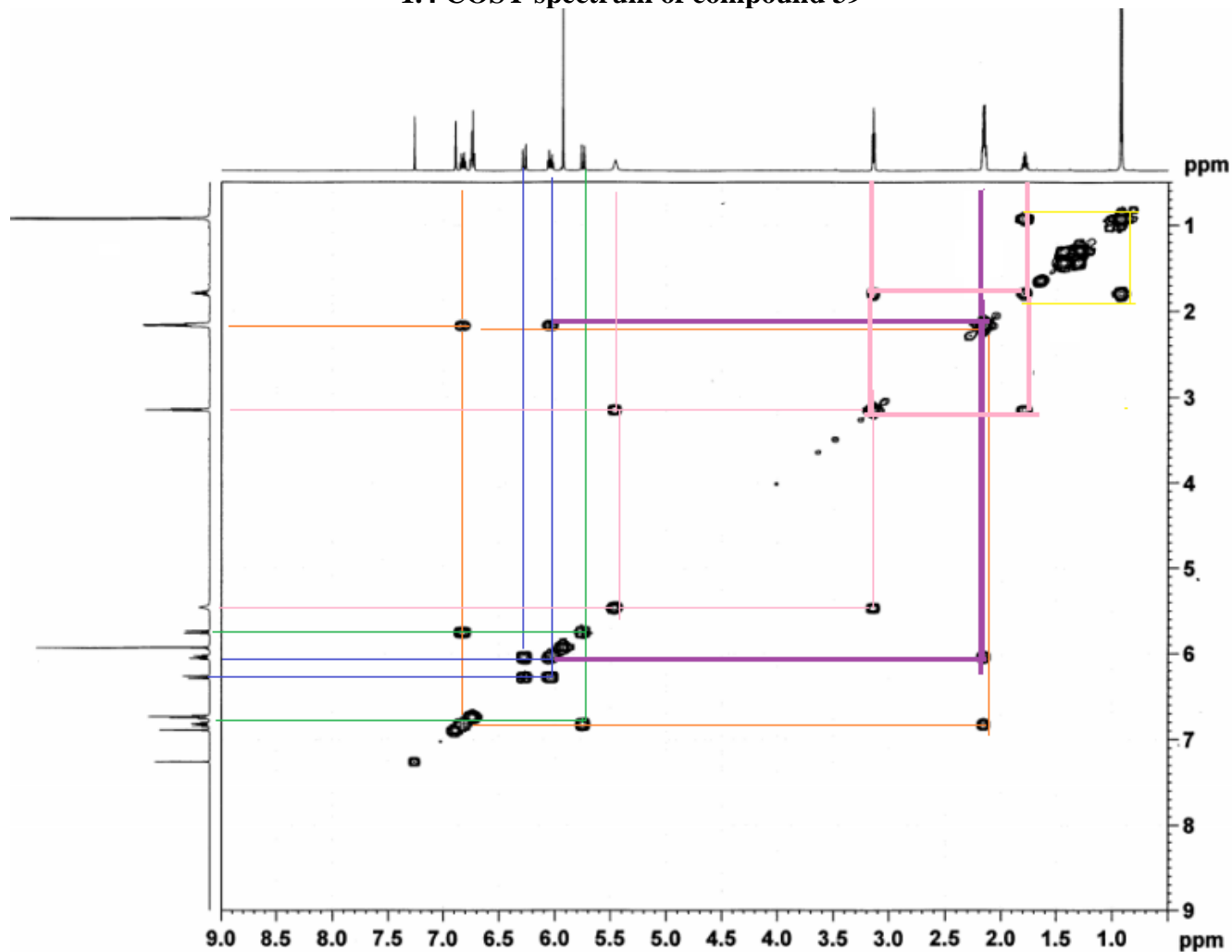
F2 - Processing parameters  
 SI 131072  
 SF 150.9272056 MHz  
 MDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

### 1.3: HSQC spectrum of compound 39





### 1.4 COSY spectrum of compound 39



```

Current Data Parameters
NAME          CHA-01
EXPNO         18
PROCNO        1

F2 - Acquisition Parameters
Date_         2020.0207
Time          12.00
INSTRUM       spect
PROBHD        5 mm CPQCI 1H-
PULPROG       cosypppgzg
TD            1024
SOLVENT       CDCl3
NS            4
DS            8
SWH           7211.539 Hz
FIDRES        7.042518 Hz
AQ            0.0709973 sec
RG            135.28
DM            69.333 usec
DE            10.00 usec
TE            295.0 K
DO            0.0000300 sec
D1            1.5000000 sec
D11           0.0300000 sec
D12           0.0002000 sec
D13           0.0000400 sec
D14           0.0002000 sec
IN0           0.00013680 sec

----- CHANNEL f1 -----
SFO1          600.230011 MHz
NUC1           1H
P0            12.00 usec
F1            12.00 usec
P17           2500.00 usec
PLW1          2.89730000 W
PLW10         0.61719000 W

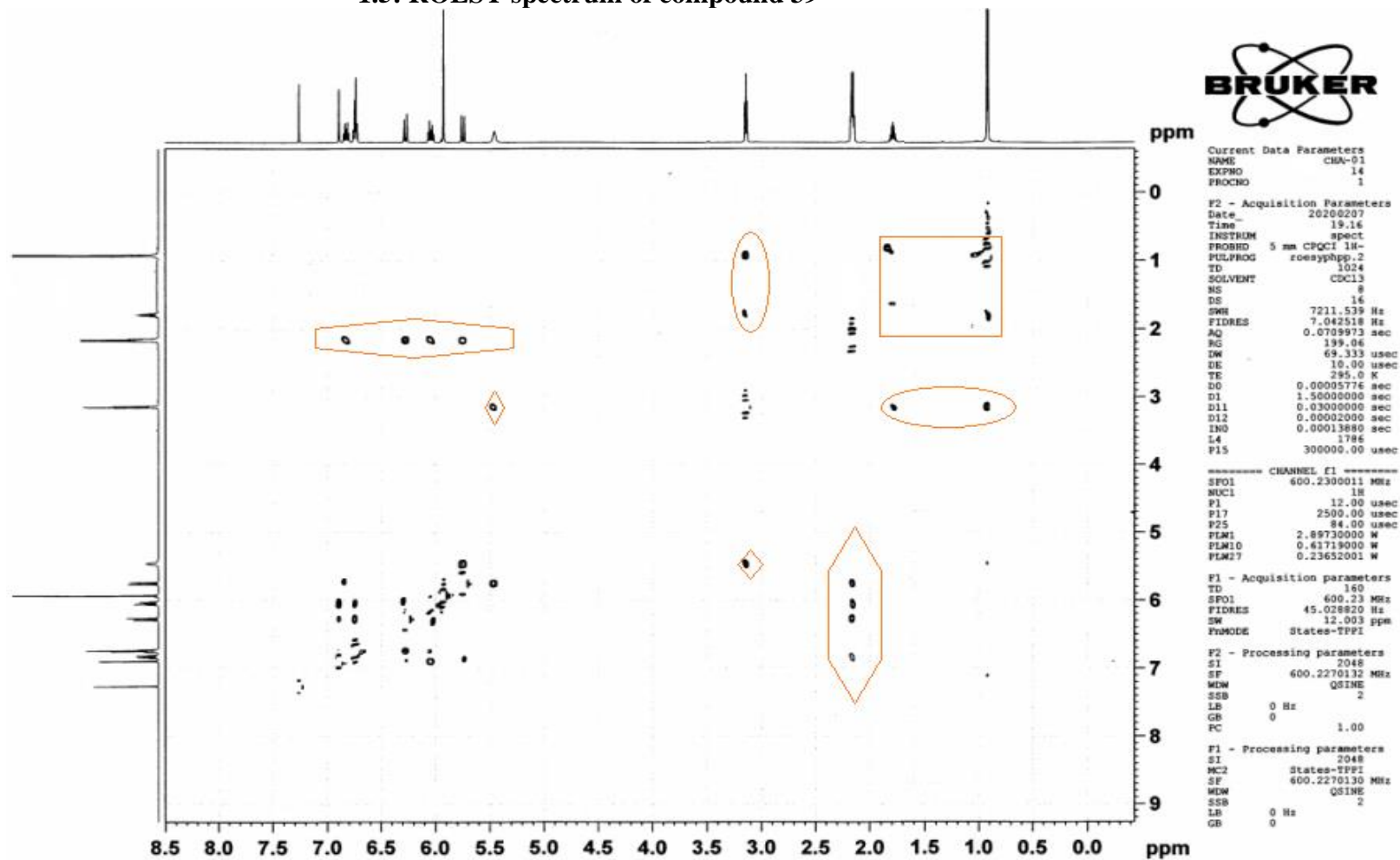
----- GRADIENT CHANNEL -----
GPMAX[1]      0.00010.100
GPE1          10.00 %
P16           1000.00 usec

F1 - Acquisition parameters
TD            160
SFO1          600.23 MHz
FIDRES        45.028620 Hz
SW            12.003 ppm
F2MODE        QF

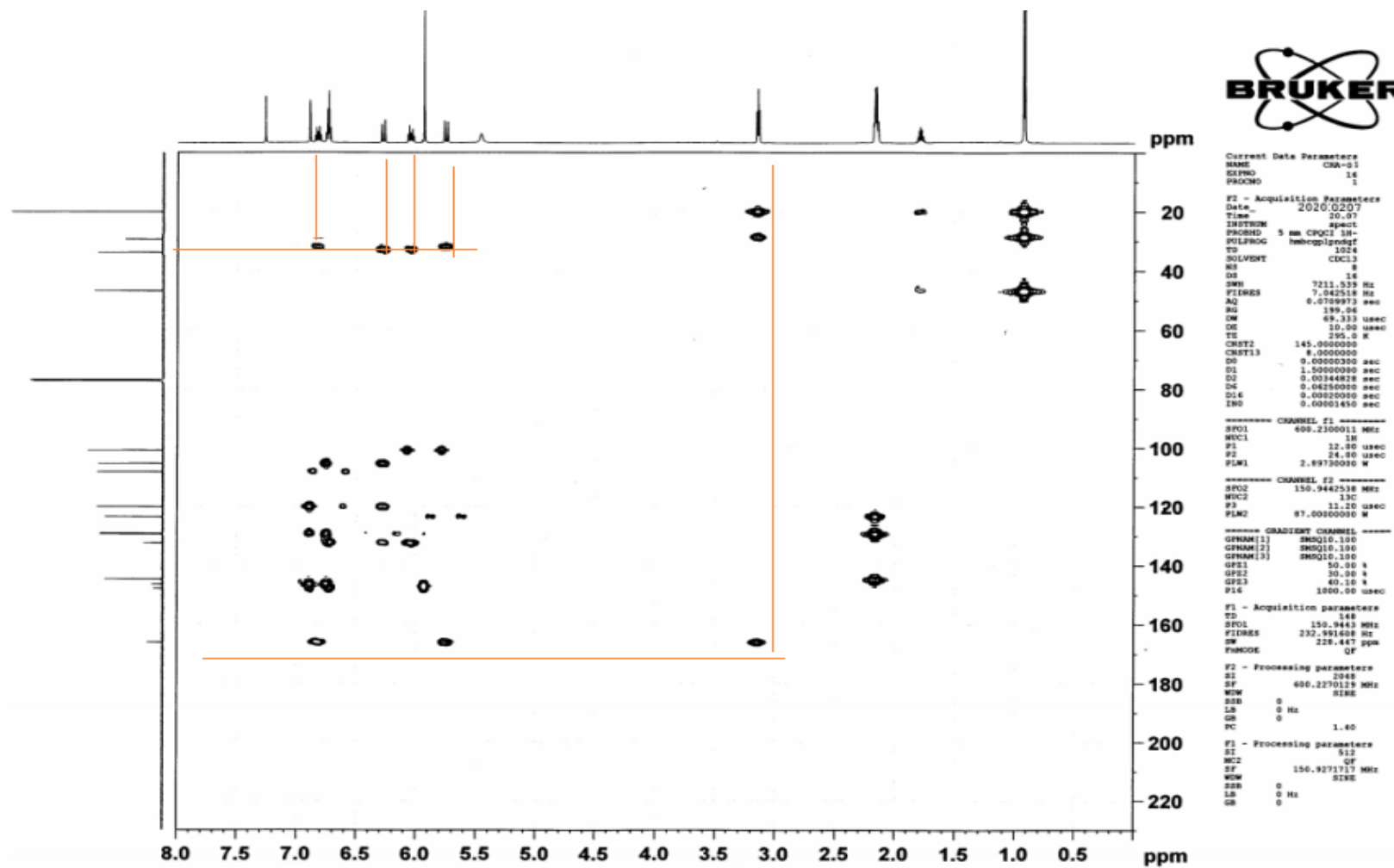
F2 - Processing parameters
SI            2048
SF            600.2270145 MHz
WDW           QZINE
SSB           0
LB            0 Hz
GB            0
PC            1.40

F1 - Processing parameters
SI            2048
MC2           QF
SF            600.2270116 MHz
WDW           QZINE
SSB           0
LB            0 Hz
GB            0
    
```

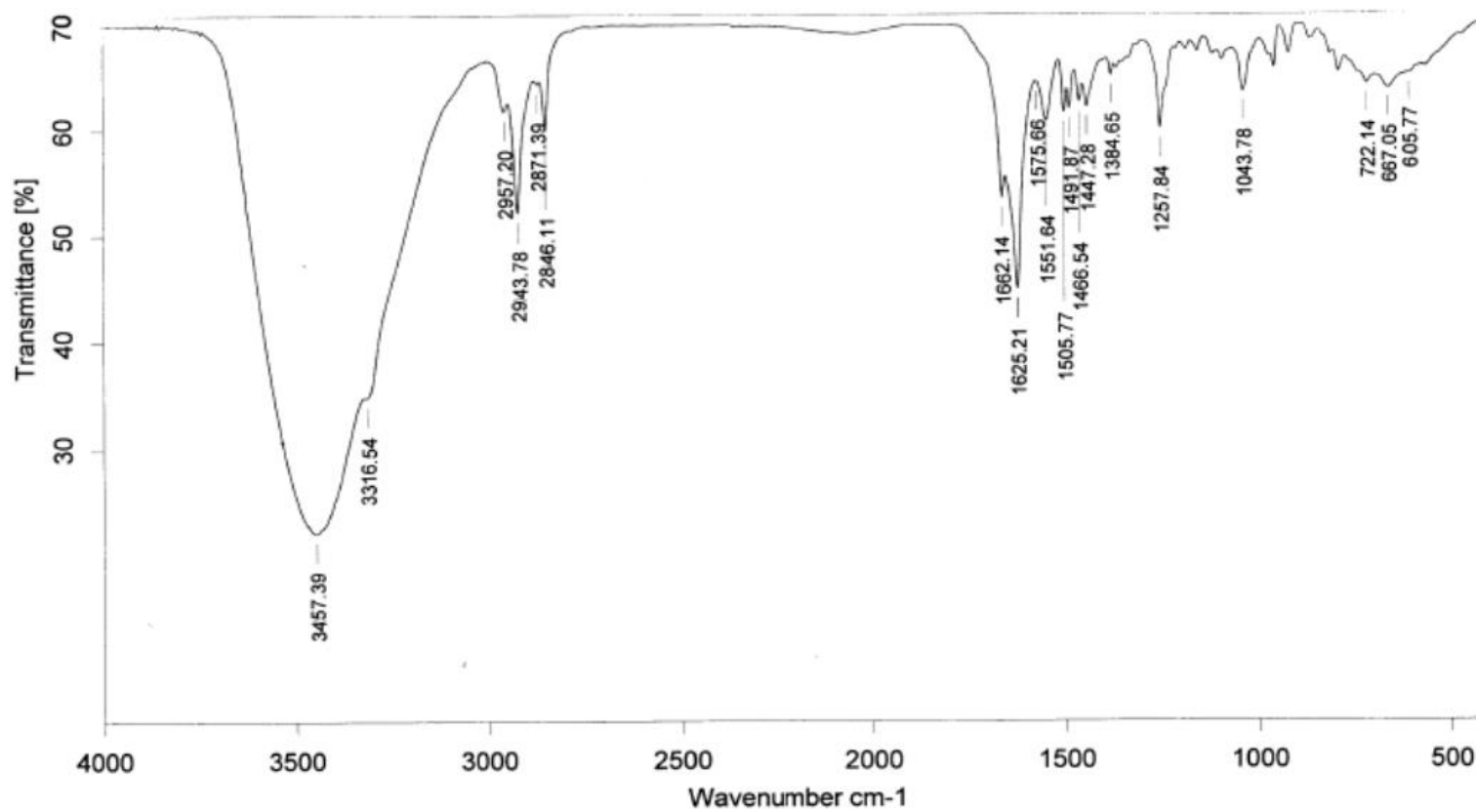
### 1.5: ROESY spectrum of compound 39



# 1.6:HMBC spectrum of compound 39



1.7: FTIR spectrum of compound 39



### 1.8: HREIMS spectrum of compound 39

#### Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -10.0, max = 120.0

Selected filters: None

Monoisotopic Mass, Odd and Even Electron Ions

20 formula(e) evaluated with 2 results within limits (up to 52 closest results for each mass)

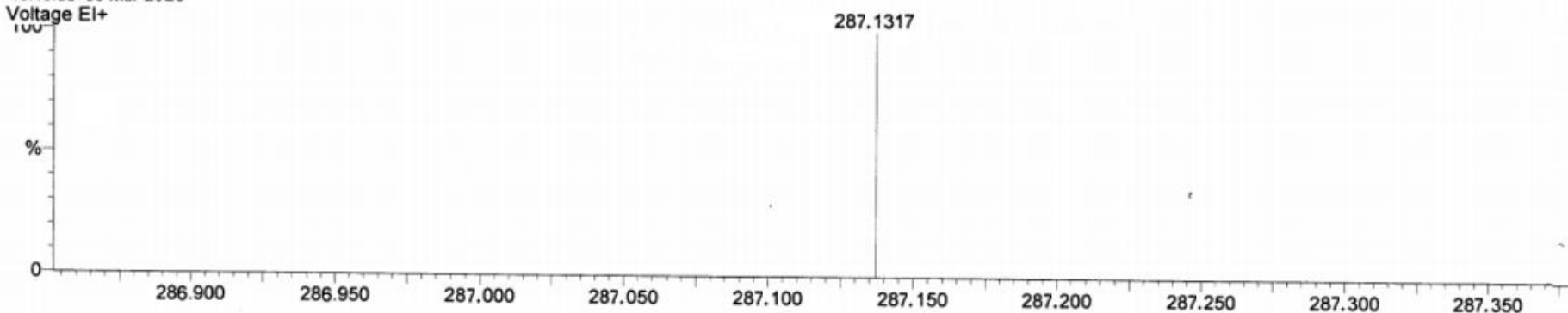
Elements Used:

C: 0-200 H: 0-400 N: 1-1 O: 2-4

cha-07

15:13:38 09-Mar-2020

Voltage EI+

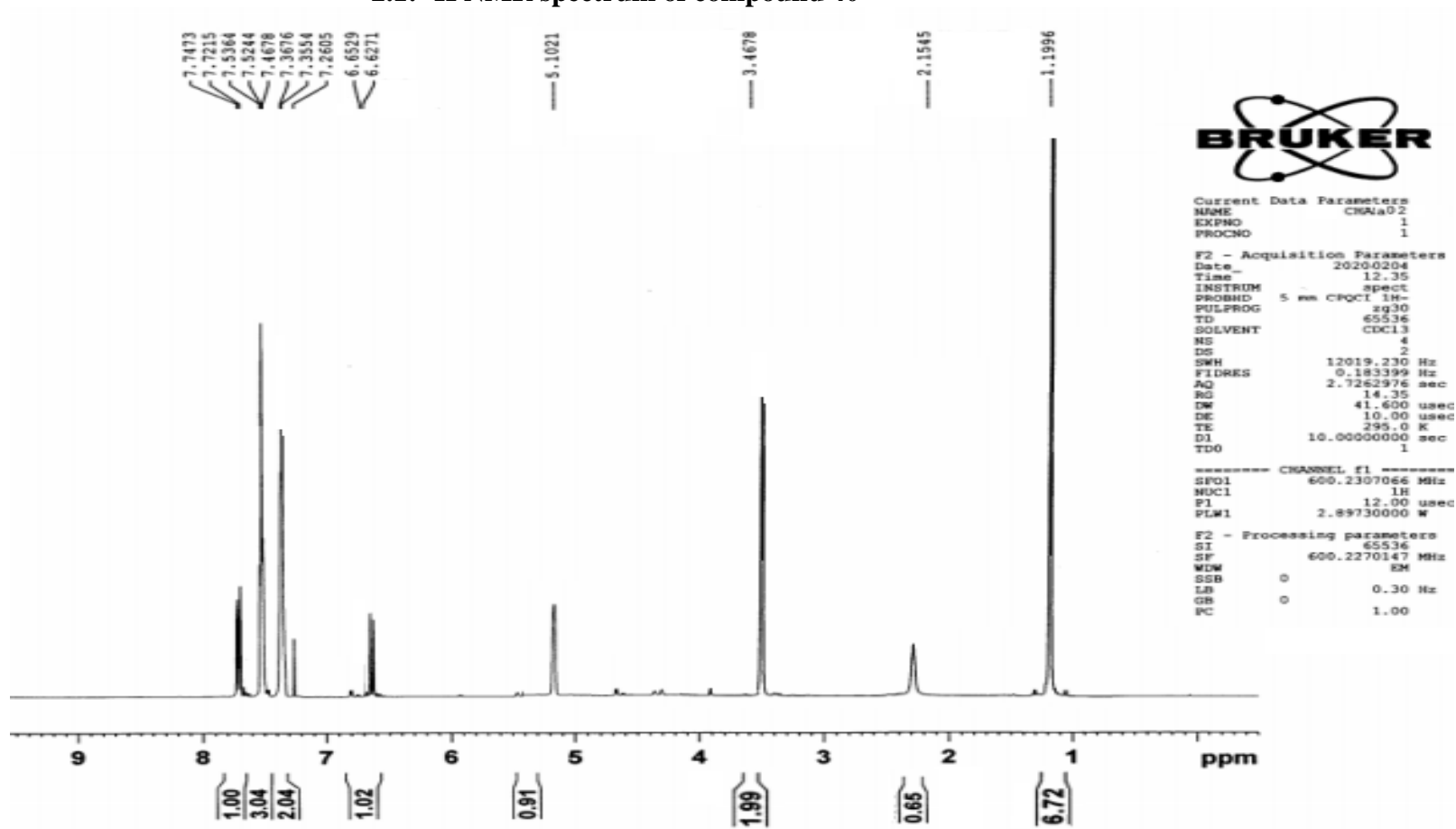


Minimum: -10.0  
Maximum: 200.0 10.0 120.0

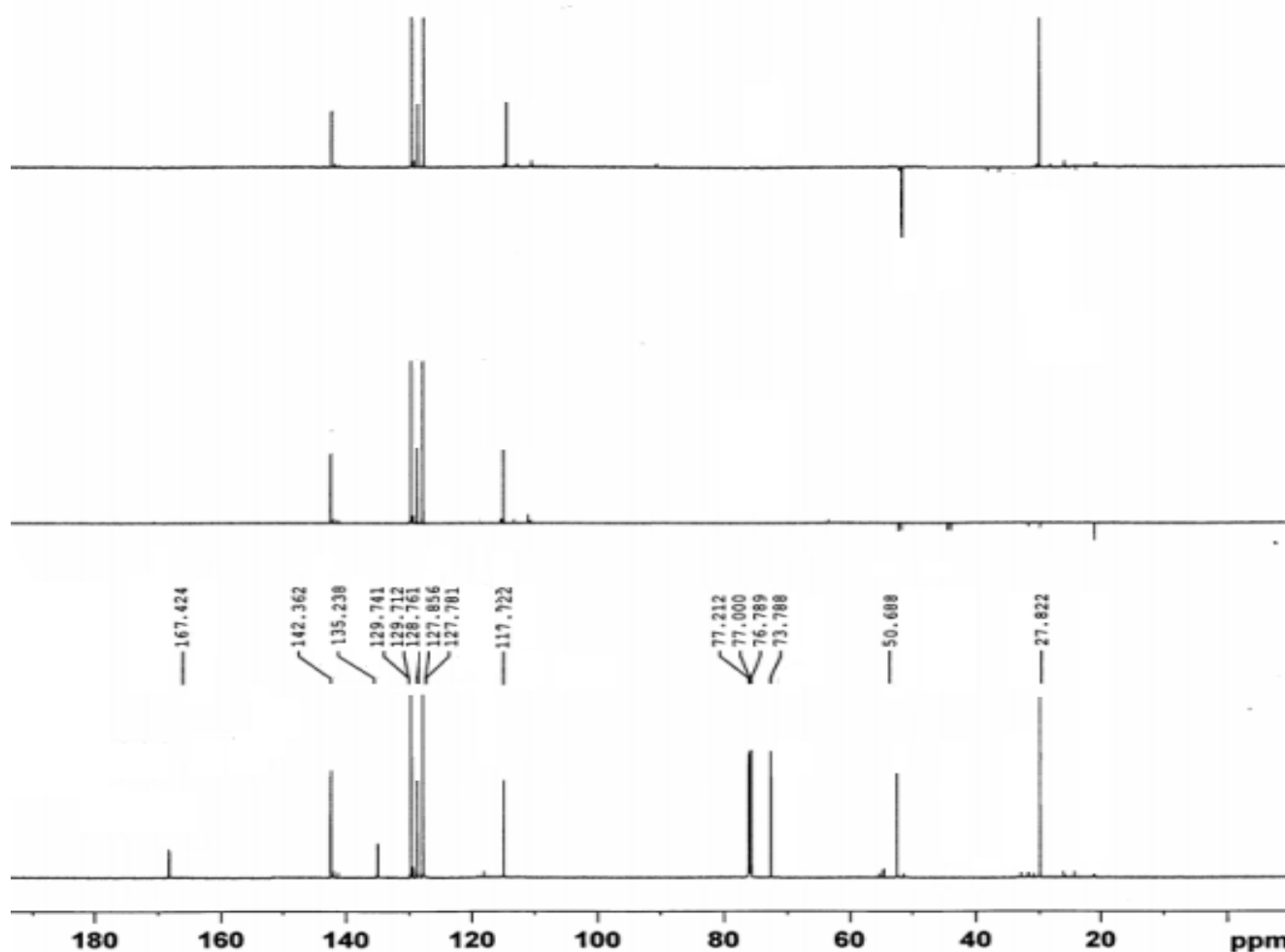
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
287.1317	287.1312	0.8	2.8	9.0	5546027.5	C17 H21 N O3

## 2. Spectra for Compound 40 (Chalybemide B)

### 2.1: $^1\text{H}$ NMR spectrum of compound 40



## 2.2 <sup>13</sup>C NMR (DEPT) spectrum of compound 40



Current Data Parameters  
 NAME CHA-02  
 EXPNO 13  
 PROCNO 1

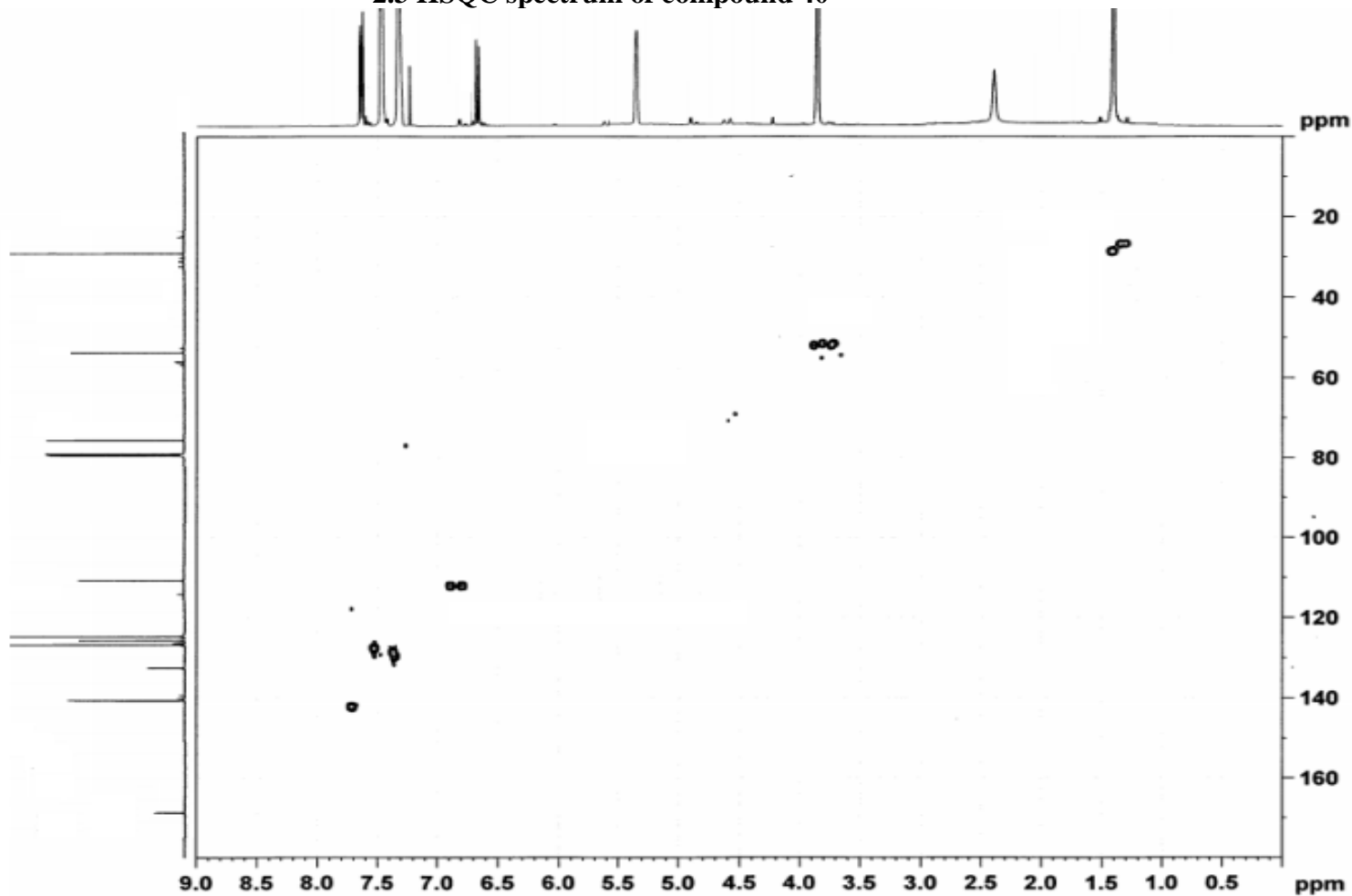
F2 - Acquisition Parameters  
 Date\_ 20200204  
 Time 13.04  
 INSTRUM spect  
 PROBHD 5 mm CPXI IN-  
 PULPROG zgpg30  
 TD 65536  
 SOLVENT CDCl3  
 NS 256  
 DS 4  
 SWH 36231.883 Hz  
 FIDRES 0.552855 Hz  
 AQ 0.9043968 sec  
 RG 139.06  
 DW 13.800 usec  
 DE 18.00 usec  
 TE 295.0 K  
 D1 2.0000000 sec  
 D11 0.0300000 sec  
 TD0 1

----- CHANNEL f1 -----  
 SFO1 150.9438010 MHz  
 NUC1 13C  
 F1 11.20 usec  
 PLW1 94.0000000 W

----- CHANNEL f2 -----  
 SFO2 600.2294009 MHz  
 NUC2 1H  
 CDFRG[2] waltz16  
 PCPD2 80.00 usec  
 PLW2 2.89730000 W  
 PLW12 0.06519000 W

F2 - Processing parameters  
 SI 131072  
 SF 150.9272109 MHz  
 WDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

### 2.3 HSQC spectrum of compound 40



```

Current Data Parameters
NAME          C9H8O2
EXPNO        20
PROCNO        1

F2 - Acquisition Parameters
Date_         20100204
Time          23.23
INSTRUM      spect
PROBHD       5 mm CPQCI 1H-
PULPROG      zgpg30
TD           1024
SOLVENT      CDCl3
NS           4
DS           16
SWH          4402.113 Hz
FIDRES      0.447374 Hz
AQ          0.0775069 sec
RG          199.04
DE          70.733 usec
TE          10.00 usec
TR          290.0 K
CR22        145.0000000
D0          0.00000000 sec
D1          1.50000000 sec
D4          0.00172414 sec
D13         0.03000000 sec
D14         0.00000000 sec
E24         0.00084207 sec
RG          0.00001440 sec
SFOPT0

----- CHANNEL f1 -----
SFO1        600.330011 MHz
NUC1         1H
P1          12.00 usec
P2          24.00 usec
P3         0 usec
P4         2.85730000 W
PULSE1

----- CHANNEL f2 -----
SFO2        150.9407824 MHz
NUC2         13C
CPDPRG2      zgpg
P3          11.20 usec
P4          22.40 usec
PCPD2       70.00 usec
PULSE2      ST.00000000 W
PULSE2      1.94009995 W

----- GRADIENT CHANNEL -----
GPRAM[1]    SMO10.100
GPRAM[2]    SMO10.100
GPRAM[3]    SMO10.100
GPRAM[4]    SMO10.100
GR1         80.00 %
GR2         20.10 %
GR3         11.00 %
GR4         -5.00 %
P16         3000.00 usec
P18         600.00 usec

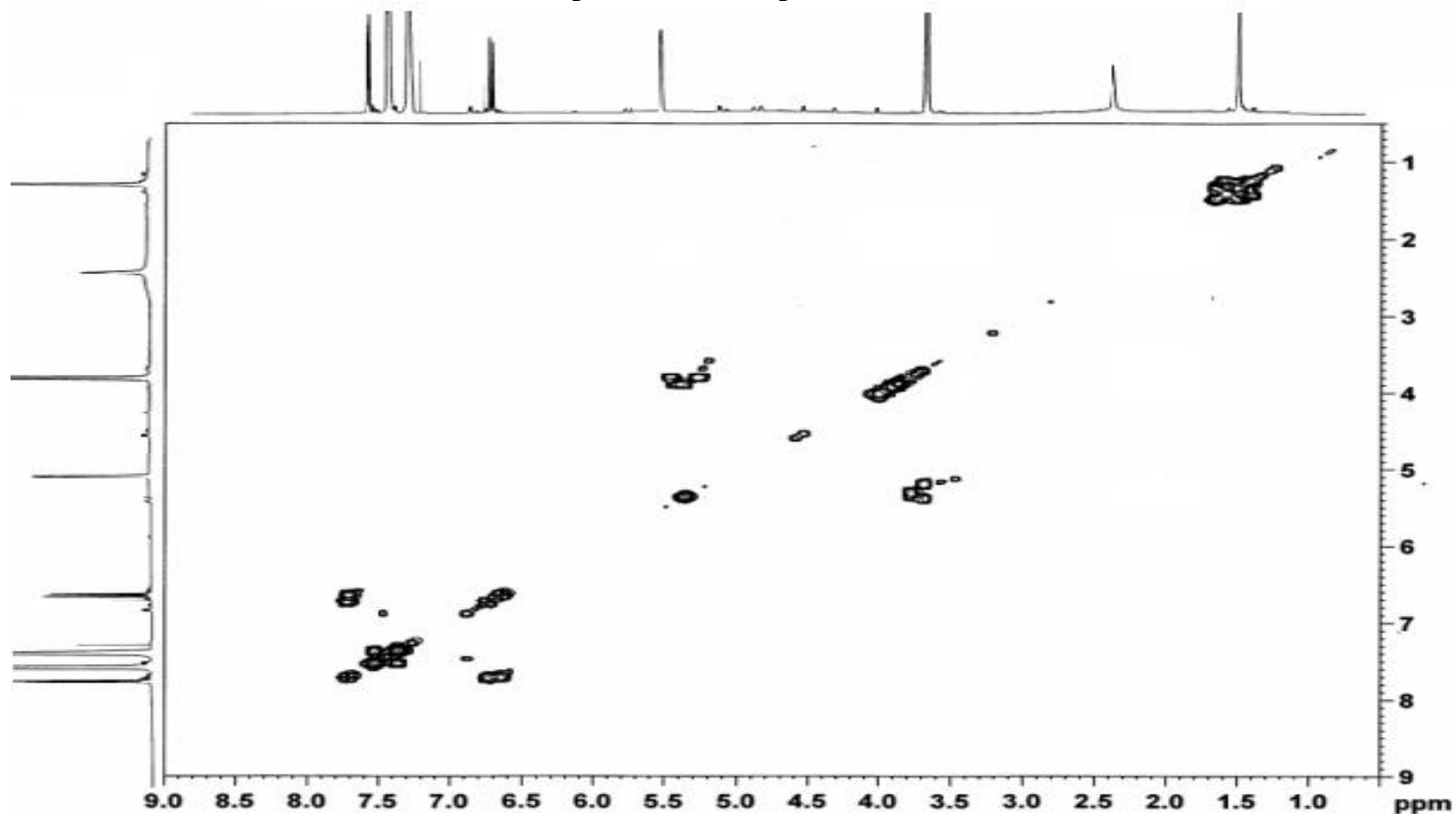
F1 - Acquisition parameters
TD          140
SFO1        150.9406 MHz
FIDRES      189.263004 Hz
SW          199.952 ppm
PROBHD      Echo-Astiecho

F2 - Processing parameters
SI          3274
SF          600.2270104 MHz
WDW         GAUSS
SSB         0
LB          0 Hz
GB          0
PC          1.40

F1 - Processing parameters
SI          512
WDW         echo-astrischo
SF          150.9272140 MHz
WDW         GAUSS
SSB         0
LB          0 Hz
GB          0
  
```



## 2.4: COSY spectrum of compound 40



```

Current Data Parameters
NAME          C14-02
EXPNO        10
PROCNO       1

F2 - Acquisition Parameters
Date_        20250904
Time_        22.22
INSTRUM      spect
PROBHD       5 mm CPQCI 1H-
PULPROG      cosygpppgf
TD           1024
SOLVENT      CDCl3
NS           4
DS           4
SWH          7211.538 Hz
FIDRES       7.042518 Hz
AQ           0.8709973 sec
RG           57.14
SM          69.333 usec
SE           19.00 usec
TE           295.0 K
DQ           0.0000300 sec
SI           1.0000000 sec
Sf1          0.0300000 sec
Sf2          0.08002000 sec
Sf3          0.00000400 sec
Df1          0.00020000 sec
IN0          0.00010000 sec

----- CHANNEL f1 -----
NUC1          600.220011 MHz
P1           12.00 usec
PL1          0.0000000
PL2          2500.00 usec
PLM1         2.89130000 W
PLM2         0.41710000 W

----- GRADIENT CHANNEL -----
GPMAX111     0.0000000
GPE1         10.00 %
P14          1000.00 usec

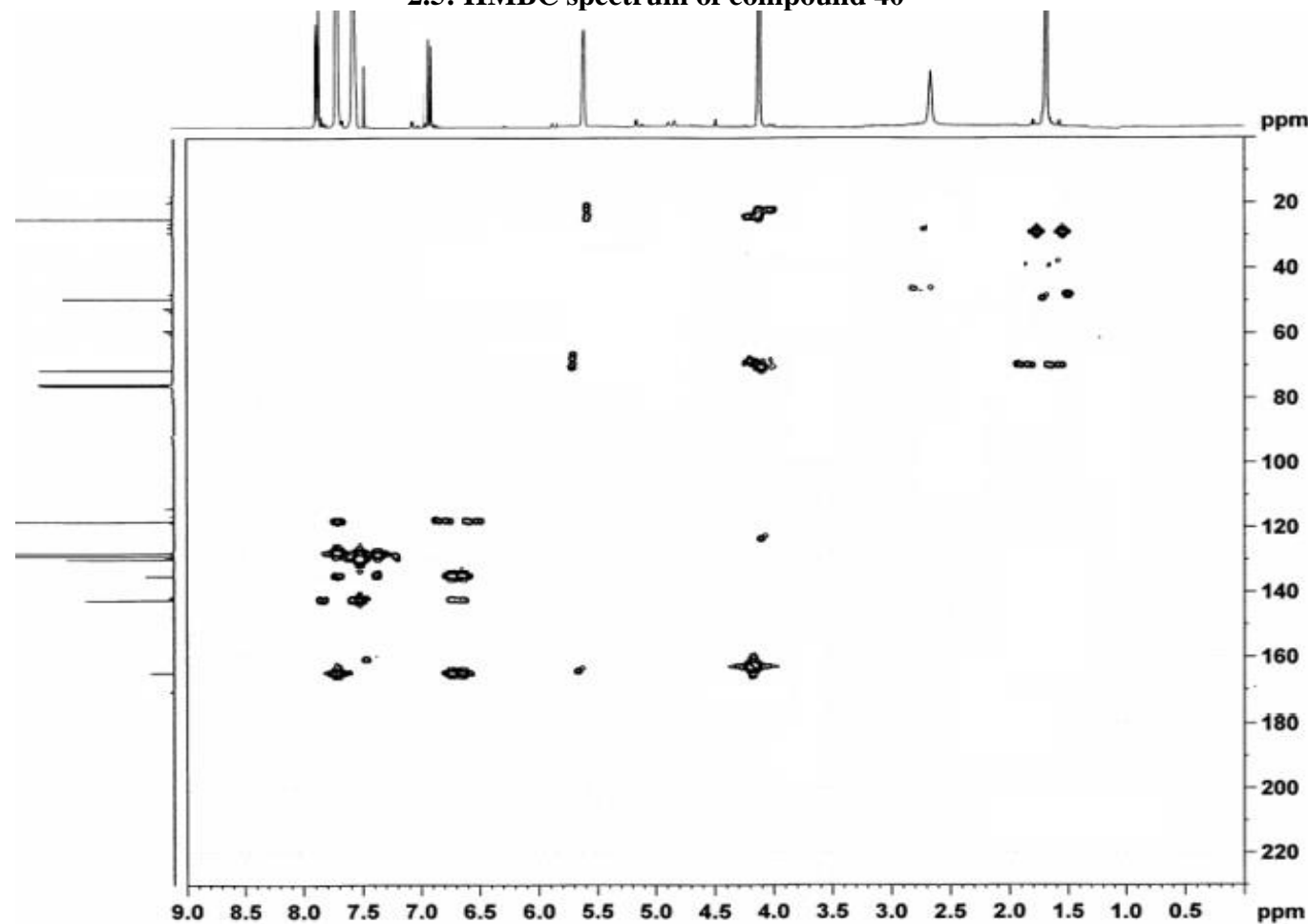
F1 - Acquisition parameters
TD           100
SFO1         600.23 MHz
FIDRES       45.028820 Hz
AQ           11.003 usec
SFO2         0
F2MODE       OF

F2 - Processing parameters
SI           2048
SF           600.2270113 MHz
WDW          Q3
SSB          0
LB           0 Hz
GB           0
PC           1.40

F1 - Processing parameters
SI           2048
SF           600.2270113 MHz
WDW          Q3
SSB          0
LB           0 Hz
GB           0

```

2.5: HMBC spectrum of compound 40



```

Current Data Parameters
NAME      C04-02
EXPNO    16
PROCNO   1

F2 - Acquisition Parameters
Date_    20200304
Time     16.10
INSTRUM spect
PROBHD   5 mm CPQCI 1H-
PULPROG zgpg30
TD       3274
SOLVENT  CDCl3
NS       2
DS       4
SWH      7211.528 Hz
FIDRES   7.042318 Hz
AQ       0.0789973 sec
RG       195.04
SWH      49.333 usec
SE       16.00 usec
TE       296.0 K
CQZT2    145.000000
CQZT13   8.000000
DQ       0.0000000 usec
S1       1.5000000 usec
S2       0.0014625 usec
S6       0.0423000 usec
S16      0.0002000 usec
S30      0.0000146 usec

----- CHANNEL f1 -----
NUC1     1H
P1       12.00 usec
P2       24.00 usec
PL1      2.09730000 W

----- CHANNEL f2 -----
NUC2     13C
P3       11.20 usec
PL2      87.0000000 W

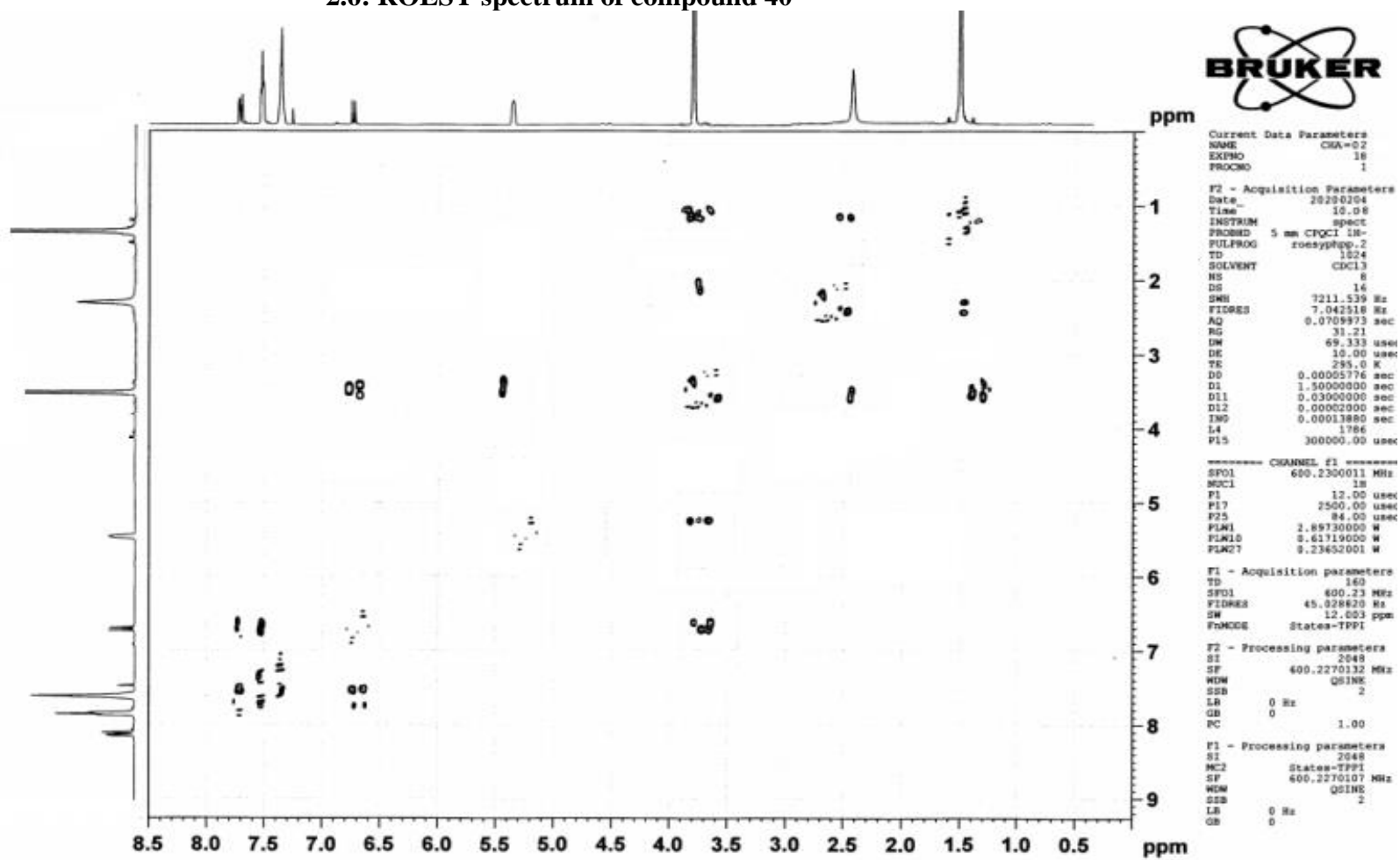
----- GRADIENT CHANNEL -----
GPMAX[1] 0.0000.100
GPMAX[2] 0.0000.100
GPMAX[3] 0.0000.100
GRF1     50.00 Hz
GRF2     36.00 Hz
GRF3     40.10 Hz
F14      1000.00 usec

F1 - Acquisition parameters
TD       144
SFO1     150.9445 MHz
FIDRES   232.991408 Hz
SWH      228.647 ppm
FWDACQ   QF

F2 - Processing parameters
SI       2048
SF       600.2270142 MHz
WDW      SINE
SSB      0
LB       0 Hz
GB       0
PC       1.40

F1 - Processing parameters
SI       612
SF       150.9271990 MHz
WDW      SINE
SSB      0
LB       0 Hz
GB       0
    
```

## 2.6: ROESY spectrum of compound 40



2.7 : HREIMS spectrum of compound 40

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -10.0, max = 120.0

Selected filters: None

Monoisotopic Mass, Odd and Even Electron Ions

11 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)

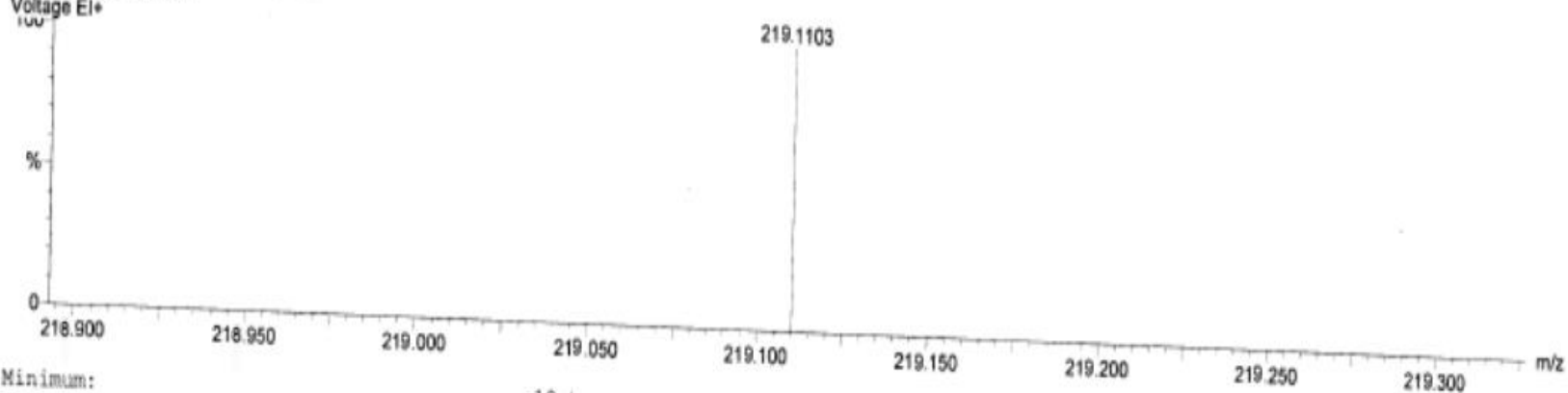
Elements Used:

C: 0-200 H: 0-400 N: 1-1 O: 1-3

cha-02

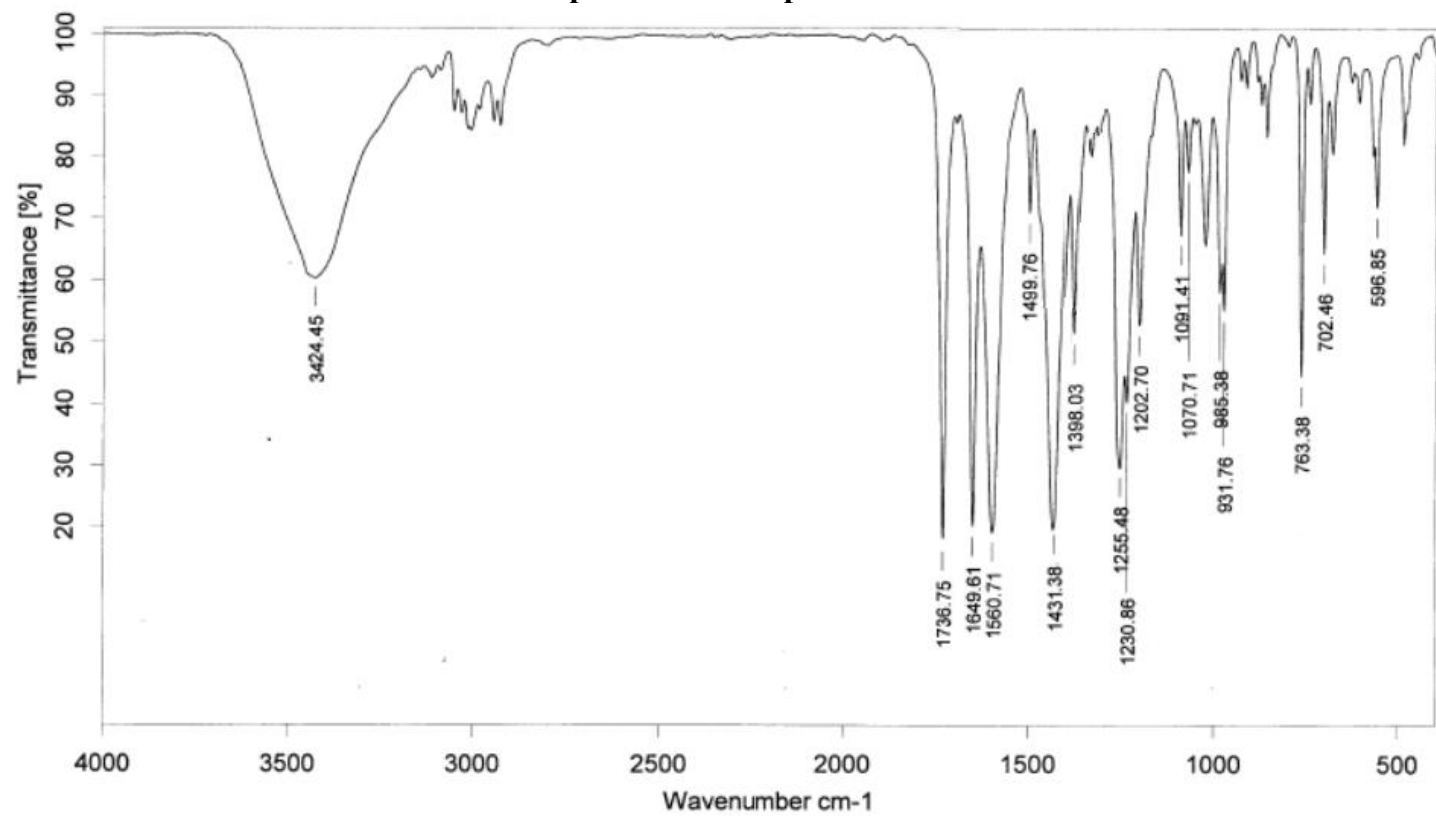
02:22:20 02-Apr-2020

Voltage E1+



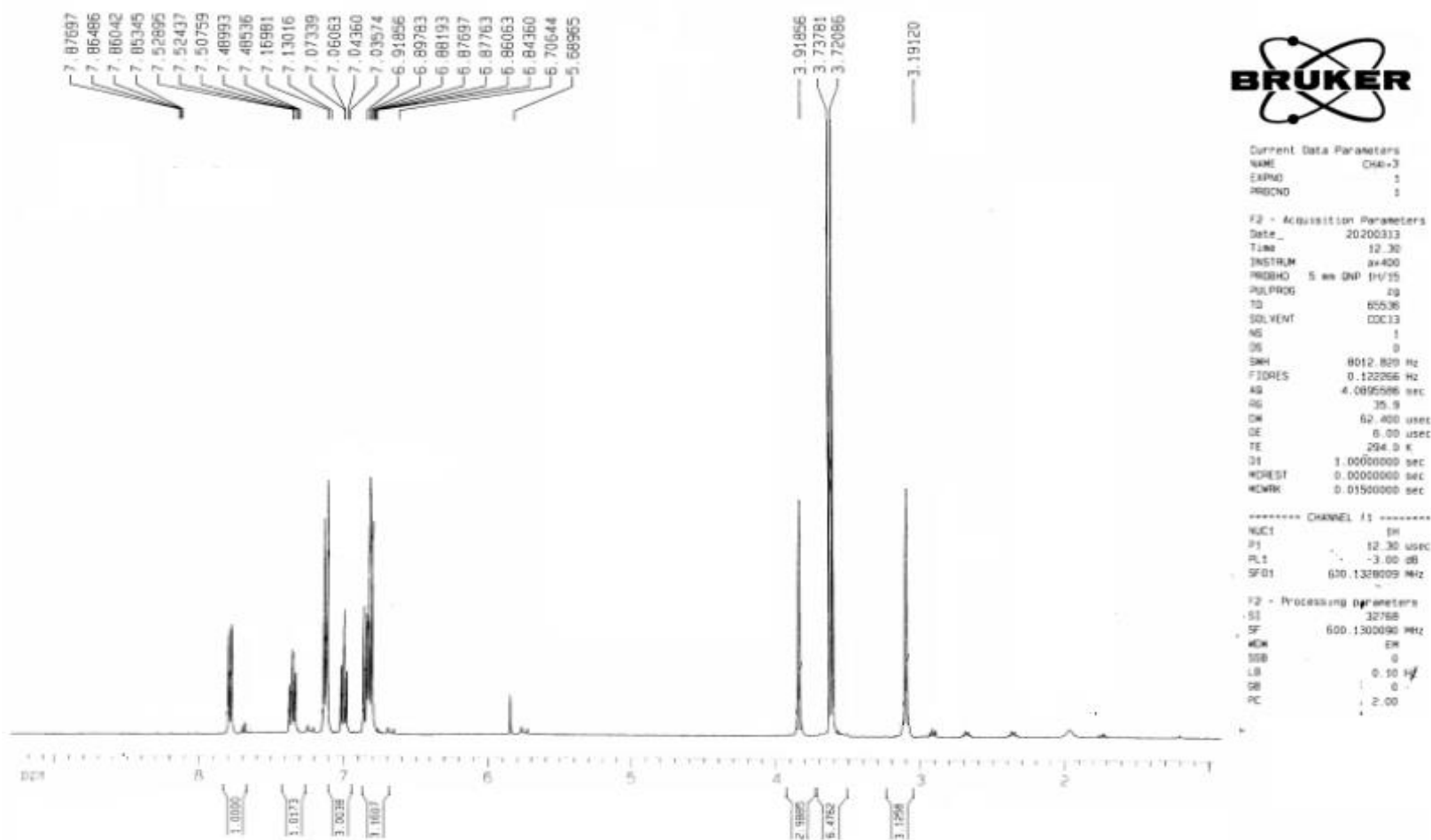
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
219.1103	219.1103	0.0	0.0	7.0	5546025.5	C13 H17 N O2

2.8: FTIR spectrum of compound 40

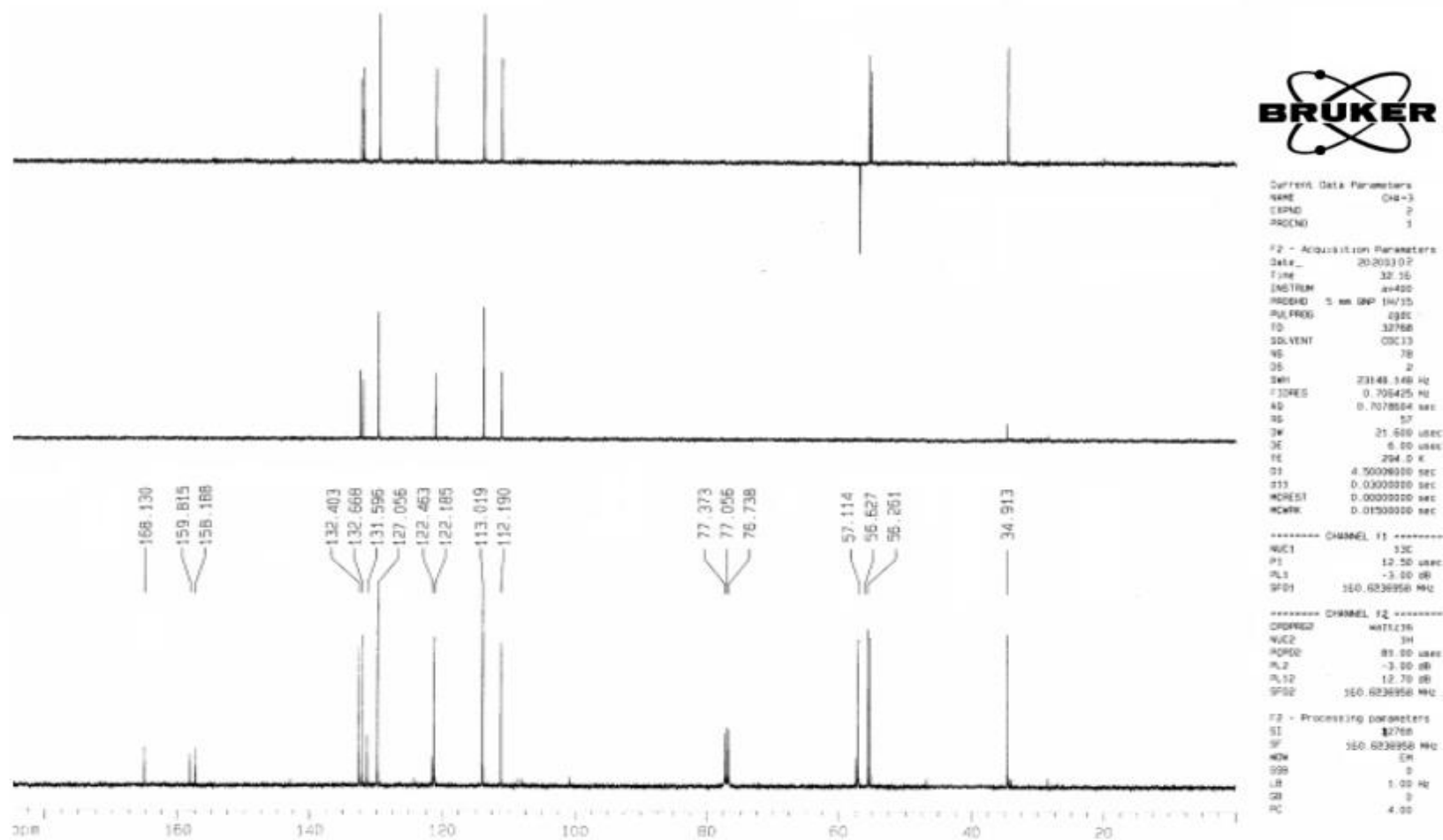


### 3. Spectra for Compound 41 (Chalybemide C)

#### 3.1: <sup>1</sup>H NMR spectrum of compound 41



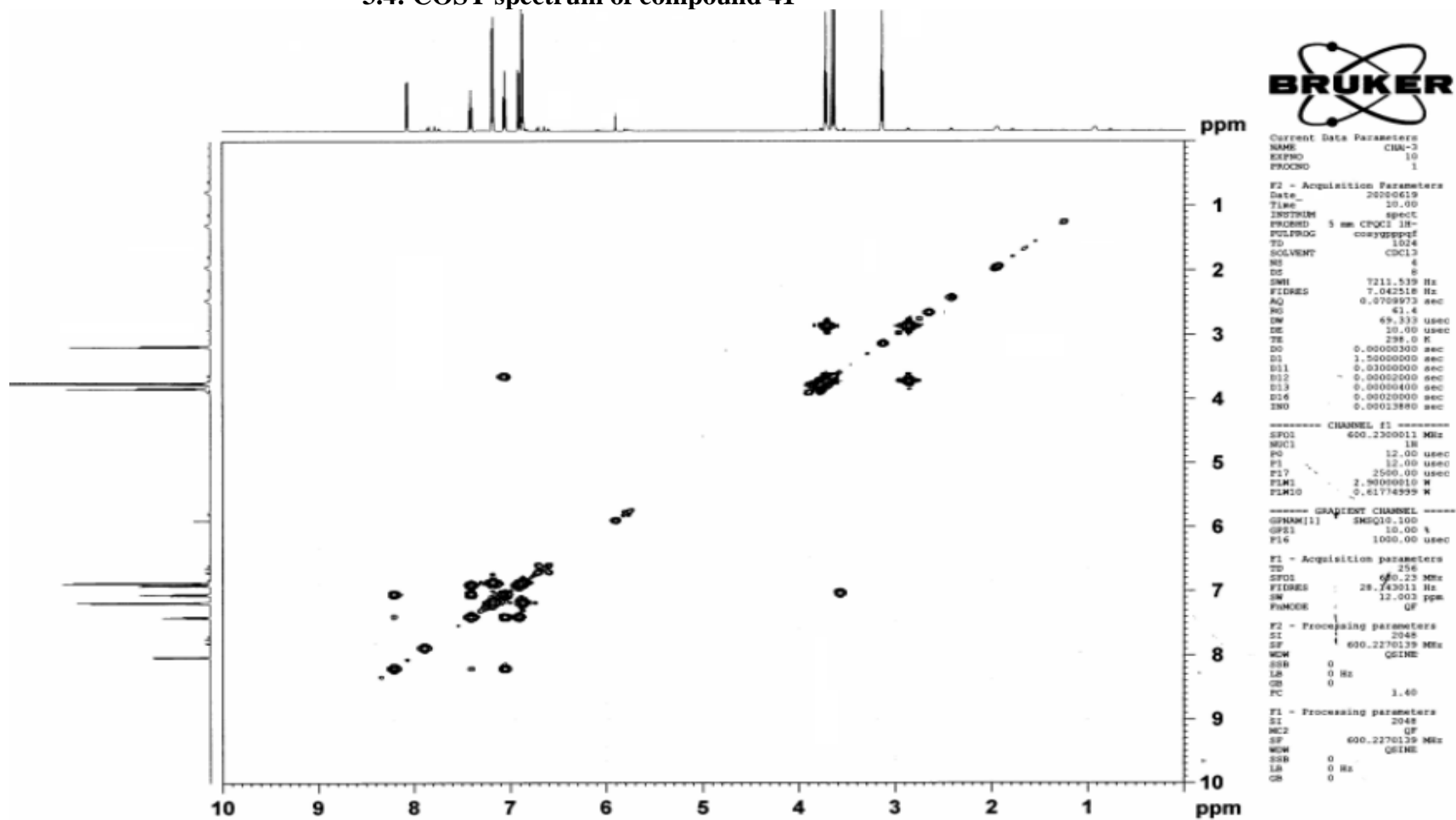
### 3.2: $^{13}\text{C}$ NMR spectrum of compound 41



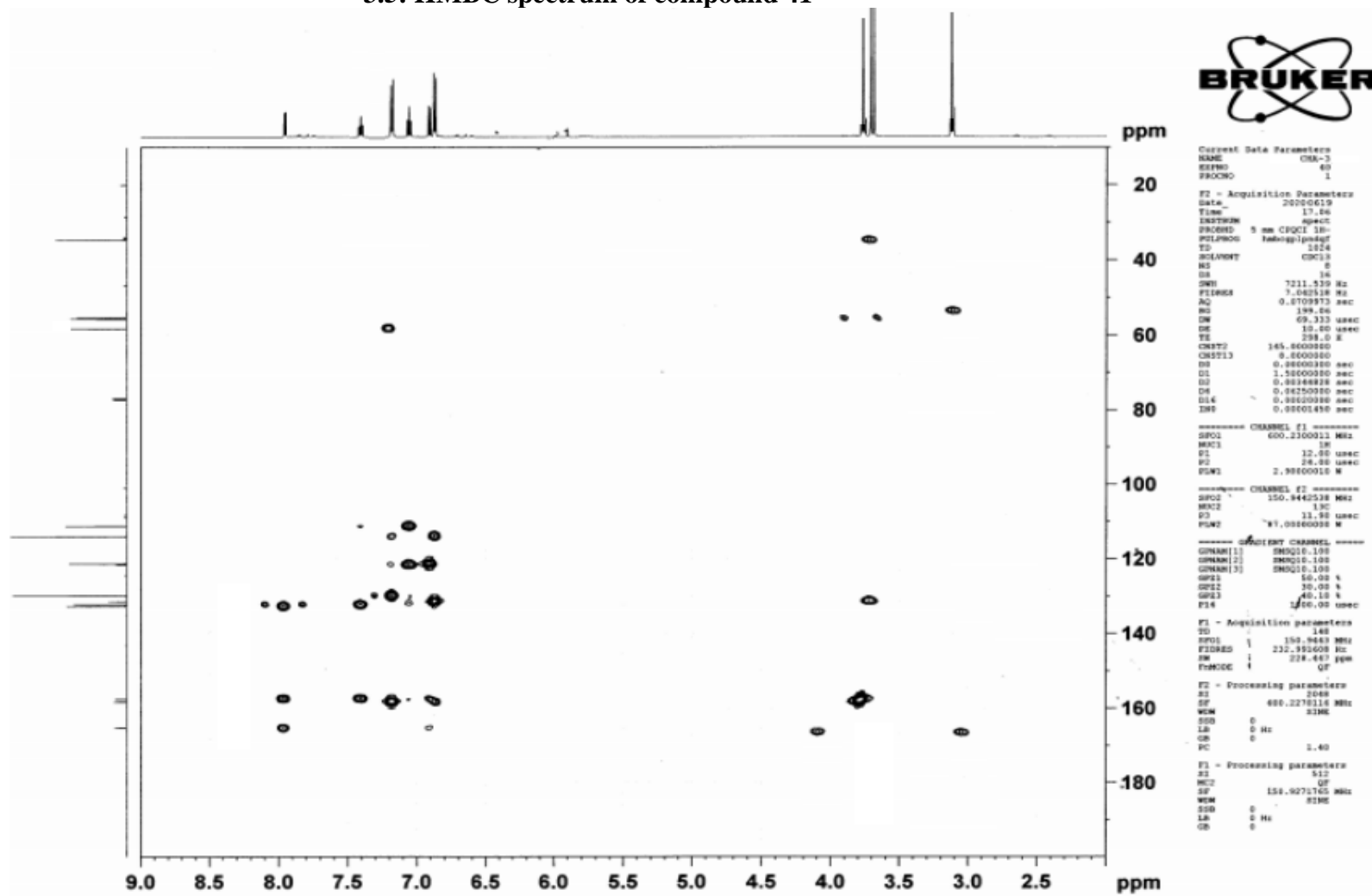




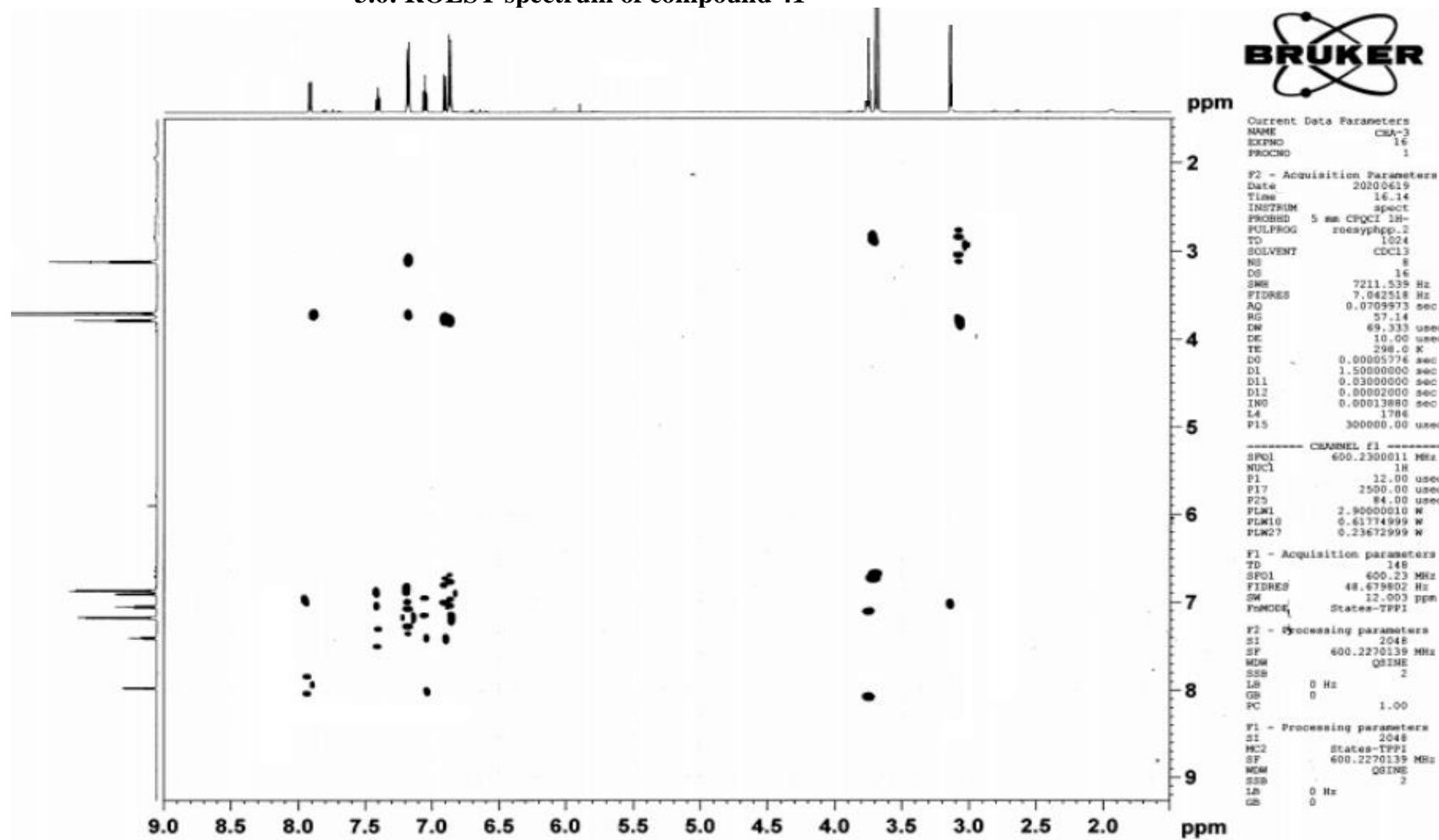
### 3.4: COSY spectrum of compound 41



### 3.5: HMBC spectrum of compound 41



### 3.6: ROESY spectrum of compound 41



### 3.7: HREIMS spectrum of compound 41

#### Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -10.0, max = 120.0

Selected filters: None

Monoisotopic Mass, Odd and Even Electron Ions

10 formula(e) evaluated with 1 results within limits (up to 21 closest results for each mass)

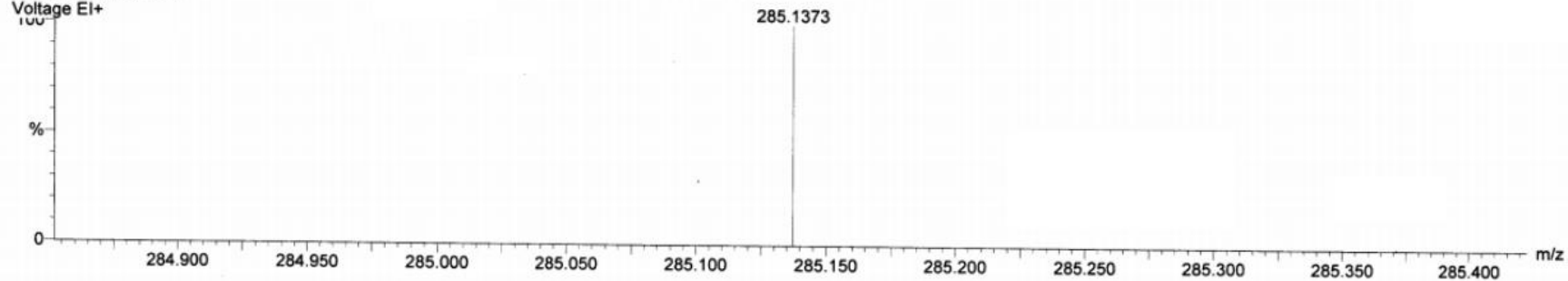
Elements Used:

C: 0-200 H: 0-400 N: 1-1 O: 2-4

cha-3

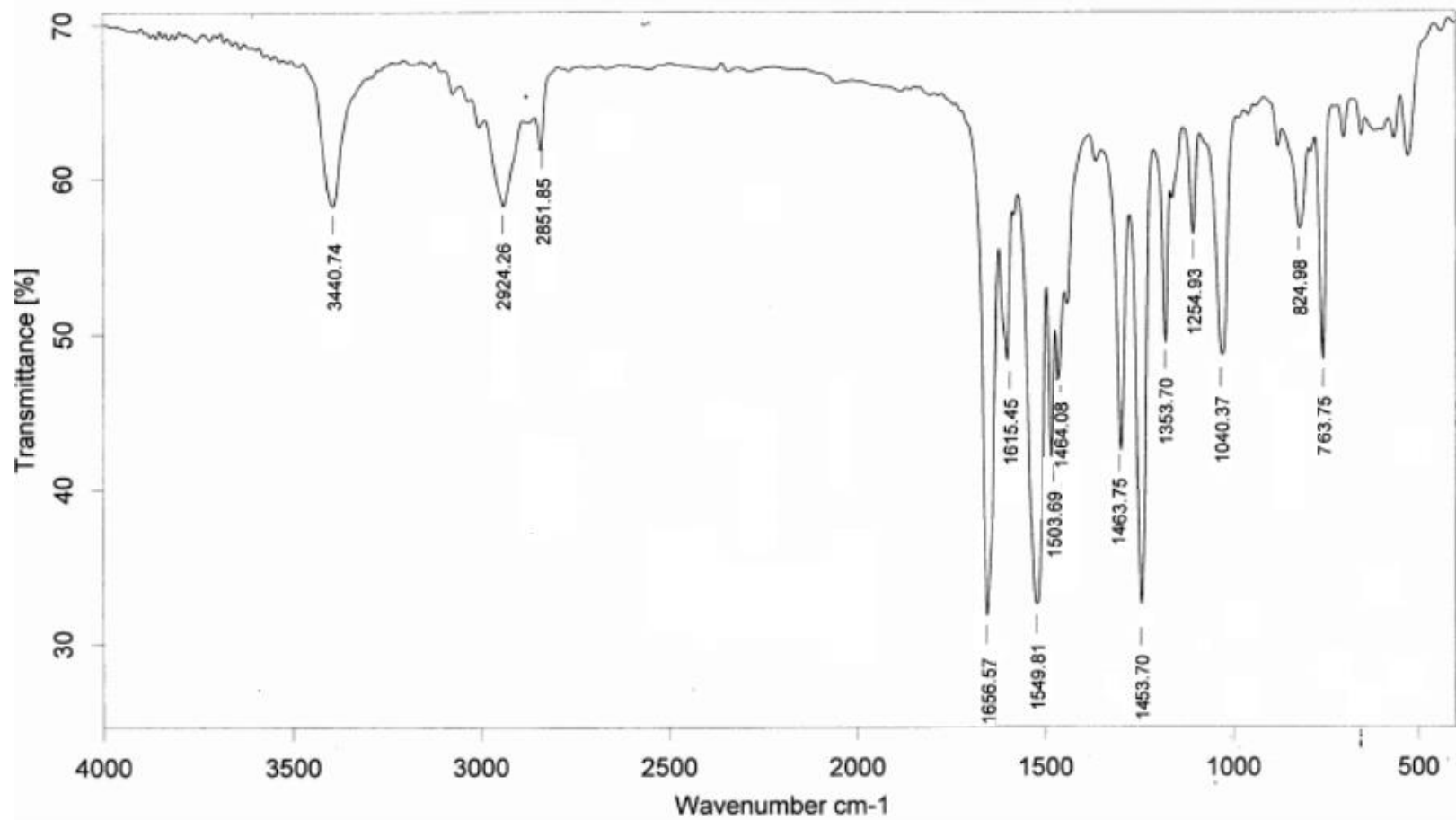
13:30:20 14-Jul-2020

Voltage EI+



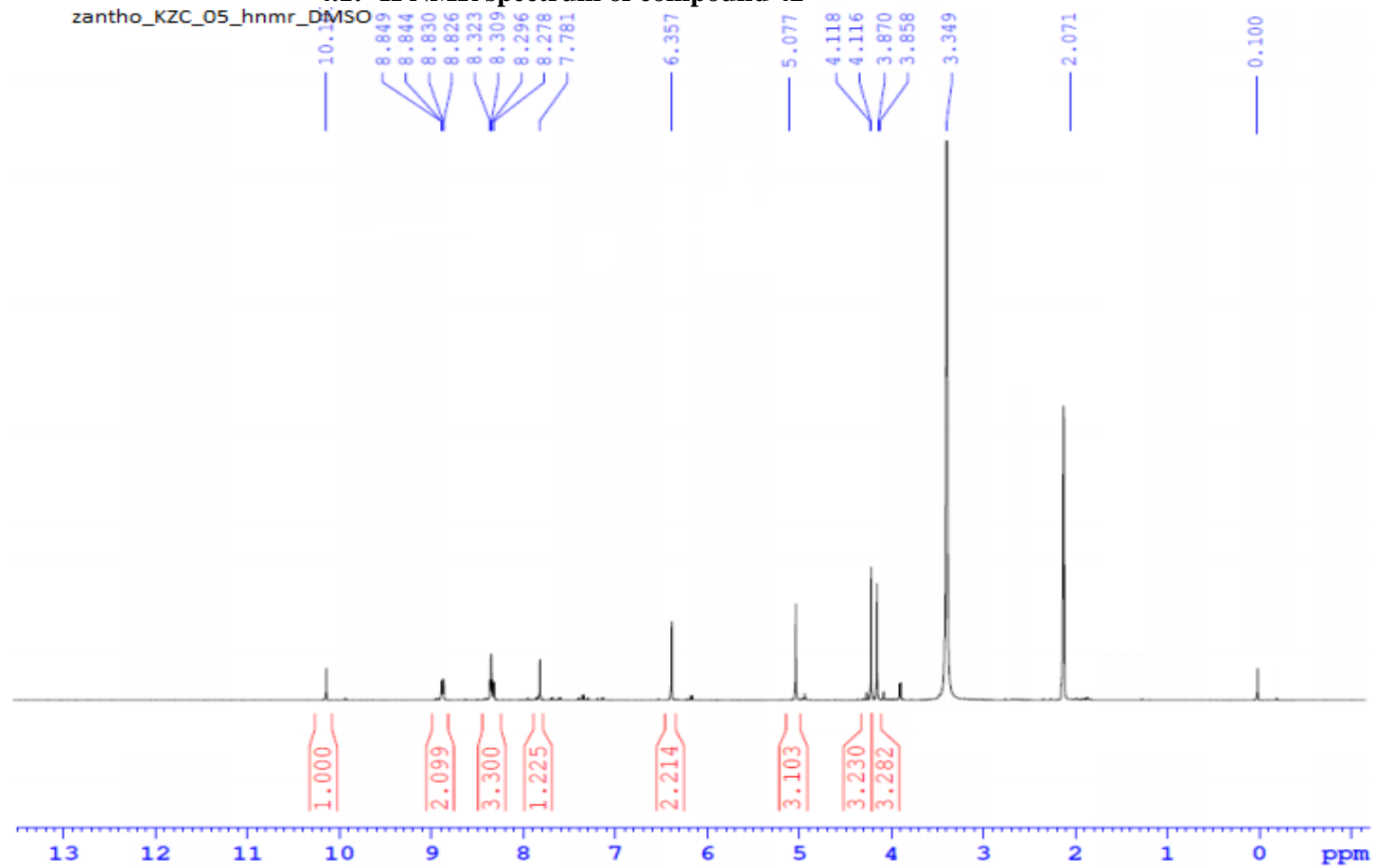
Minimum:				-10.0		
Maximum:	200.0	10.0		120.0		
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
285.1373	285.1365	0.8	2.8	9.0	5546027.5	C17 H19 N O3

3.8: FTIR spectrum of compound 41

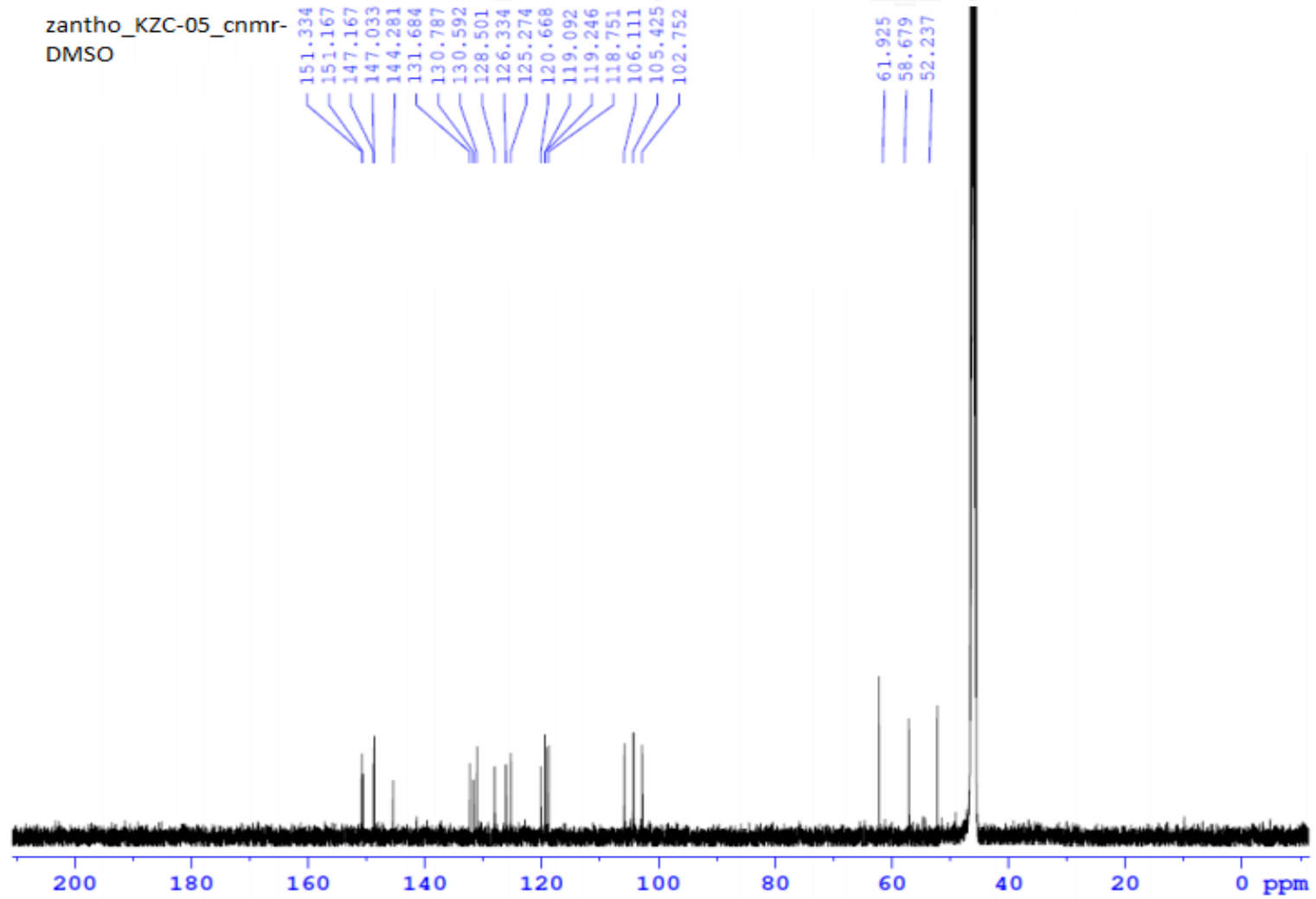


#### 4. Spectra for Compound 42 (Phenanthridine A)

##### 4.1: $^1\text{H}$ NMR spectrum of compound 42

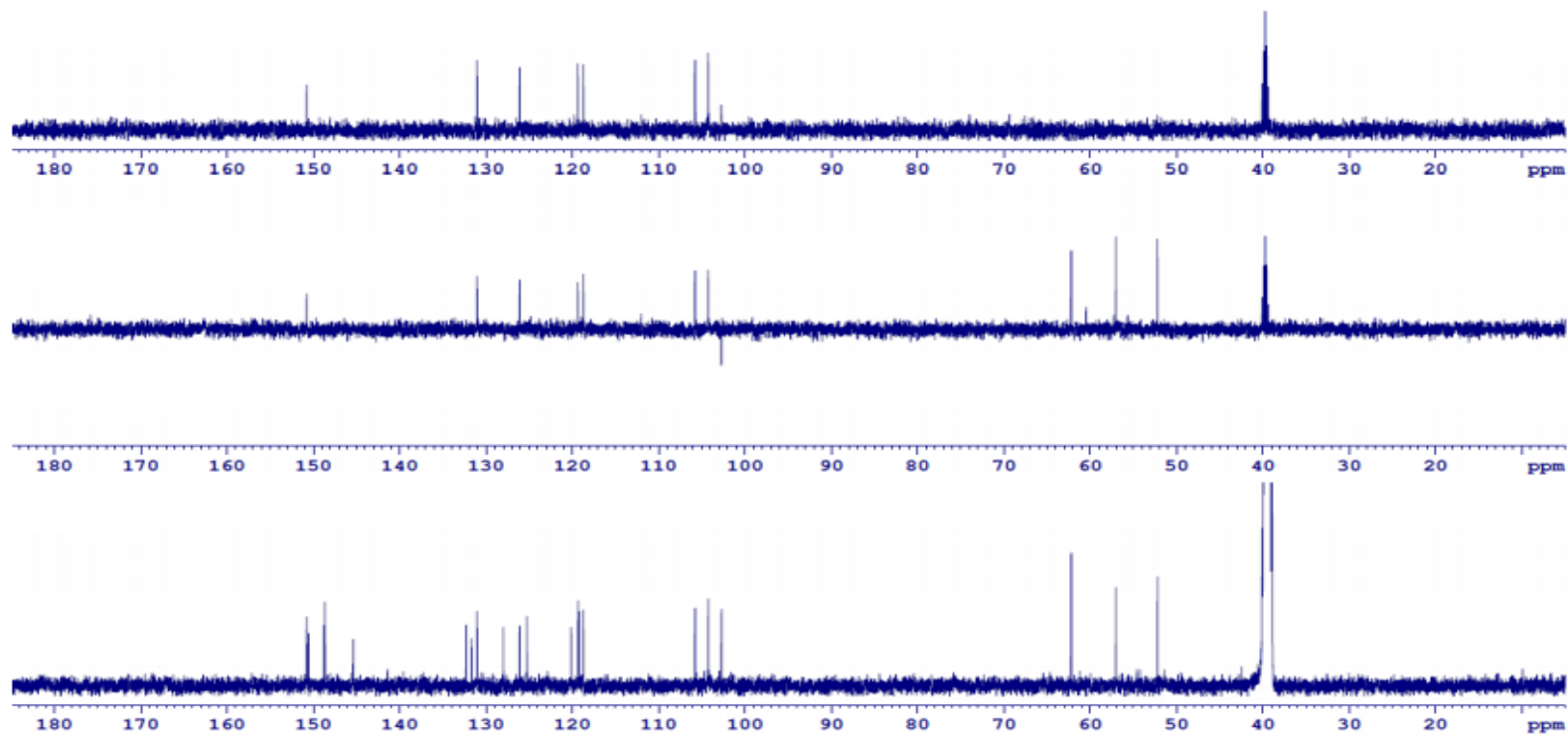


4.2:  $^{13}\text{C}$  NMR spectrum of compound 42



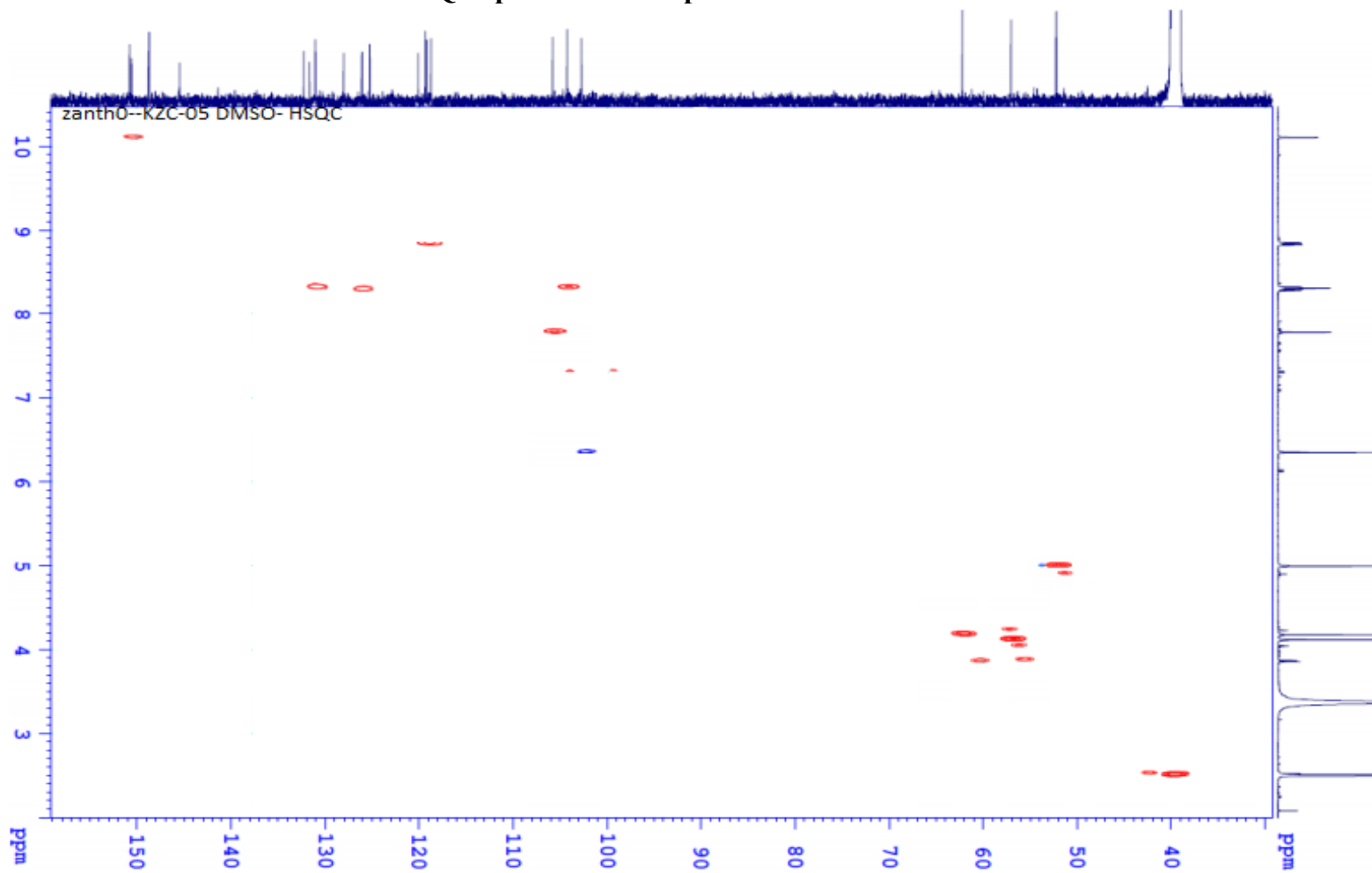
### 4.3: DEPT NMR spectrum of compound 42

Zanth-KZC\_05-DMSO DEPT

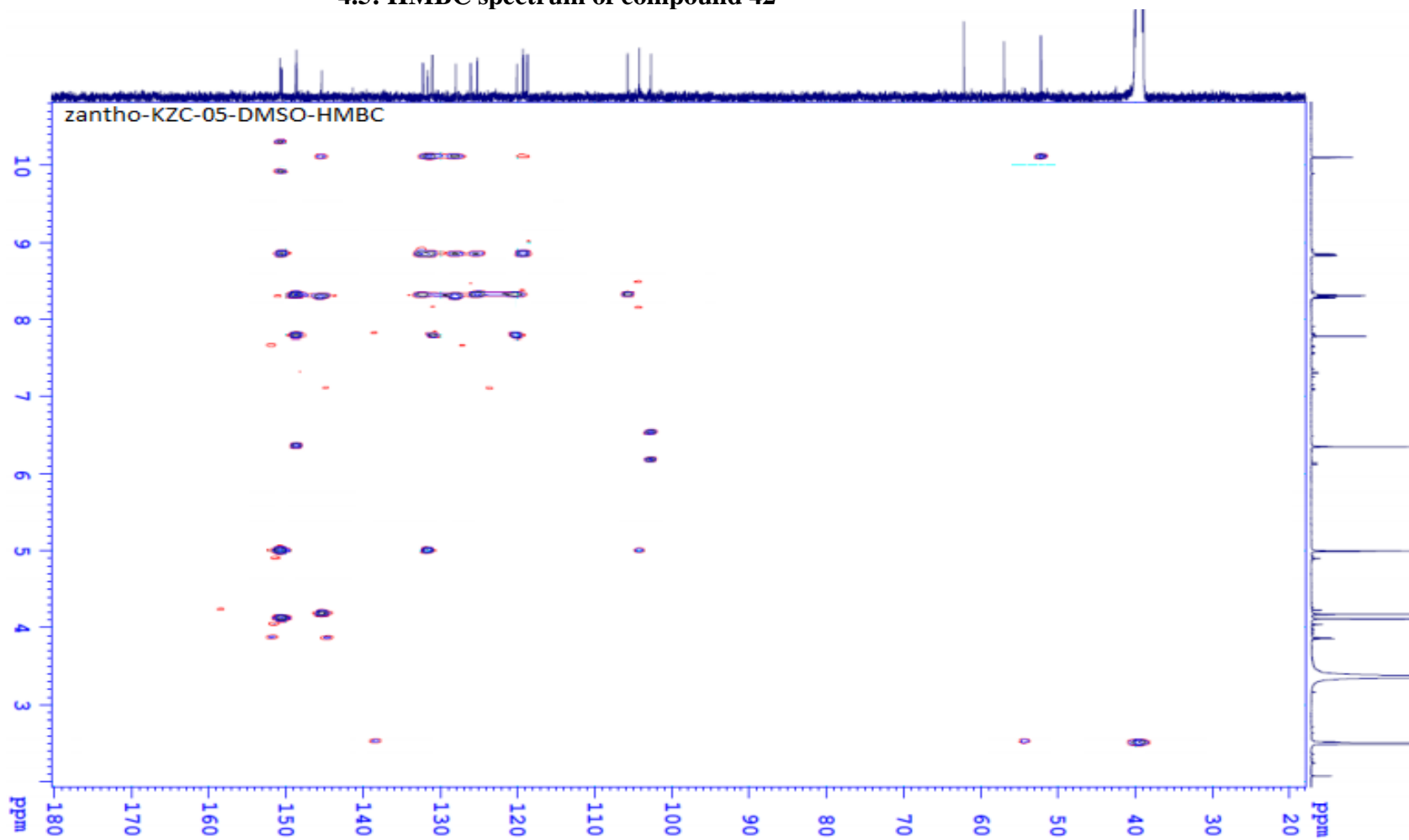




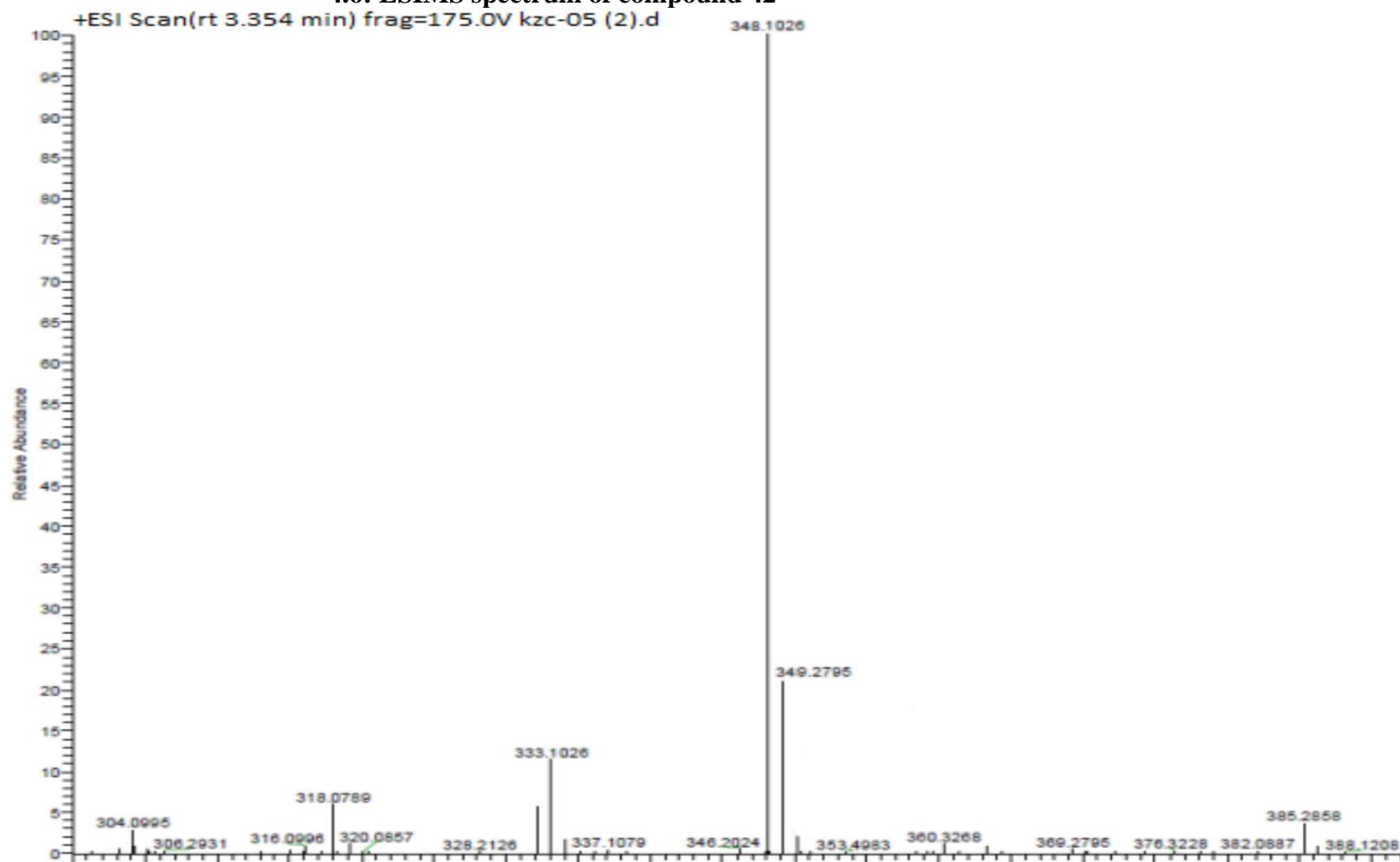
#### 4.4: HSQC spectrum of compound 42



4.5: HMBC spectrum of compound 42

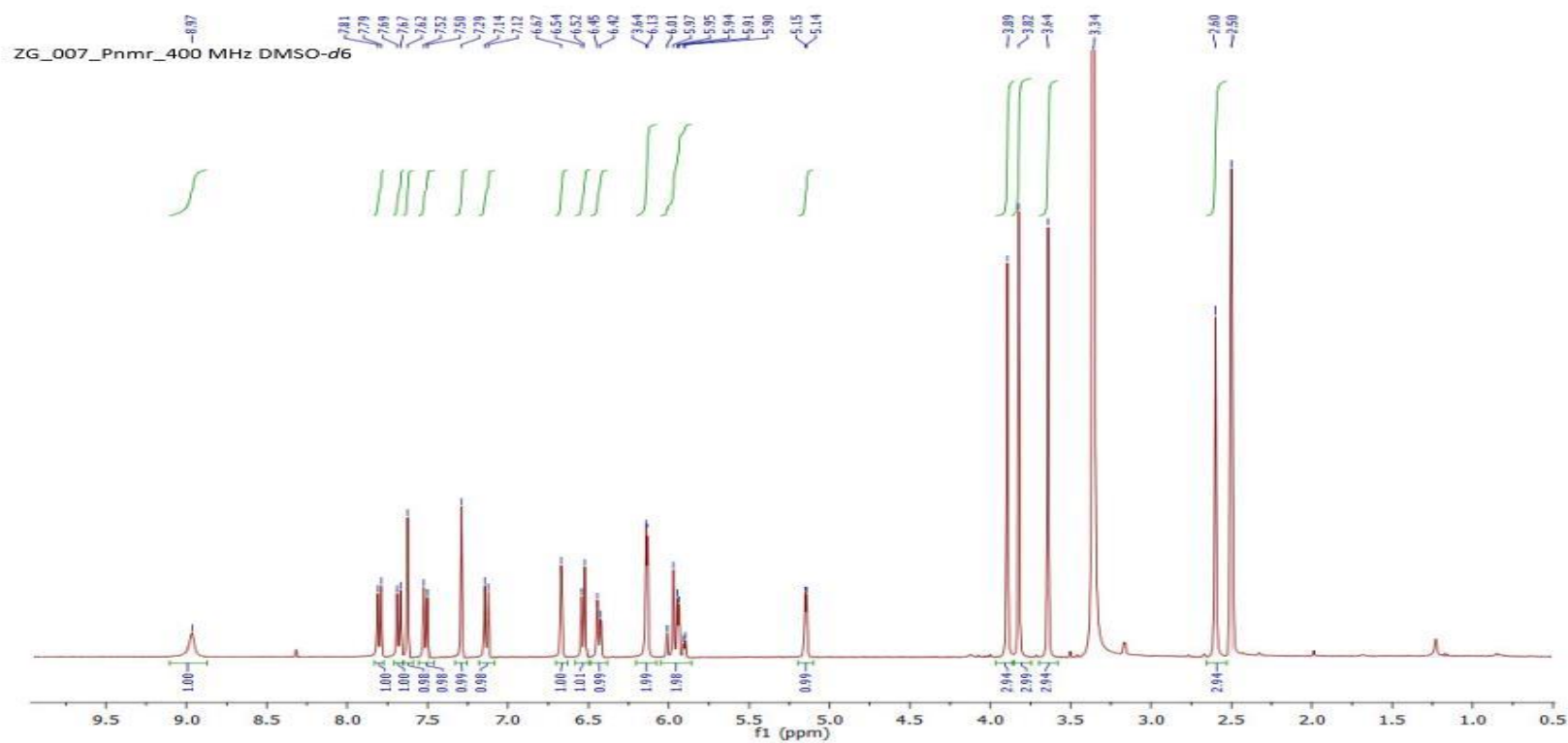


#### 4.6: ESIMS spectrum of compound 42

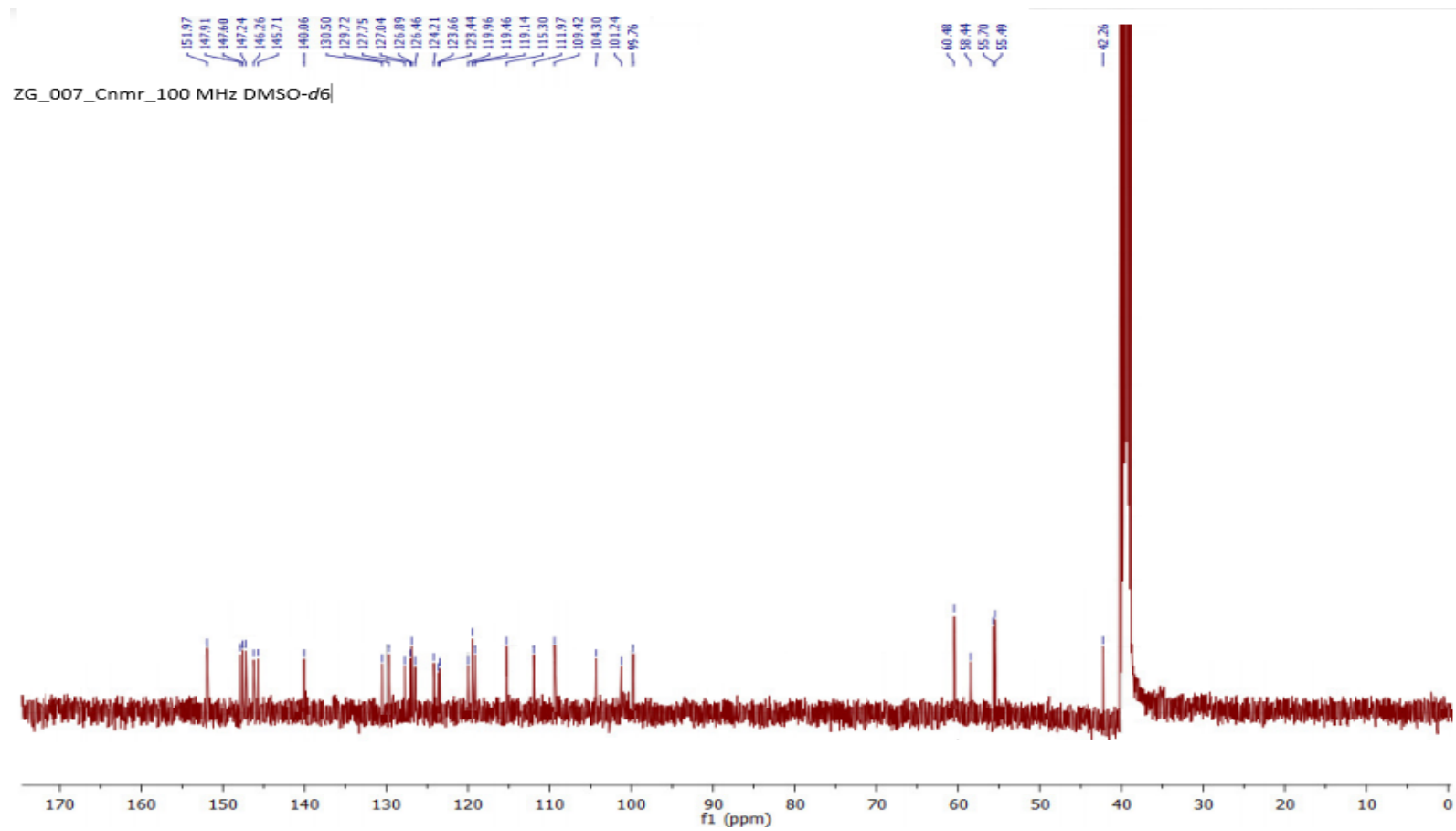


## 5. Spectra for Compound 43 (Zanthocapsine)

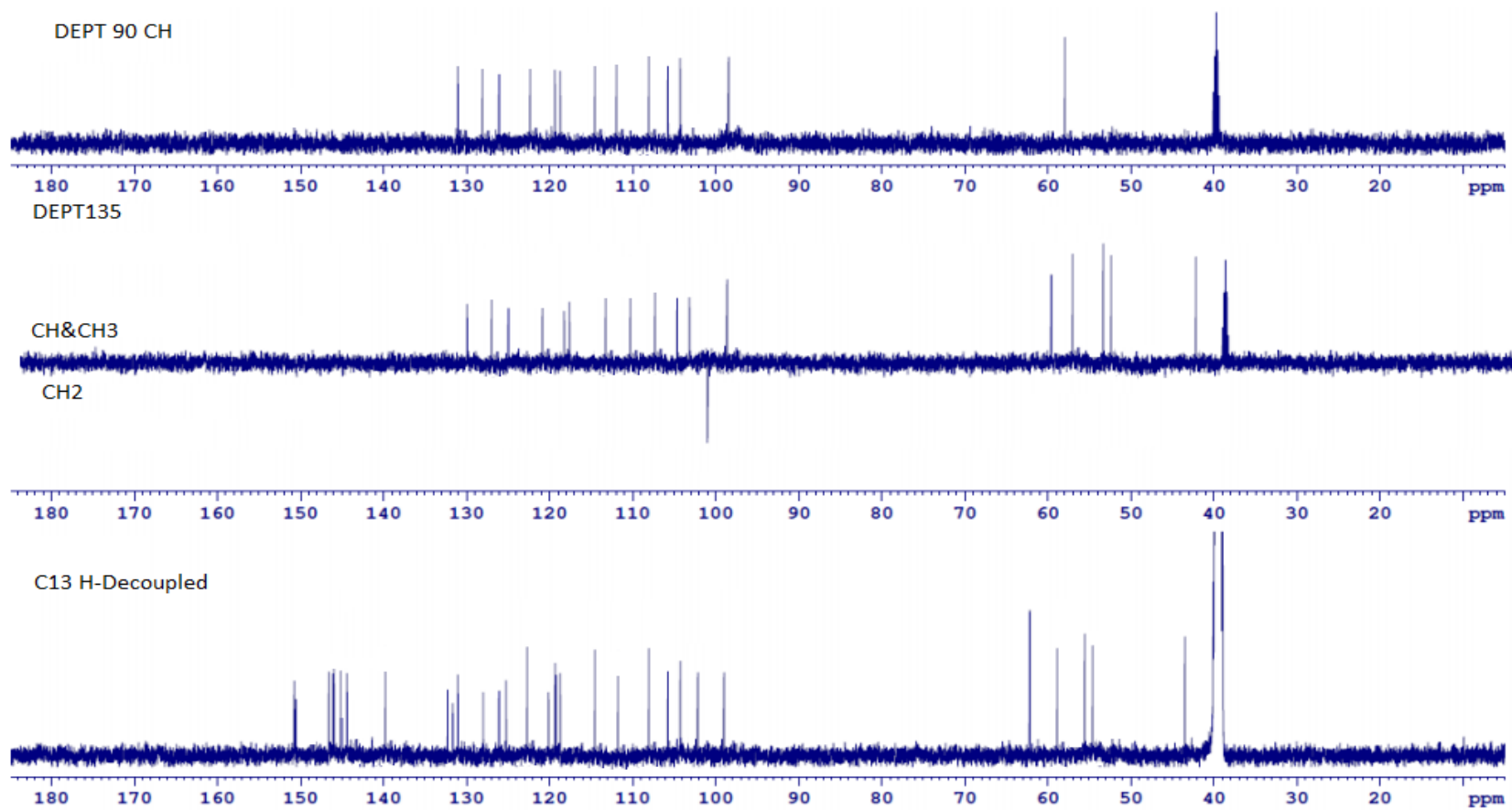
### 5.1: $^1\text{H}$ NMR spectrum of compound 43



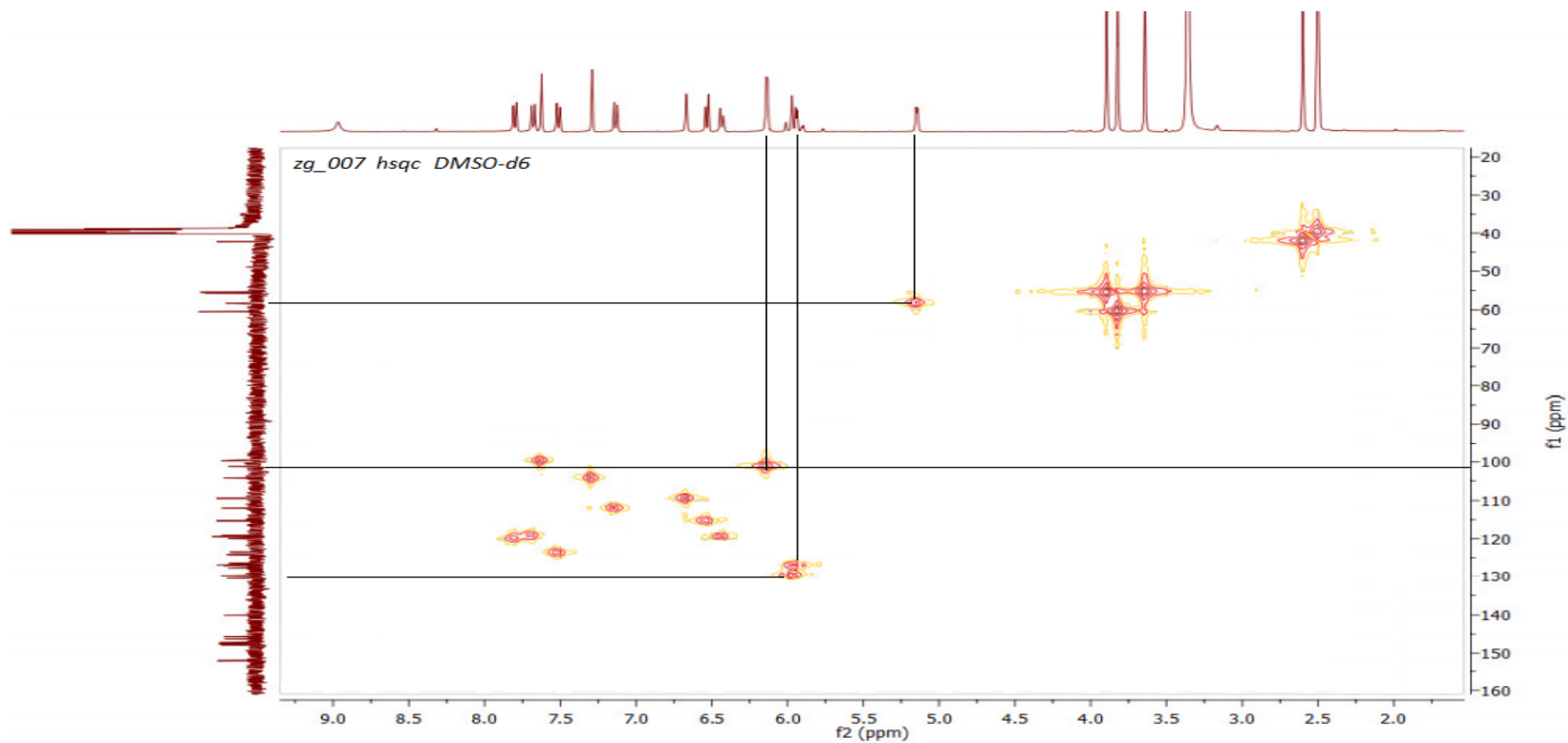
## 5.2: <sup>13</sup>C NMR spectrum of compound 43



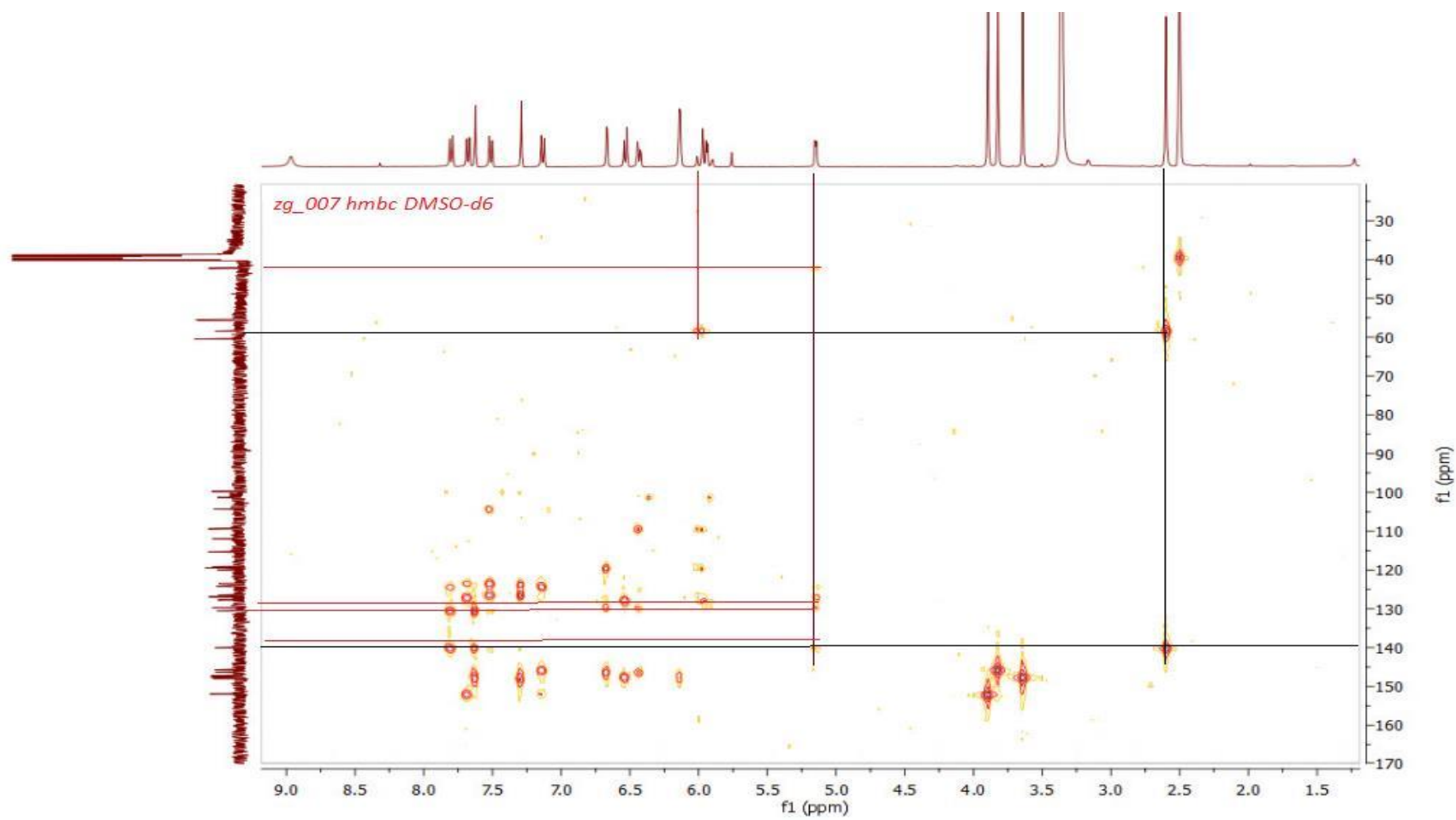
### 5.3: DEPT NMR spectrum of compound 43



### 5.4: HSQC spectrum of compound 43

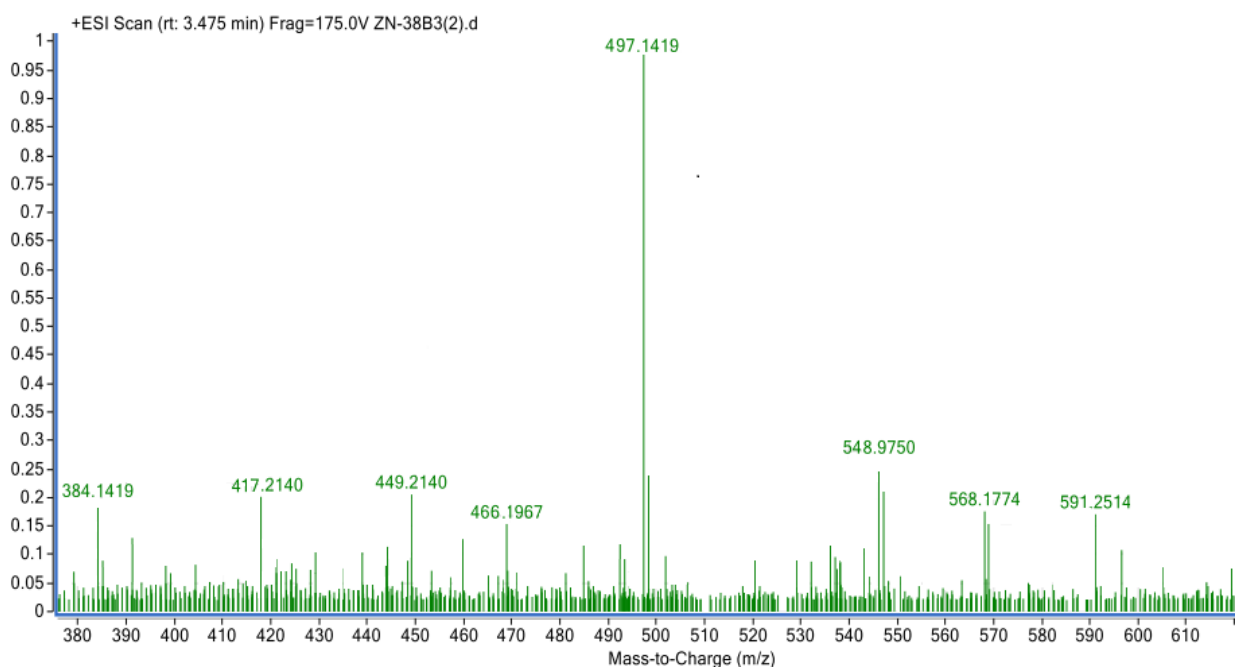


### 5.5: HMBC spectrum of compound 43



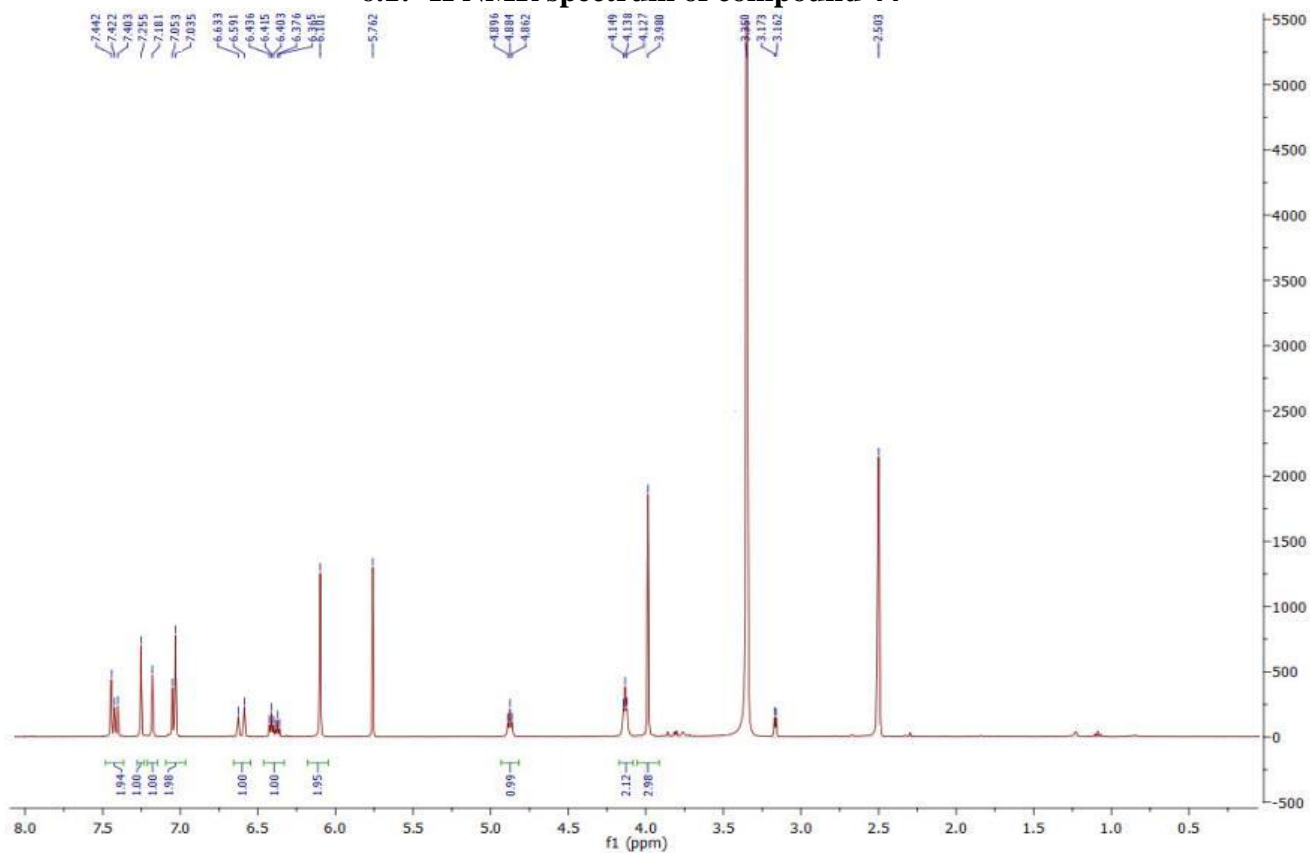


## 5.6: ESIMS spectrum of compound 43

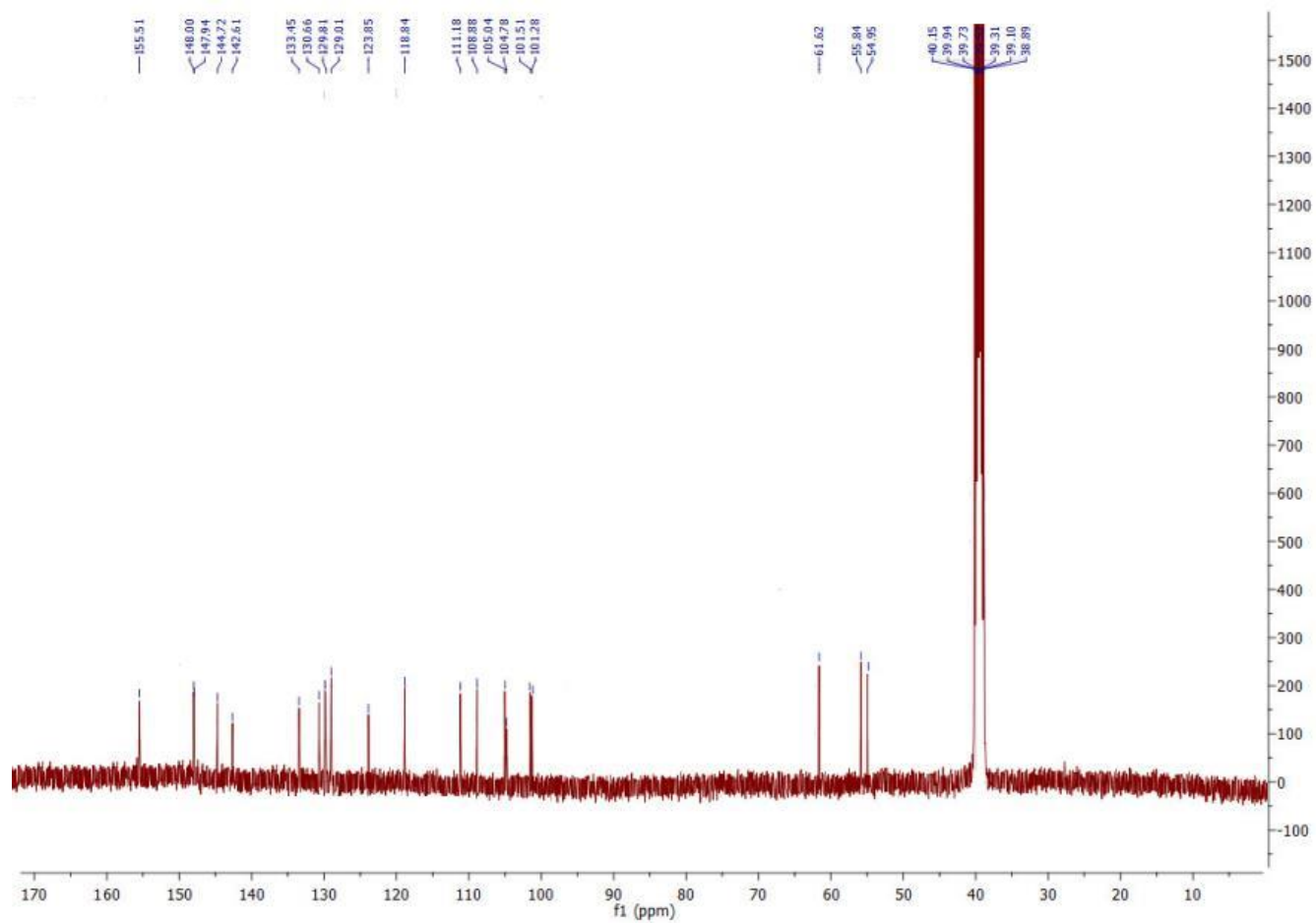


## 6. Spectra for Compound 44 (Zanthocapsol)

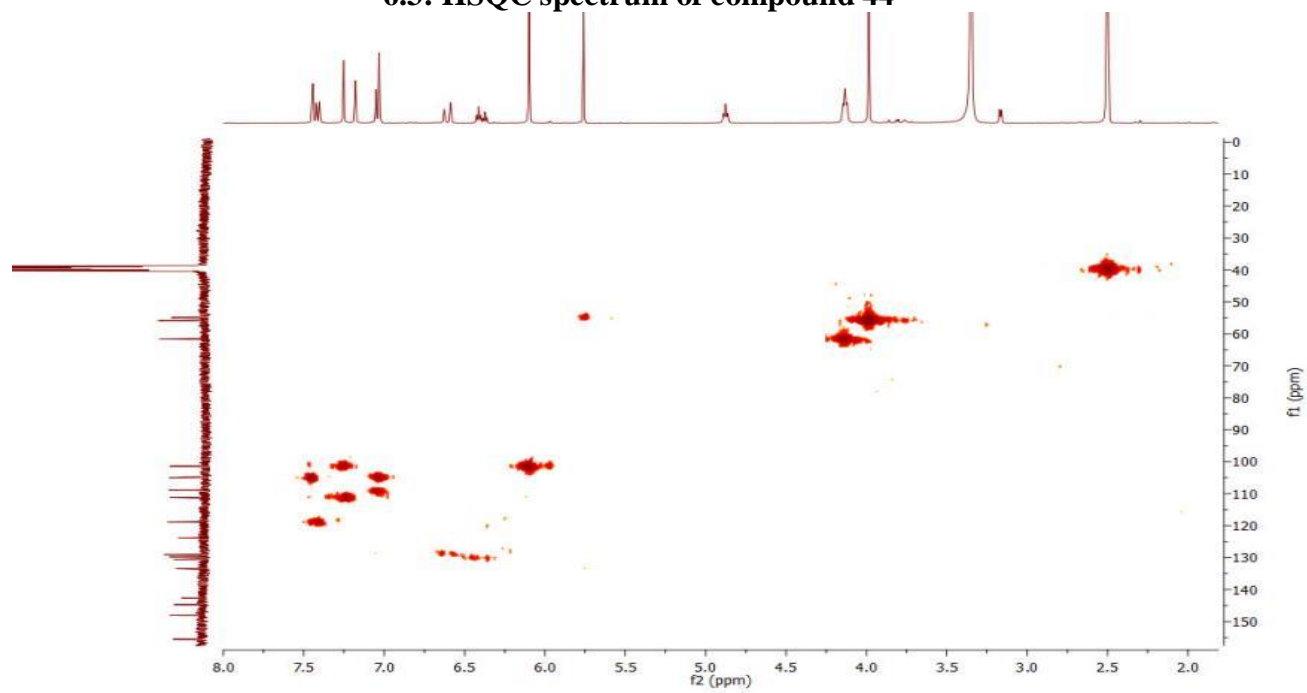
### 6.1: $^1\text{H}$ NMR spectrum of compound 44



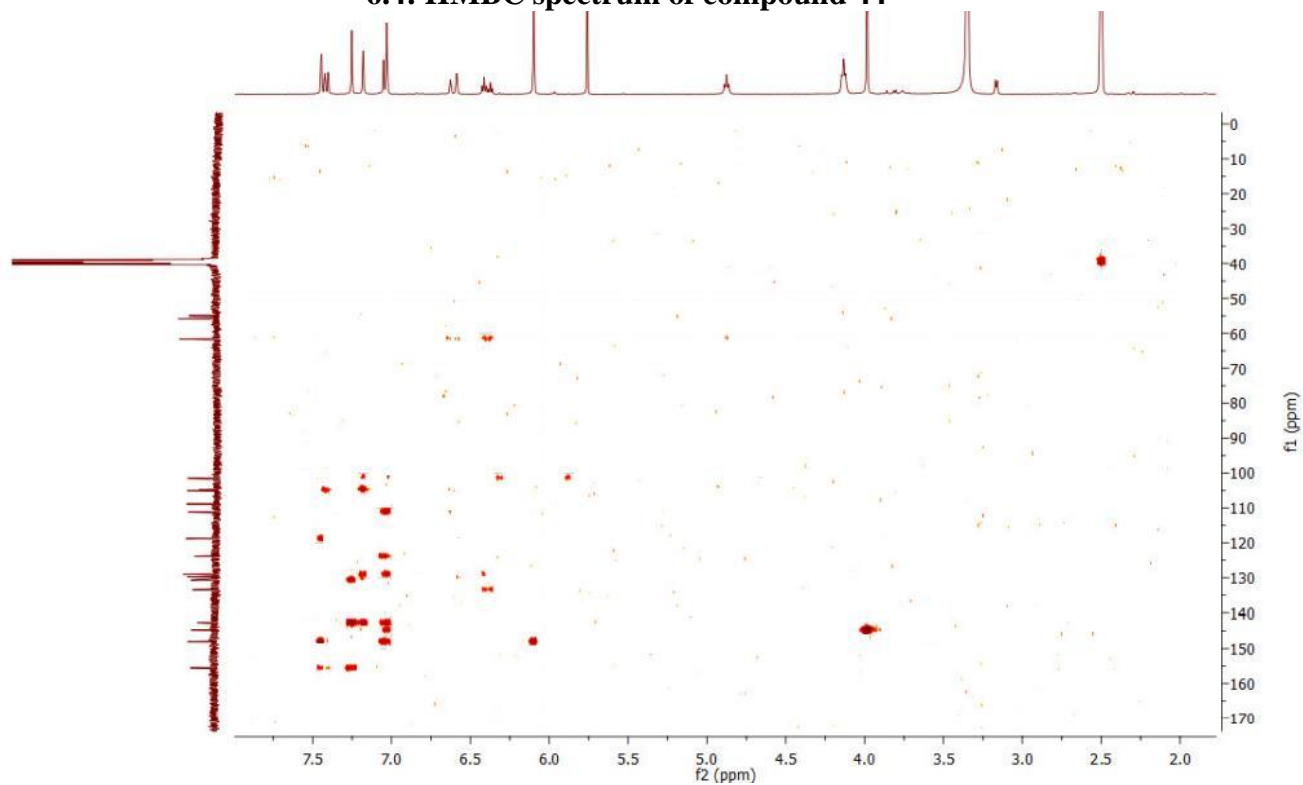
### 6.2: $^{13}\text{C}$ NMR spectrum of compound 44



### 6.3: HSQC spectrum of compound 44

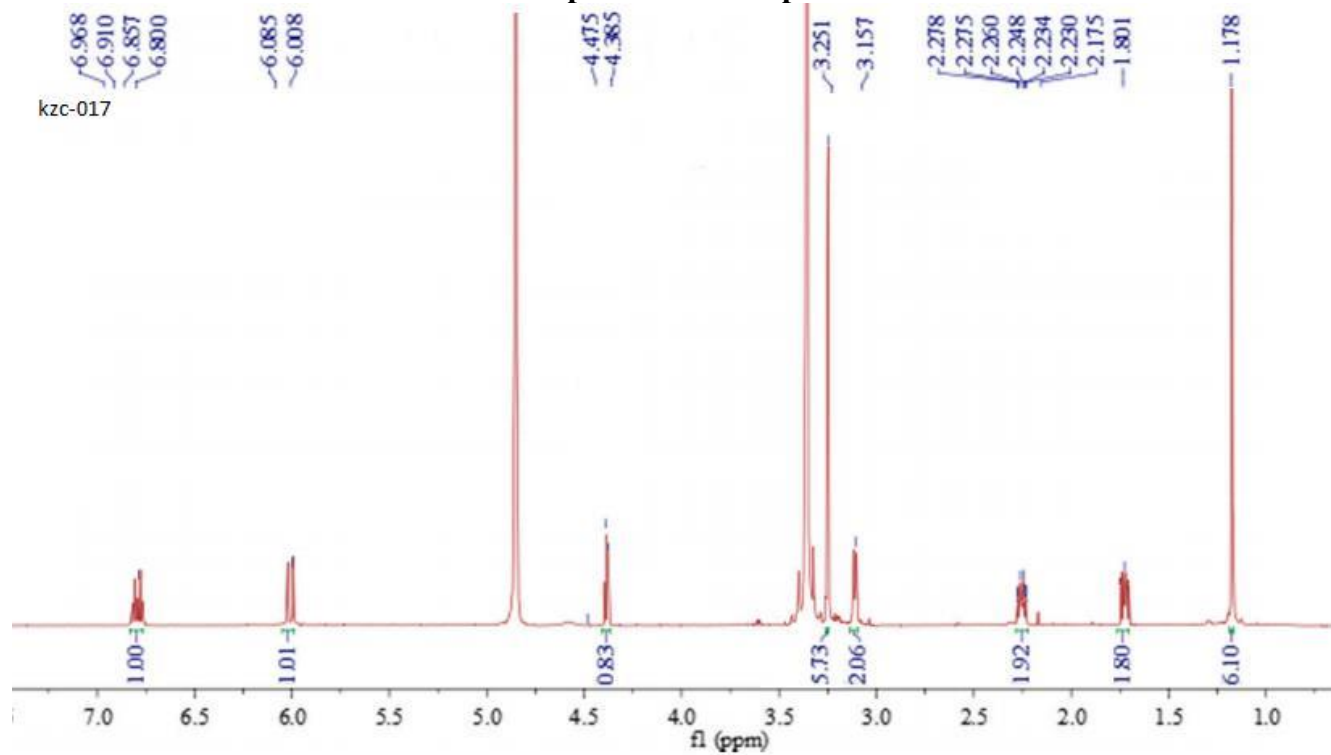


### 6.4: HMBC spectrum of compound 44

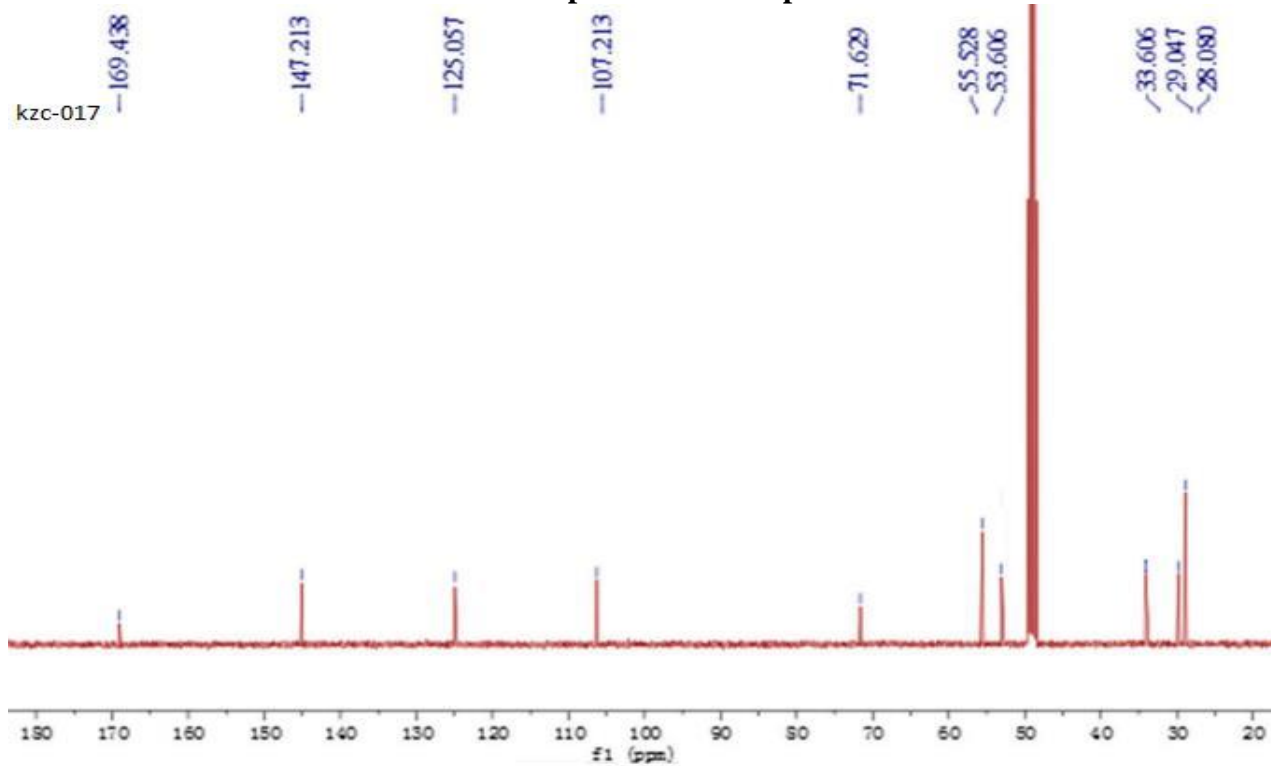


## 7. Spectra for Compound 45 (Zanthoamide D)

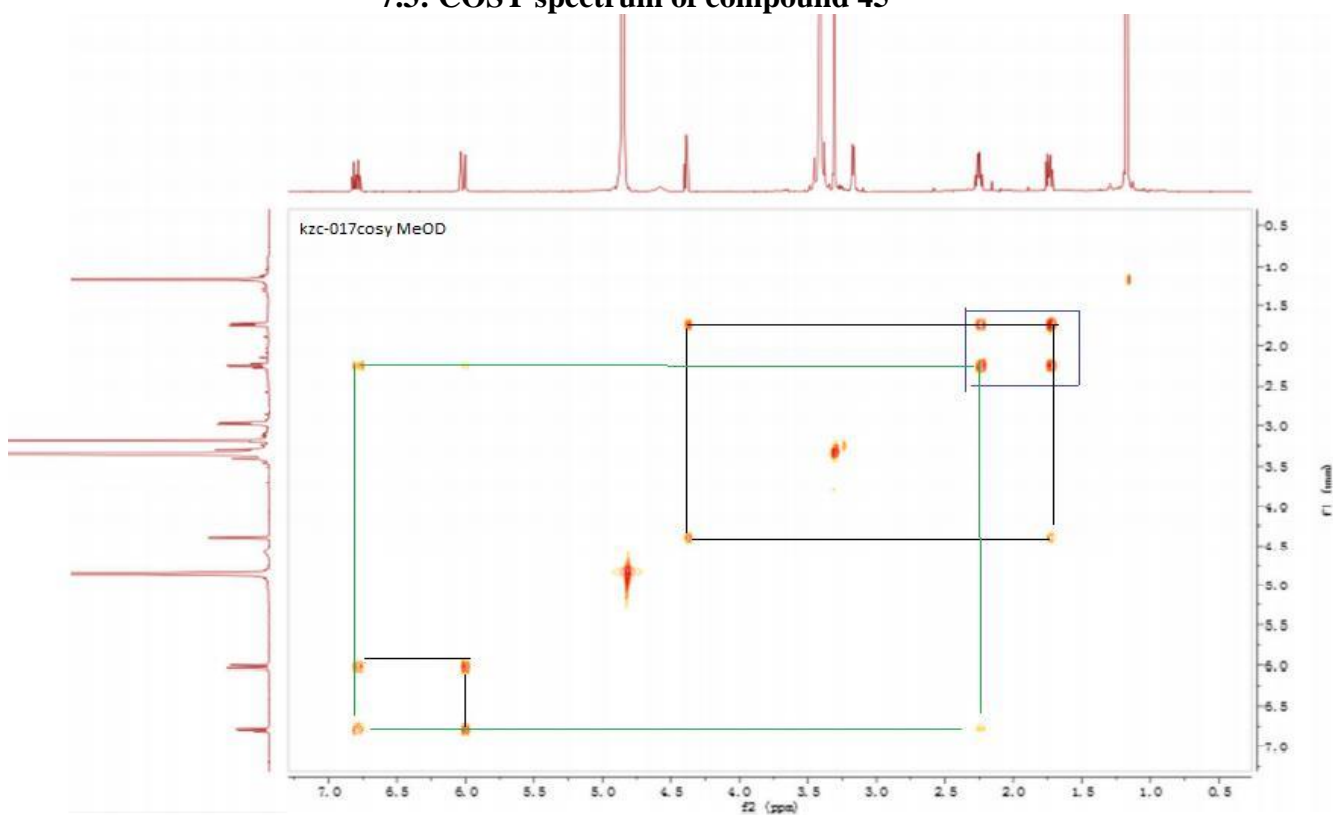
### 7.1: $^1\text{H}$ NMR spectrum of compound 45



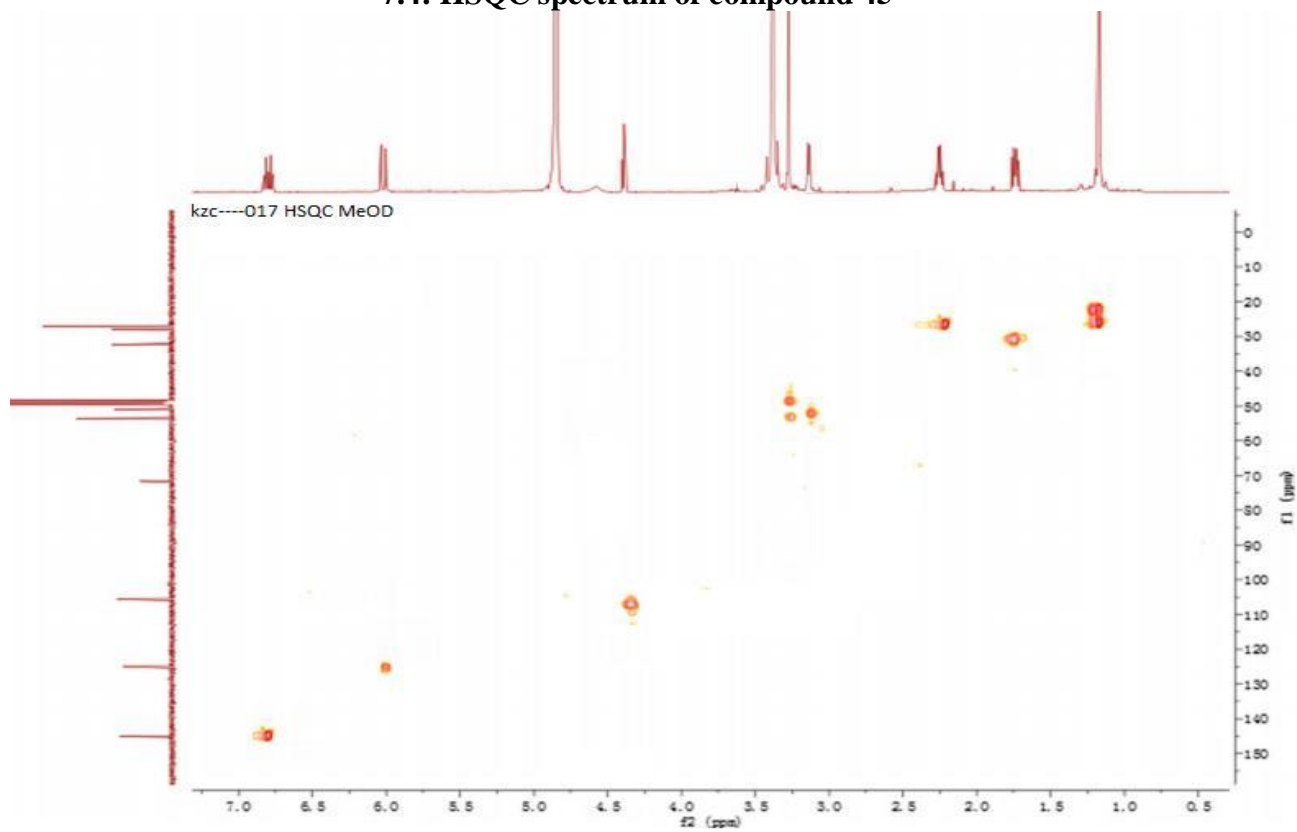
### 7.2: $^{13}\text{C}$ NMR spectrum of compound 45



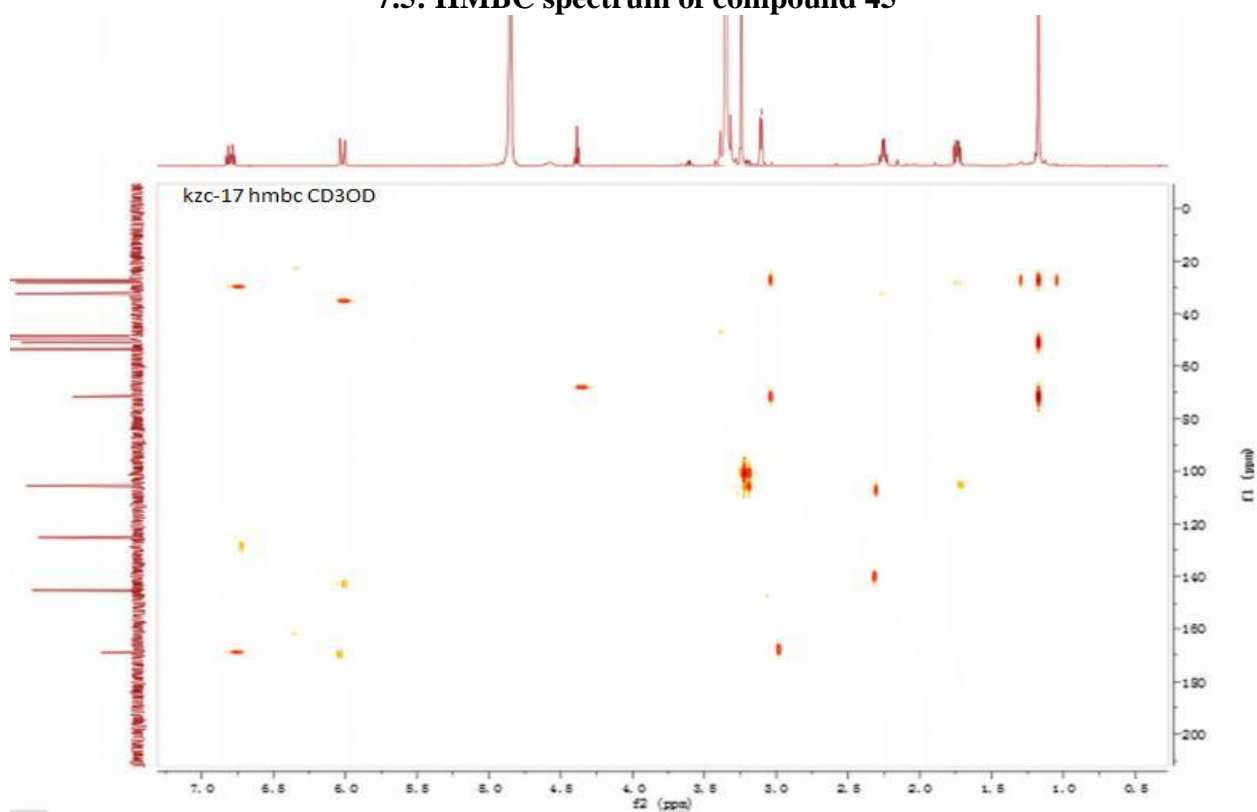
### 7.3: COSY spectrum of compound 45



### 7.4: HSQC spectrum of compound 45



### 7.5: HMBC spectrum of compound 45

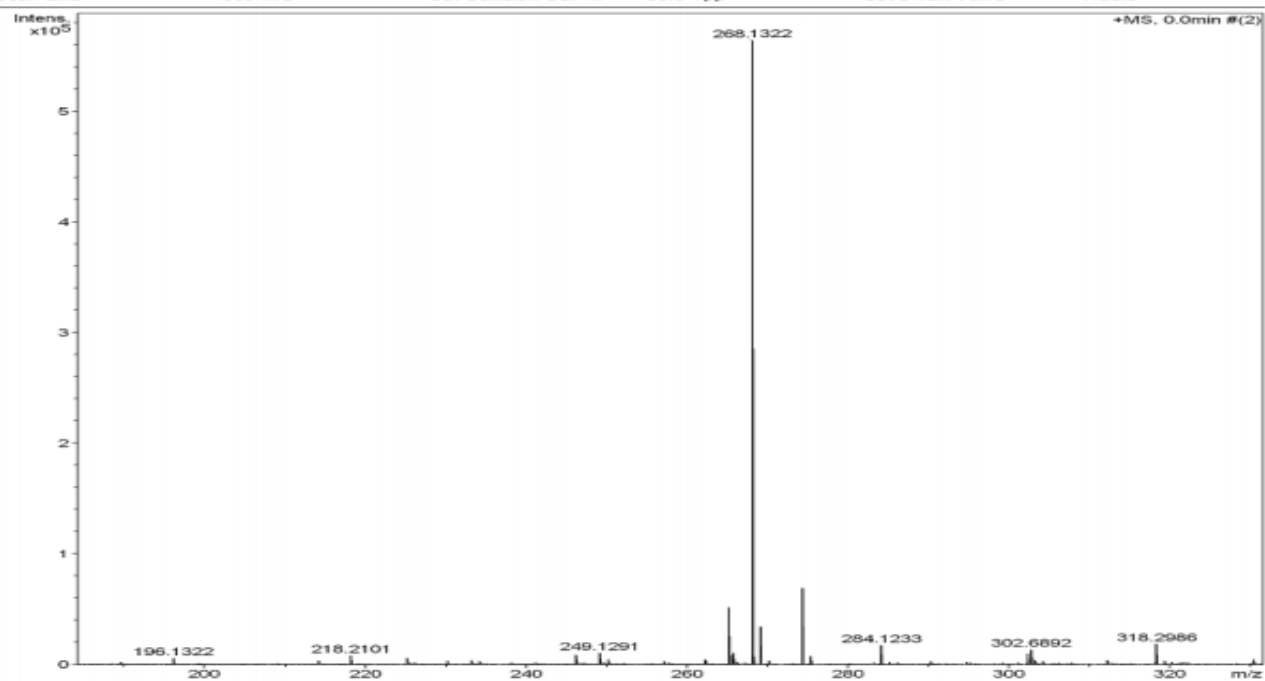


## 7.6: ESIMS spectrum of compound 45

### ESIMS Report

<b>Analysis Info</b>		<b>Acquisition Date</b>	3/22/20 21 10:12:03 AM		
<b>Analysis Name</b>	D:\Data\SFM\20 21\0722\j-4-1.d	<b>Operator</b>	SFM		
<b>Method</b>	pos_low_1000_0531.m	<b>Instrument</b>	maXis		
<b>Sample Name</b>	kzc_17		10103		

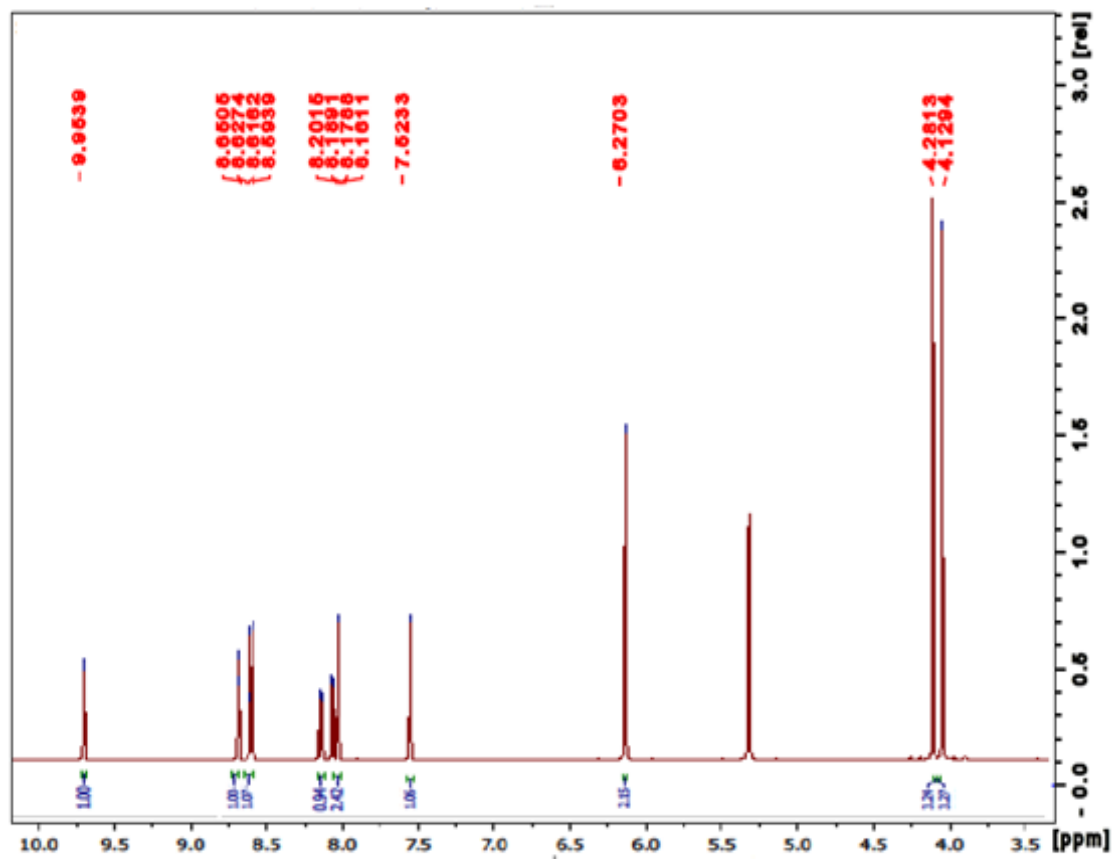
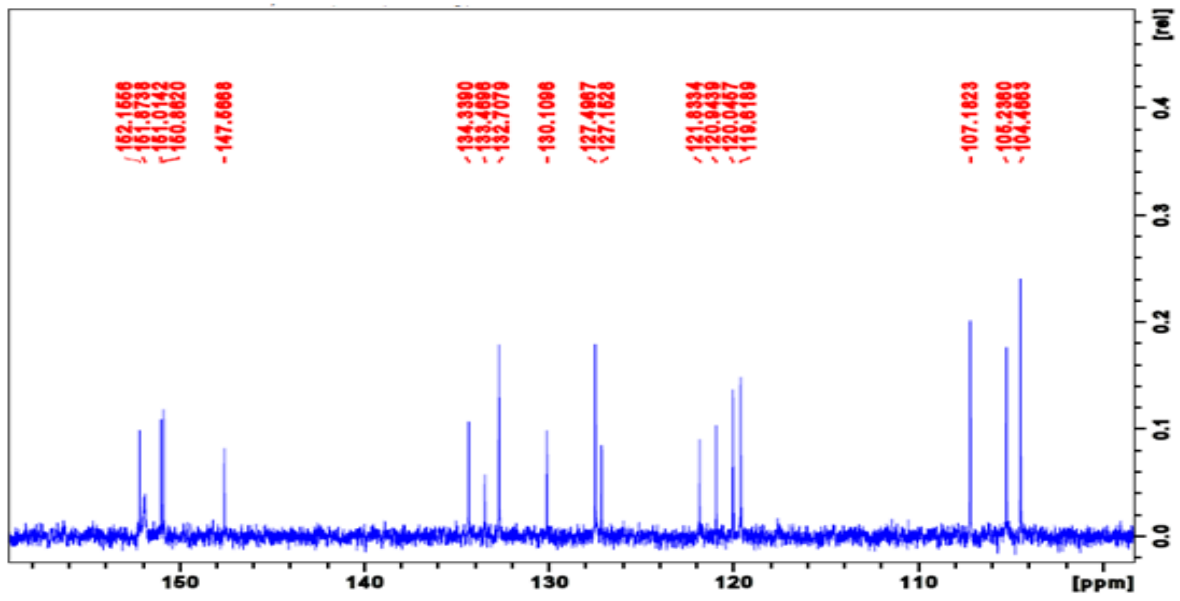
<b>Acquisition Parameter</b>					
<b>Source Type</b>	ESI	<b>Ion Polarity</b>	Positive	<b>Set Nebulizer</b>	0.4 Bar
<b>Focus</b>	Not active	<b>Set Capillary</b>	4500 V	<b>Set Dry Heater</b>	180 °C
<b>Scan Begin</b>	100 m/z	<b>Set End Plate Offset</b>	-500 V	<b>Set Dry Gas</b>	4.0 l/min
<b>Scan End</b>	1000 m/z	<b>Set Collision Cell RF</b>	80.0 Vpp	<b>Set Divert Valve</b>	Waste



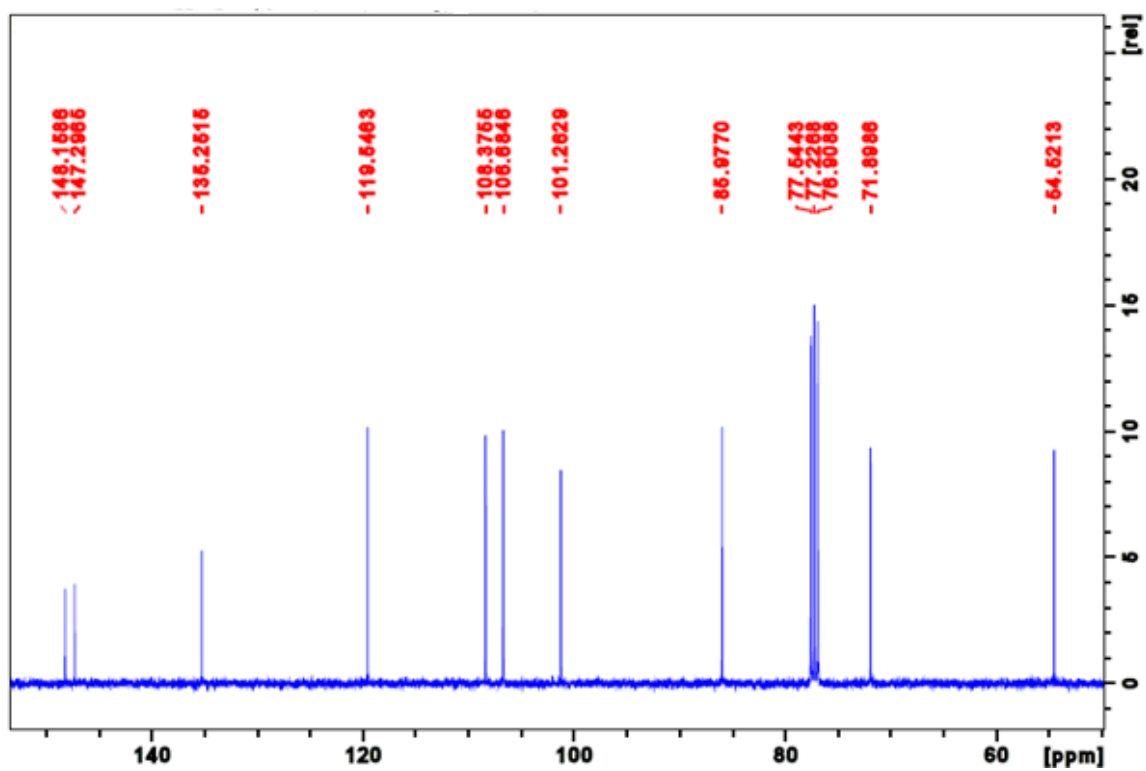
Bruker Compass DataAnalysis 4.0

printed: 3/22/20 21 10:12:03 AM

8. Spectra for Compound 9 (Norchelerythrine)  
 8.1: <sup>1</sup>H and <sup>13</sup>C NMR spectrum of compound 9



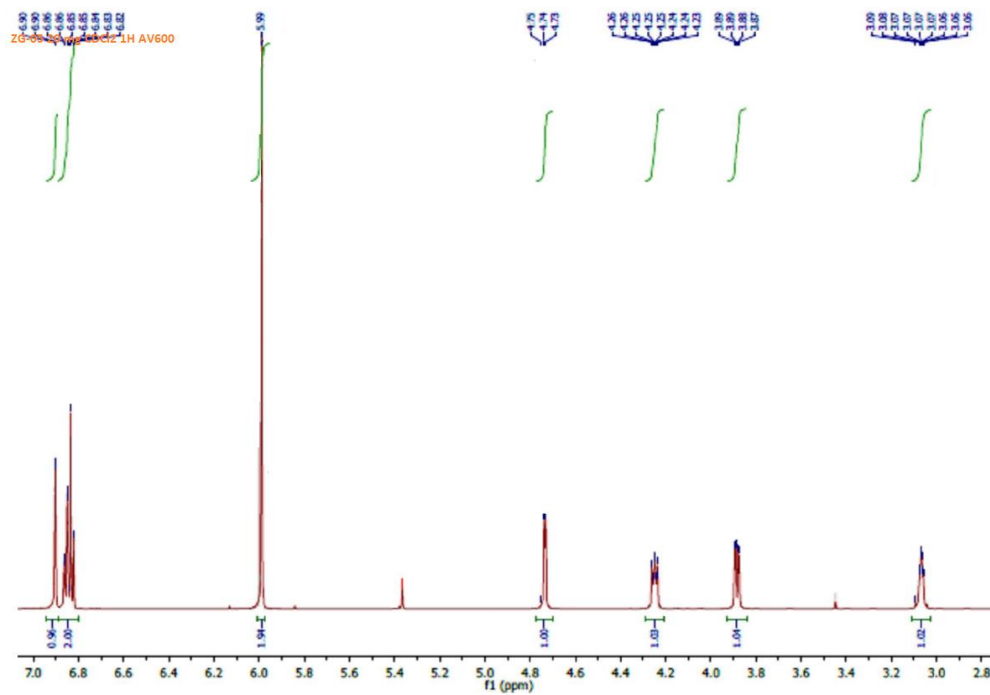
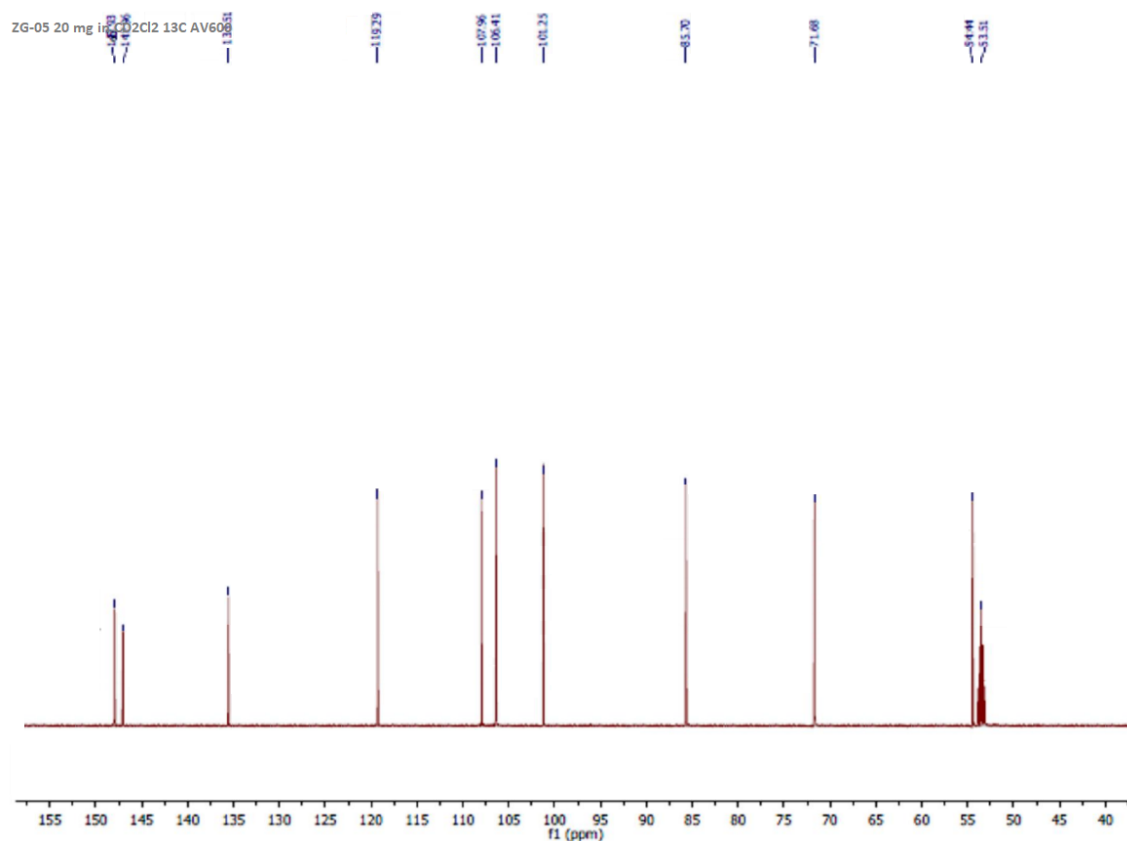
9. Spectra for Compound 10 (2,3-epoxy-6,7-methylenedioxy coniferyl alcohol)  
9.1:  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectrum of compound 10





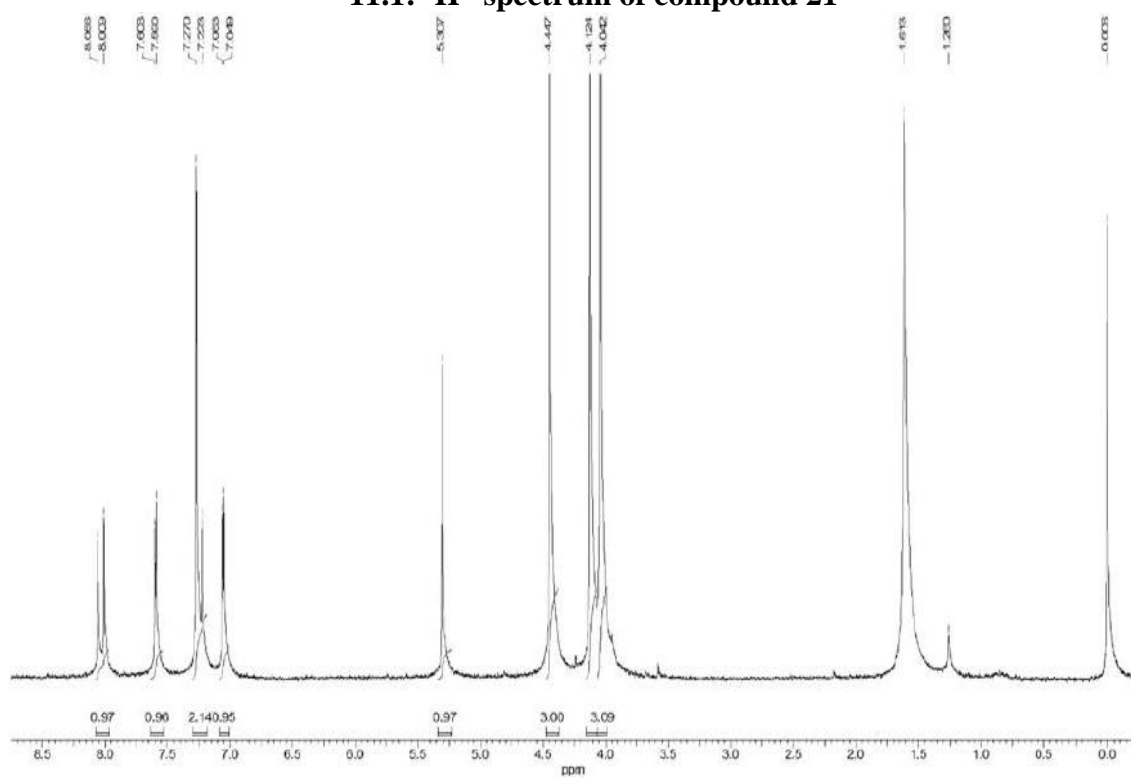
## 10. Spectra for Compound 46 (Sesamine)

### 10.1: $^1\text{H}$ and $^{13}\text{C}$ NMR spectrum of compound 46

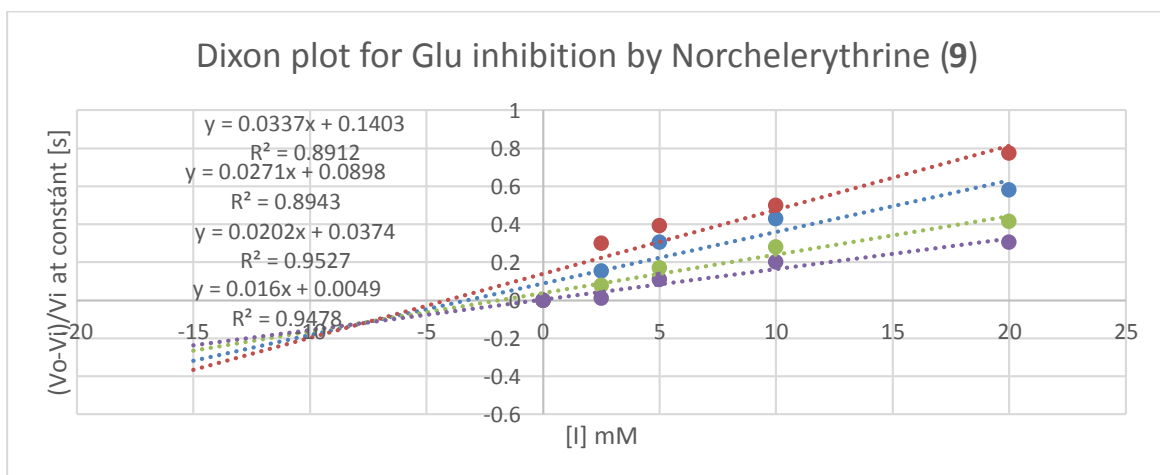
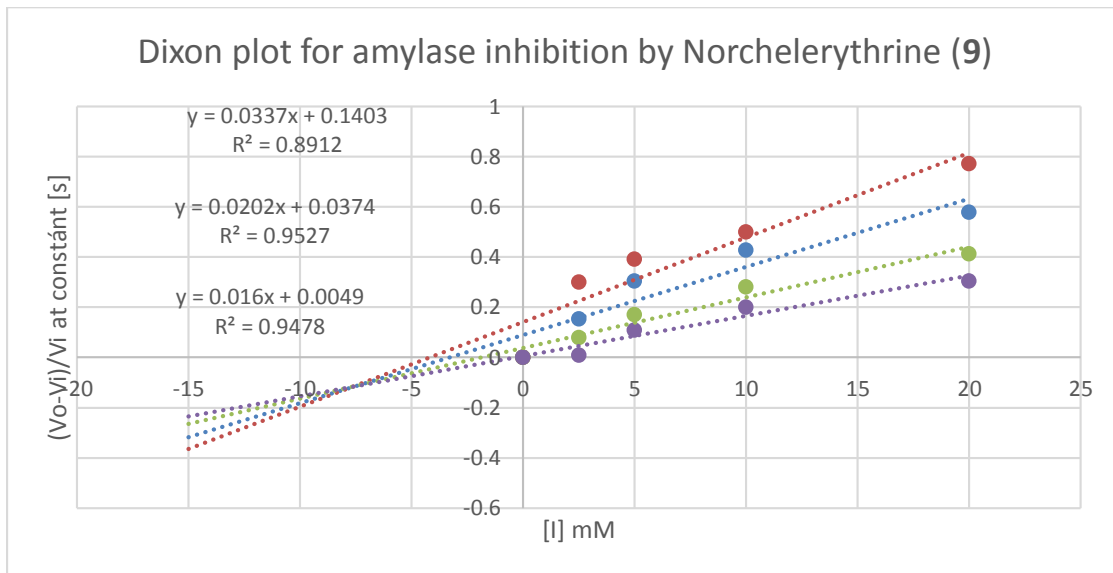
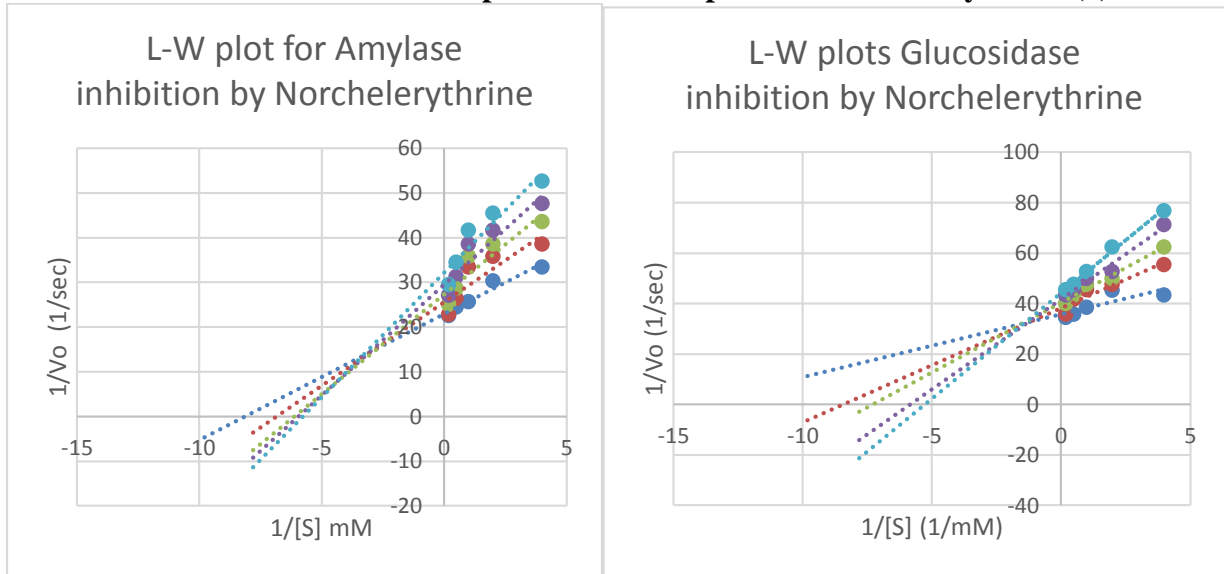


## 11. Spectra for Compound 21 (Skimmianine)

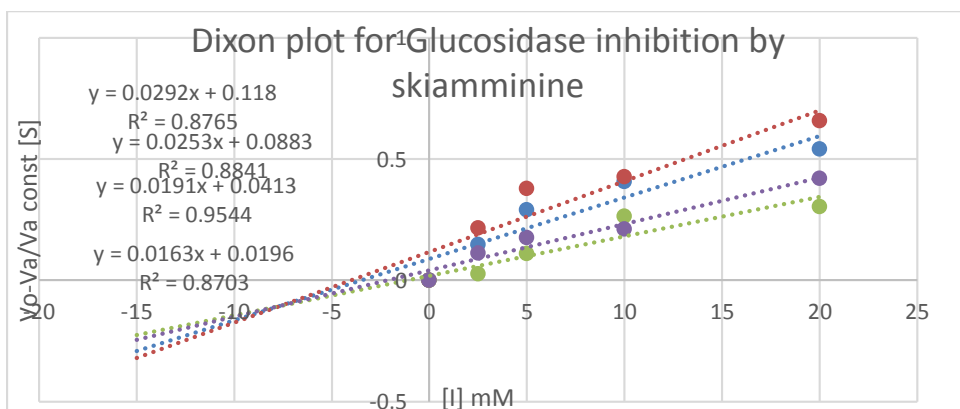
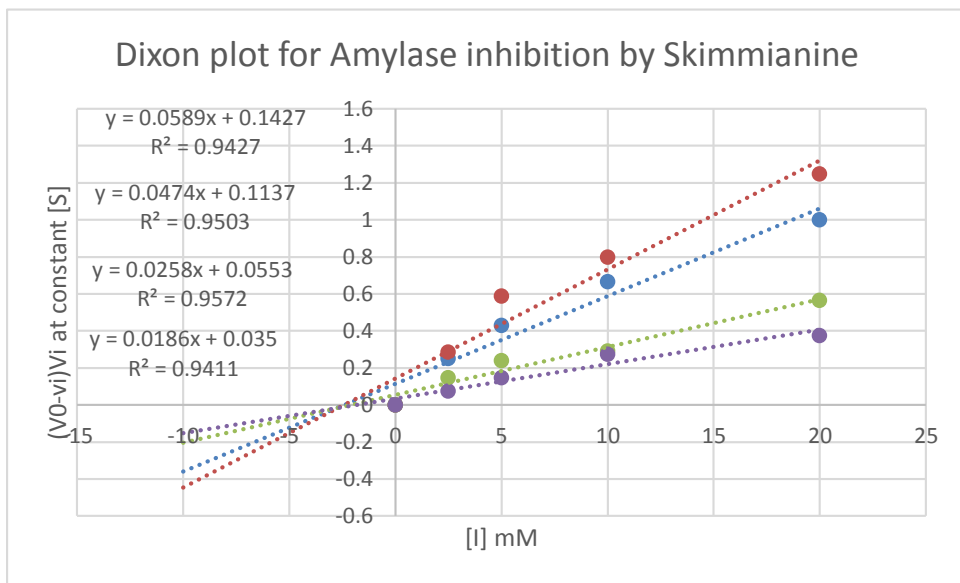
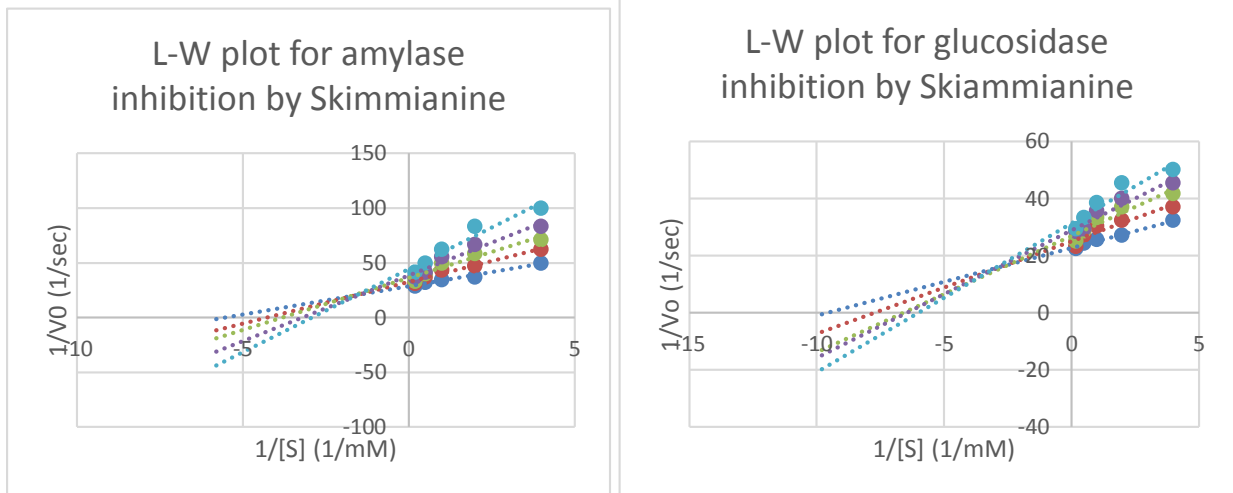
### 11.1: $^1\text{H}$ spectrum of compound 21



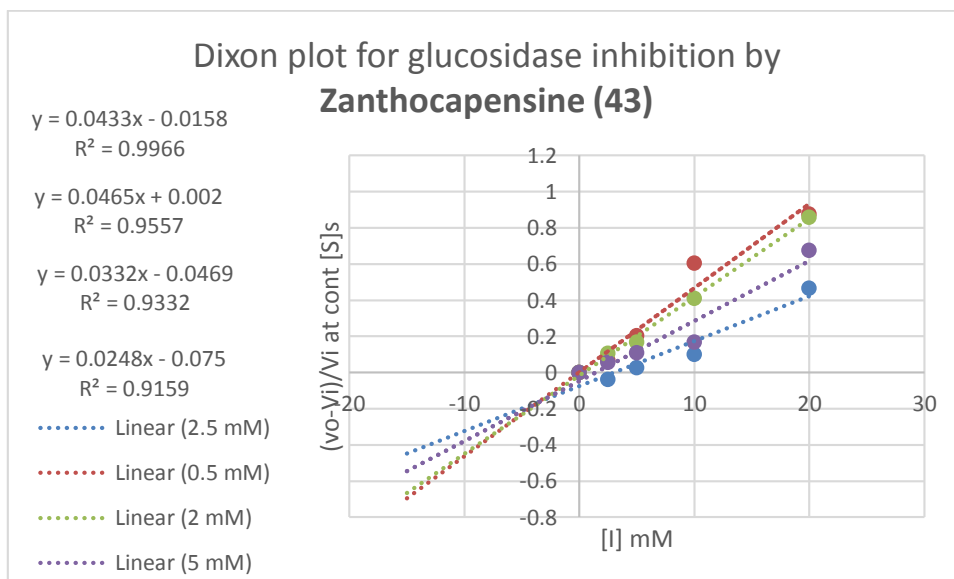
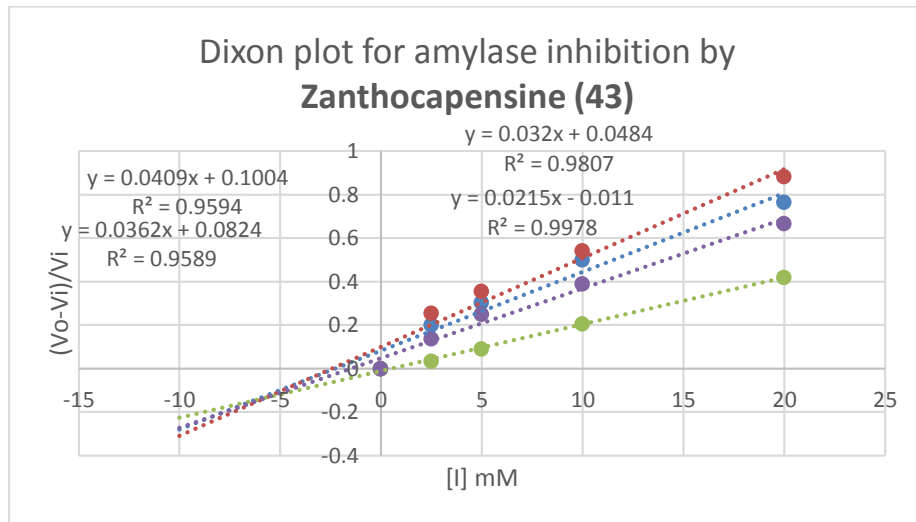
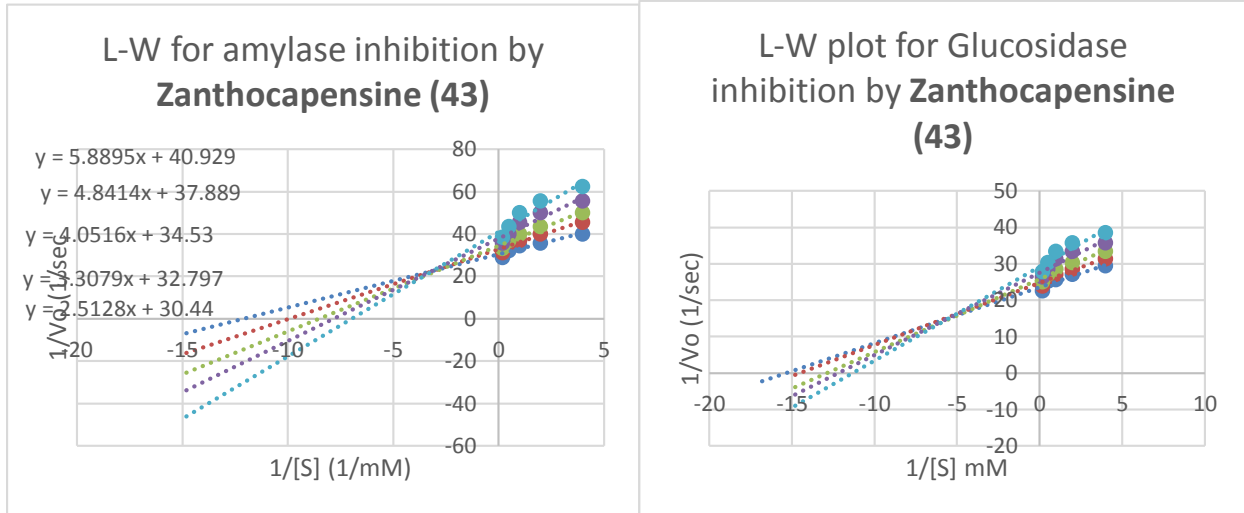
**12. Lineweaver-Burk plots and Dixon secondary plots**  
**12.1: Lineweaver-Burk plots and Dixon plots for Norchelerythrine (9)**



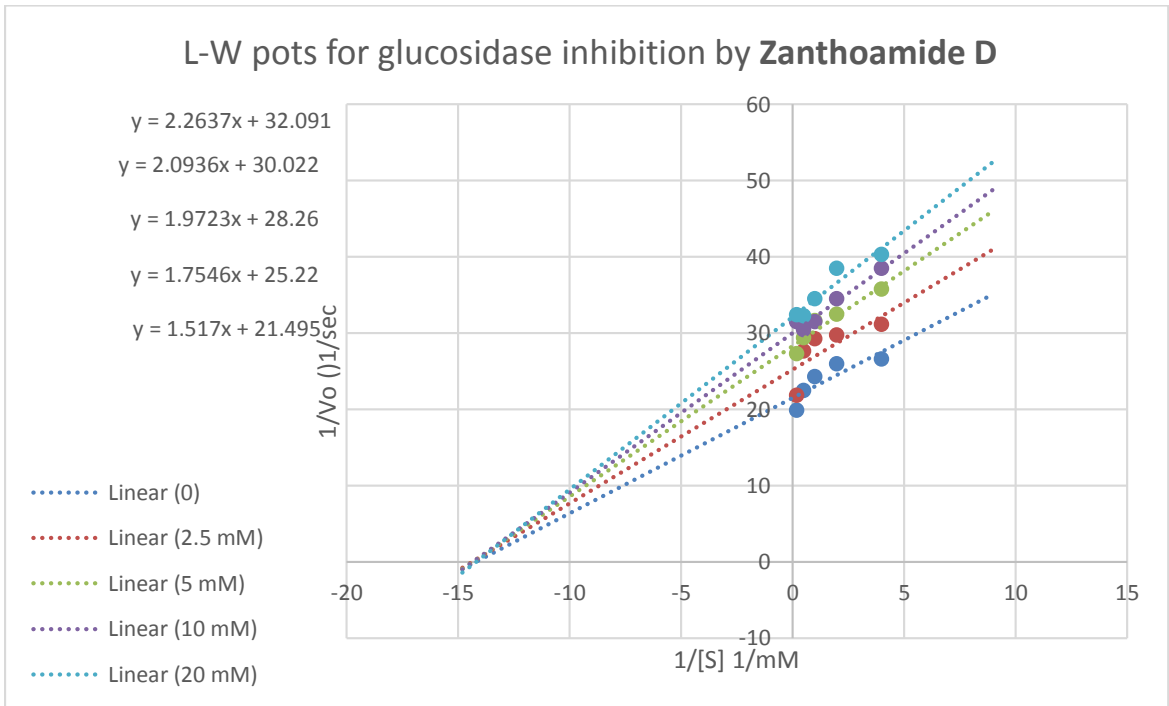
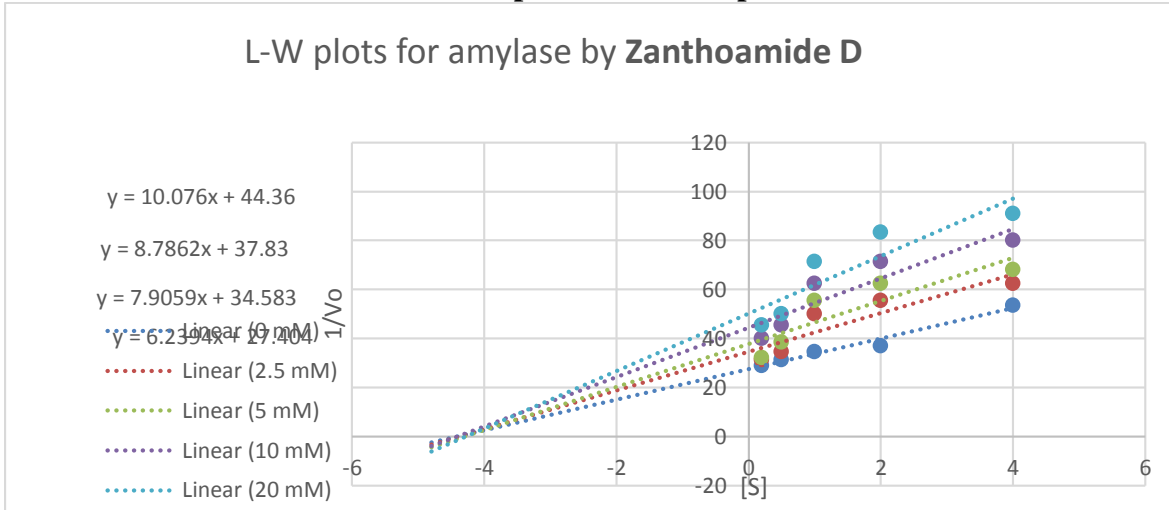
### 12.2: Lineweaver-Burk plots and Dixon plots for Skimmianine (21)



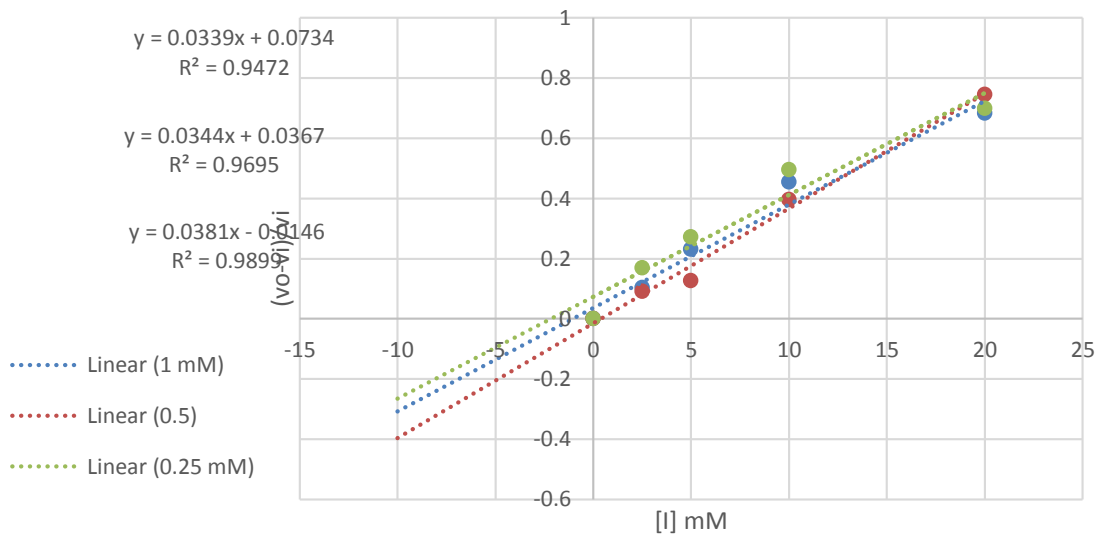
### 12.3: Lineweaver-Burk plots and Dixon plots for Zanthocapsine (43)



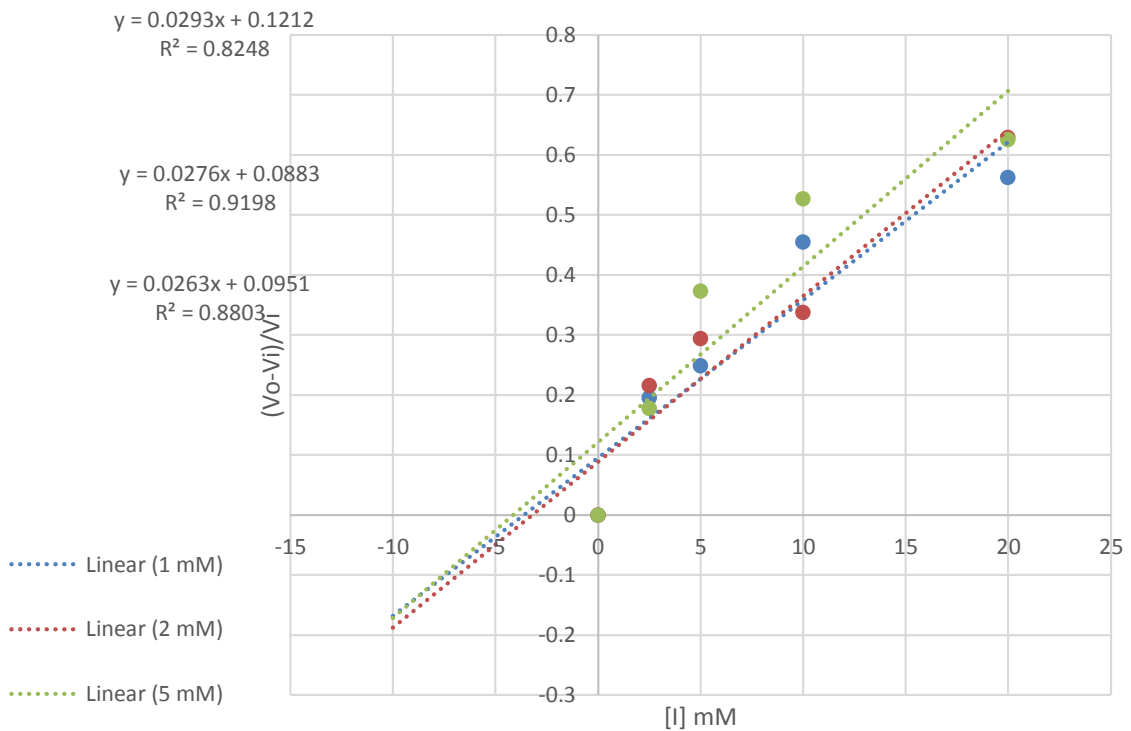
### 12.4: Lineweaver-Burk plots and Dixon plots for Zanthoamide D



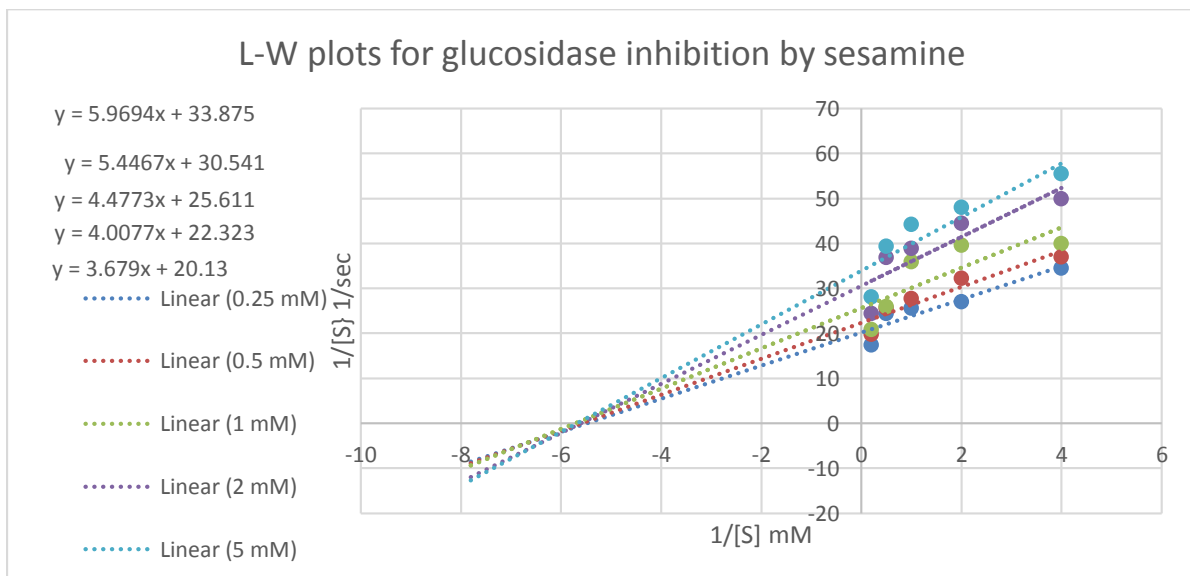
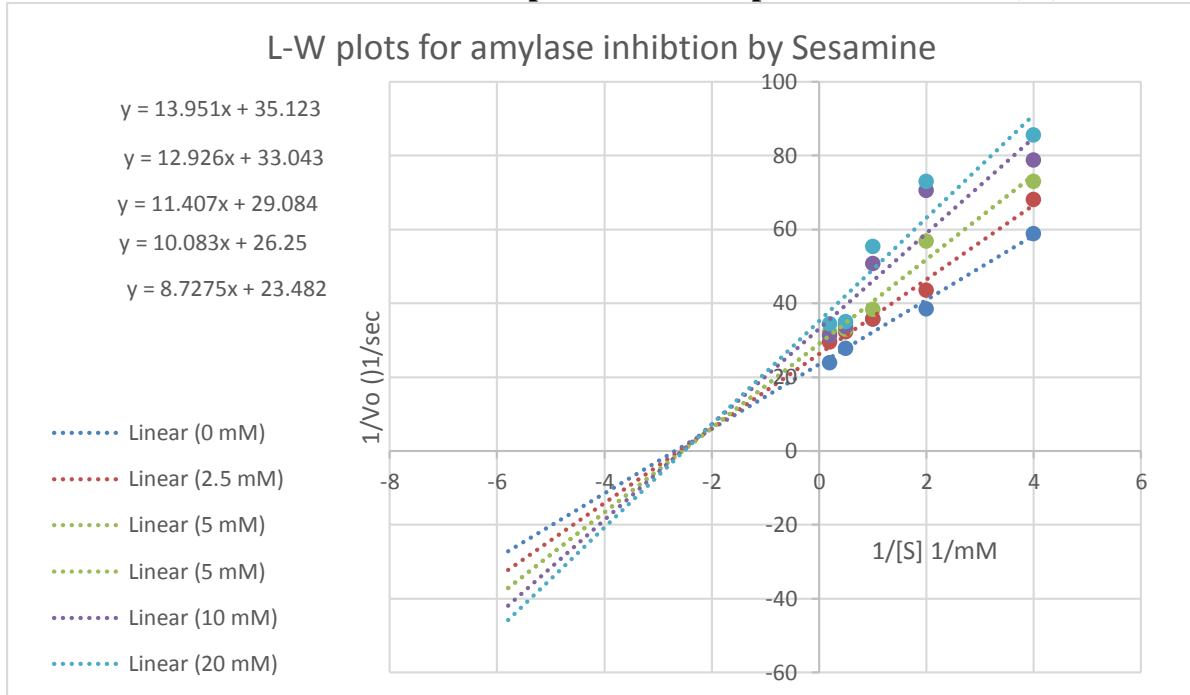
### Dixon plots for amylase inhibition by Zanthoamide D



### Dixon plots for glucosidase inhibition by Zanthoamide D



## 12.5: Lineweaver-Burk plots and Dixon plots for Sesamine (46)





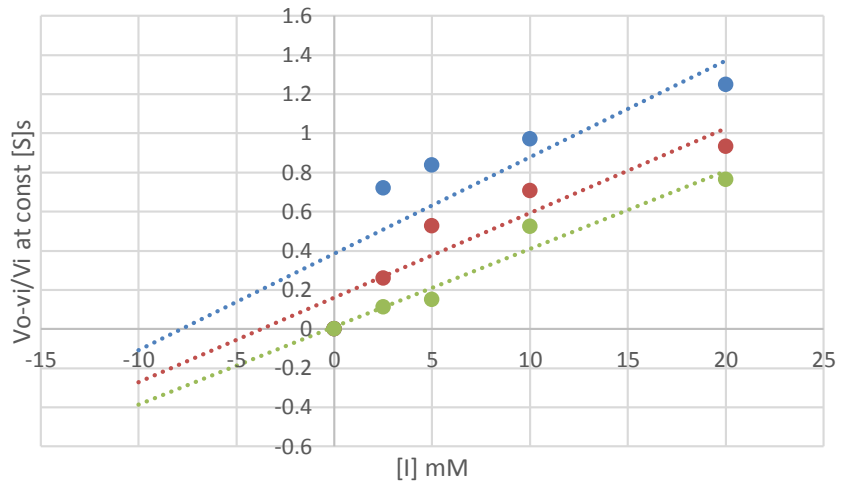
### Dixon plots for amylase inhibition by Sesamine

$y = 0.0494x + 0.3846$   
 $R^2 = 0.7029$

$y = 0.0432x + 0.1609$   
 $R^2 = 0.8707$

$y = 0.0397x + 0.0115$   
 $R^2 = 0.9552$

- ..... Linear (1 mM)
- ..... Linear (0.5 mM)
- ..... Linear (0.25 mM)



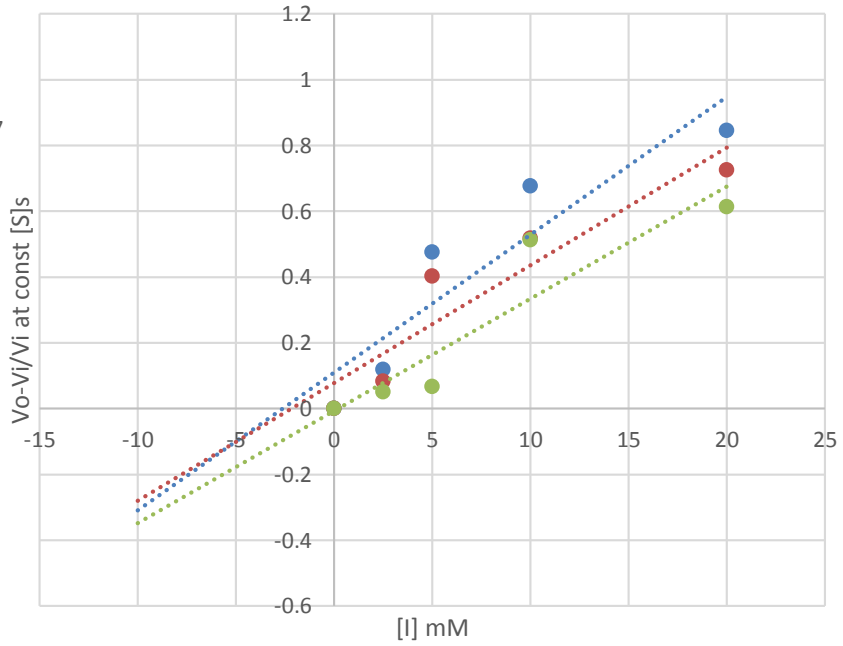
### Dixon plots for glucosidase by sesamine

$y = 0.0419x + 0.109$   
 $R^2 = 0.8483$

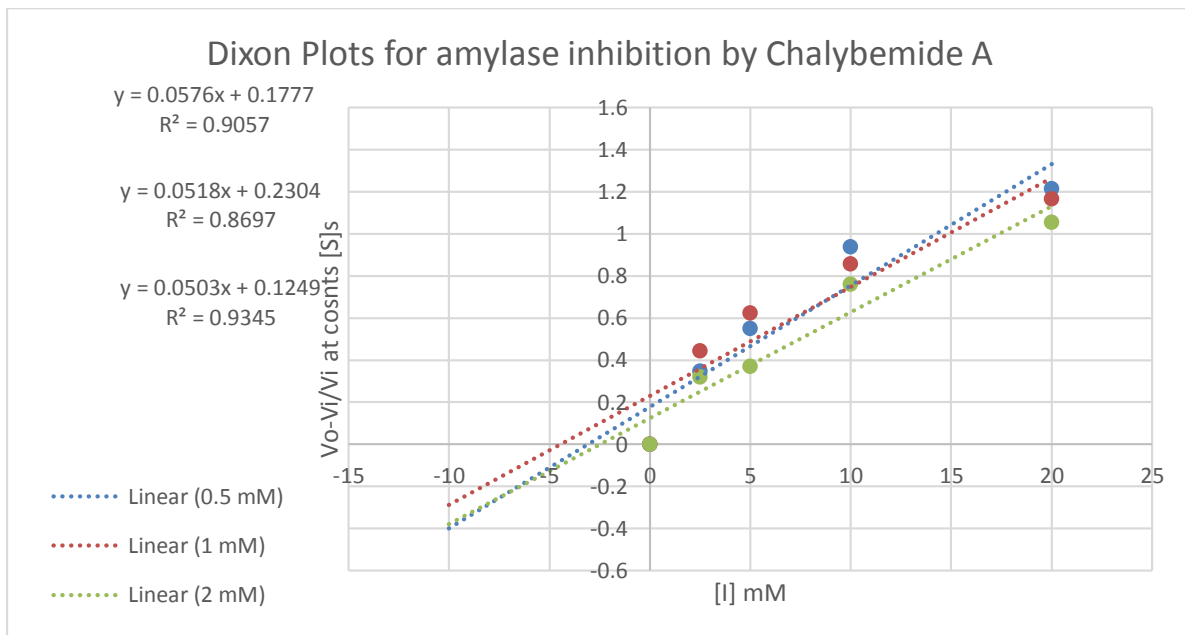
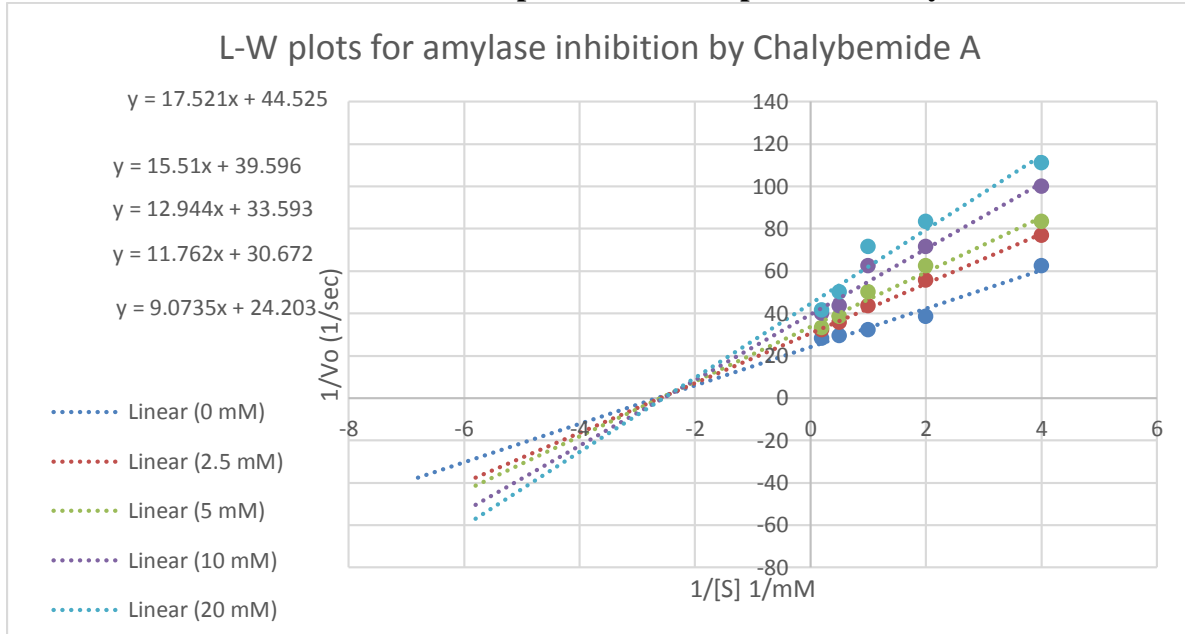
$y = 0.0358x + 0.0777$   
 $R^2 = 0.8748$

$y = 0.0341x - 0.0069$   
 $R^2 = 0.8644$

- ..... Linear (0.5 mM)
- ..... Linear (1 mM)
- ..... Linear (2 mM)



## 12.6: Lineweaver-Burk plots and Dixon plots for Chalybemide A



### L-W plots for glucosidase inhibition by chalybemide A

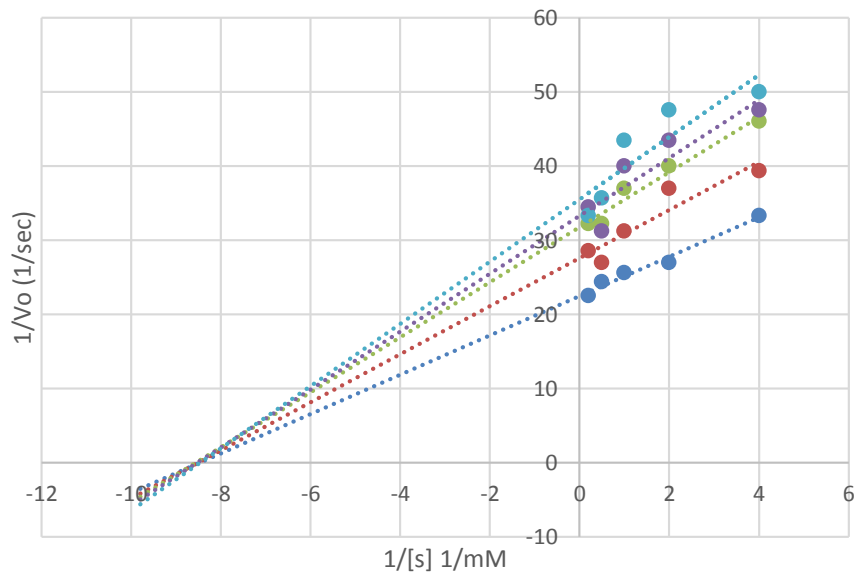
$$y = 3.9054x + 33.352$$

$$y = 3.7097x + 31.814$$

$$y = 3.2462x + 27.652$$

$$y = 2.6545x + 22.498$$

- ..... Linear (0 mM)
- ..... Linear (2.5 mM)
- ..... Linear (5 mM)
- ..... Linear (10 mM)
- ..... Linear (20 mM)



### Dixon Plots for glucosidases inhibitions by chalybemide A

$$y = 0.0499x + 0.0704$$

$$R^2 = 0.9824$$

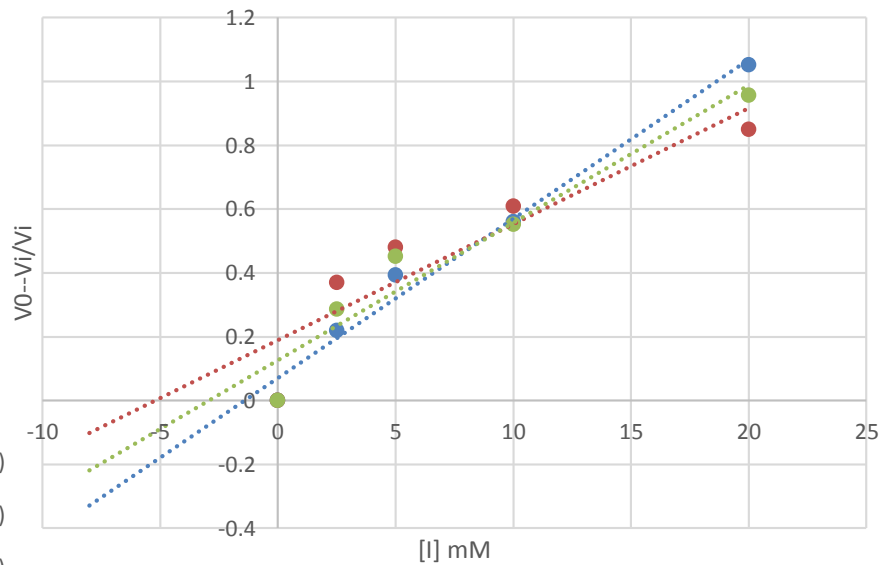
$$y = 0.0364x + 0.1888$$

$$R^2 = 0.8394$$

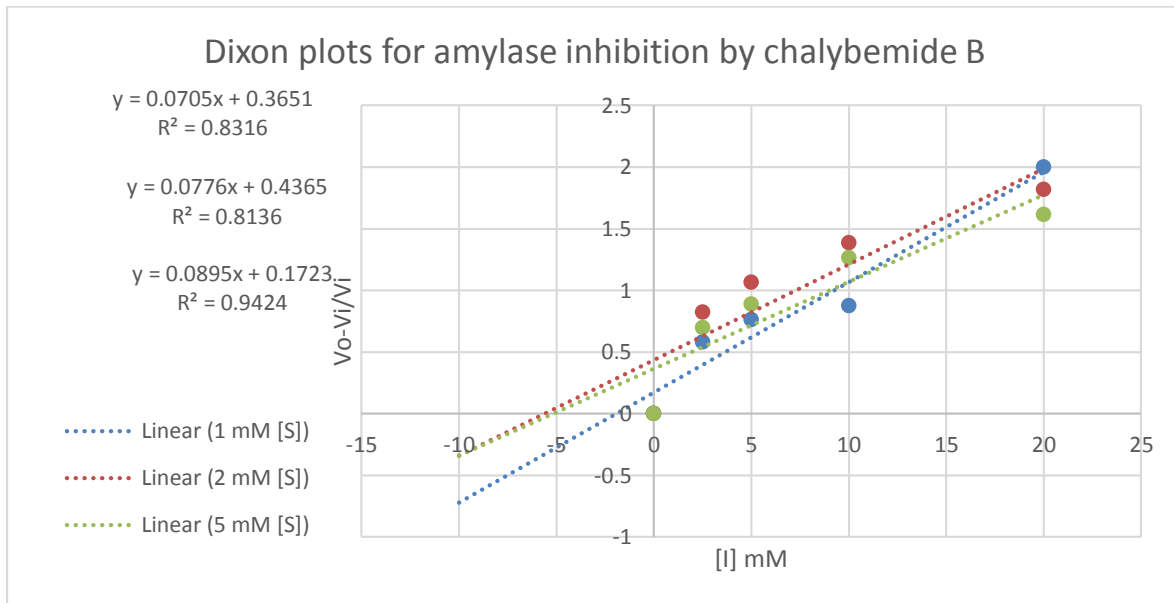
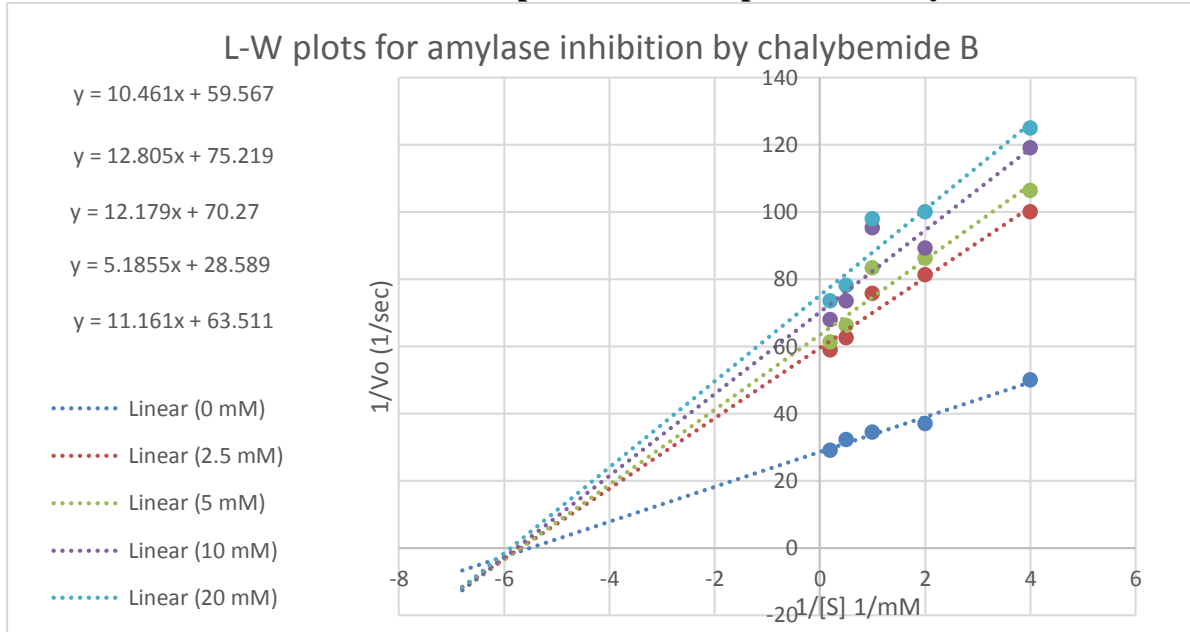
$$y = 0.0431x + 0.1258$$

$$R^2 = 0.9361$$

- ..... Linear (1 mM)
- ..... Linear (2 mM)
- ..... Linear (5 mM)



### 12.7: Lineweaver-Burk plots and Dixon plots for Chalybemide B



### L-W plots for glucosidase inhibition by Chaylbemide B

$$y = 4.8434x + 30.094$$

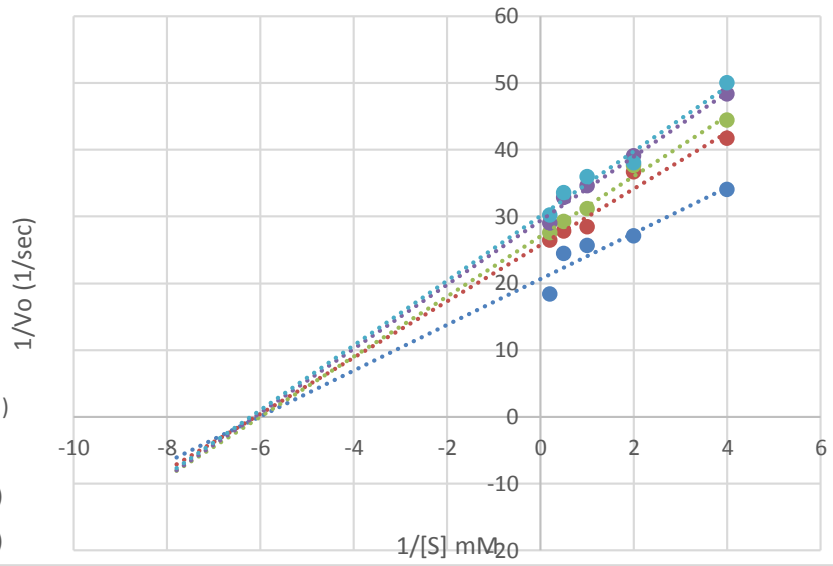
$$y = 4.7906x + 29.372$$

$$y = 4.5038x + 27.06$$

$$y = 4.2059x + 25.711$$

$$y = 3.4201x + 20.626$$

- ..... Linear (0 mM [I])
- ..... Linear (2.5 mM [I])
- ..... Linear (5 mM [I])
- ..... Linear (10 mM [I])
- ..... Linear (20 mM [I])



### Dixon plots for glucosidase inhibition by chalybemide B

$$y = 0.0211x + 0.0743$$

$$R^2 = 0.9184$$

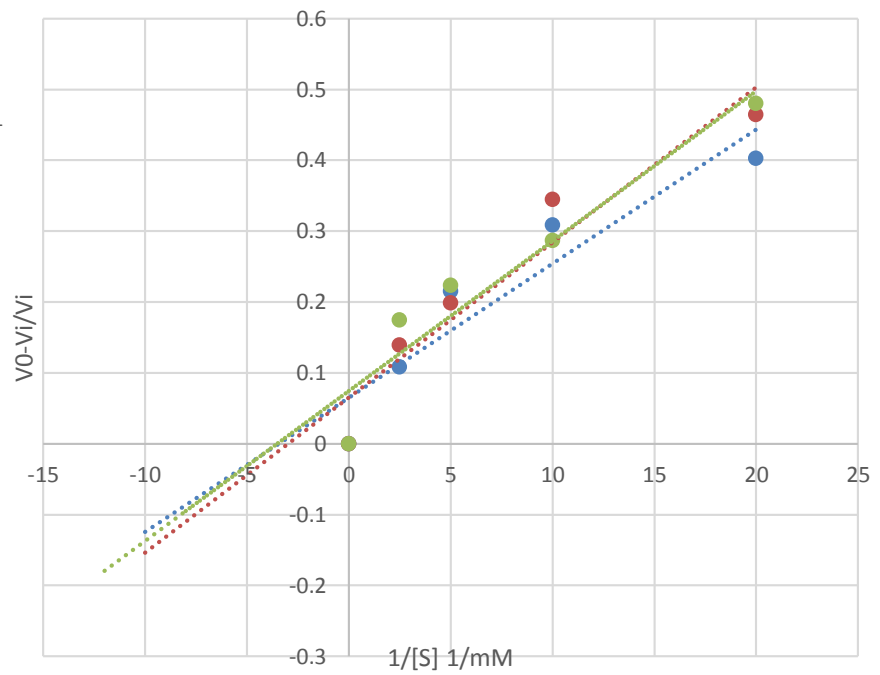
$$y = 0.0219x + 0.0651$$

$$R^2 = 0.9207$$

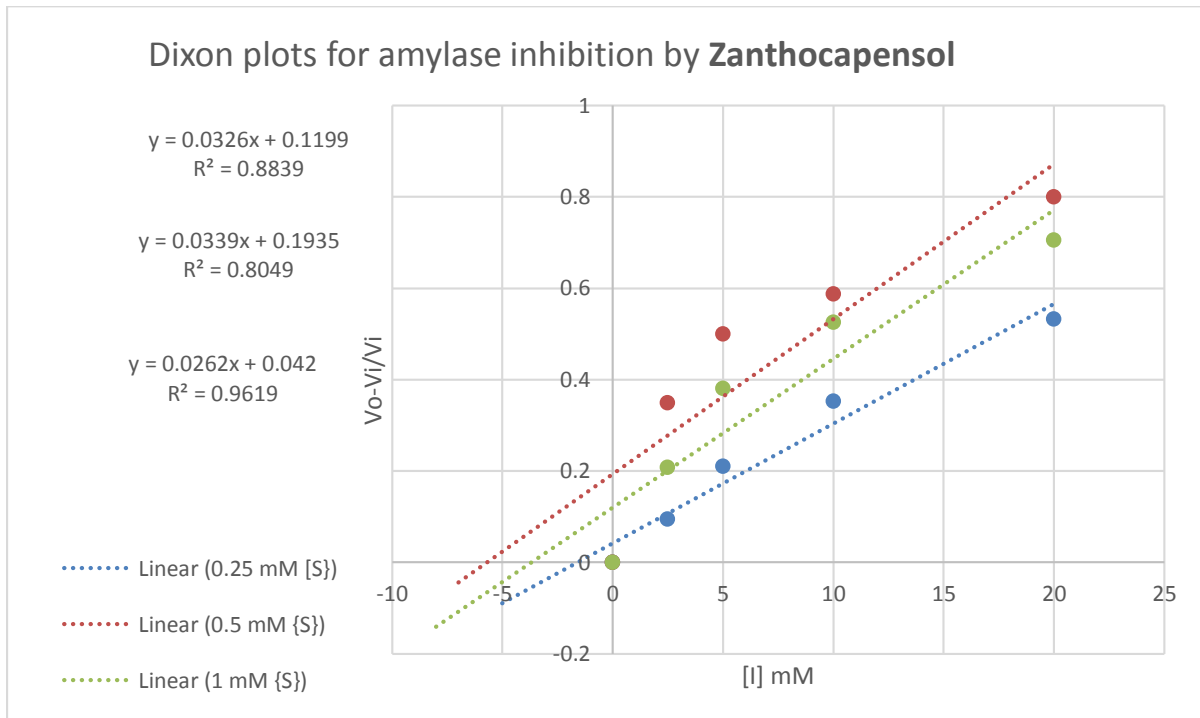
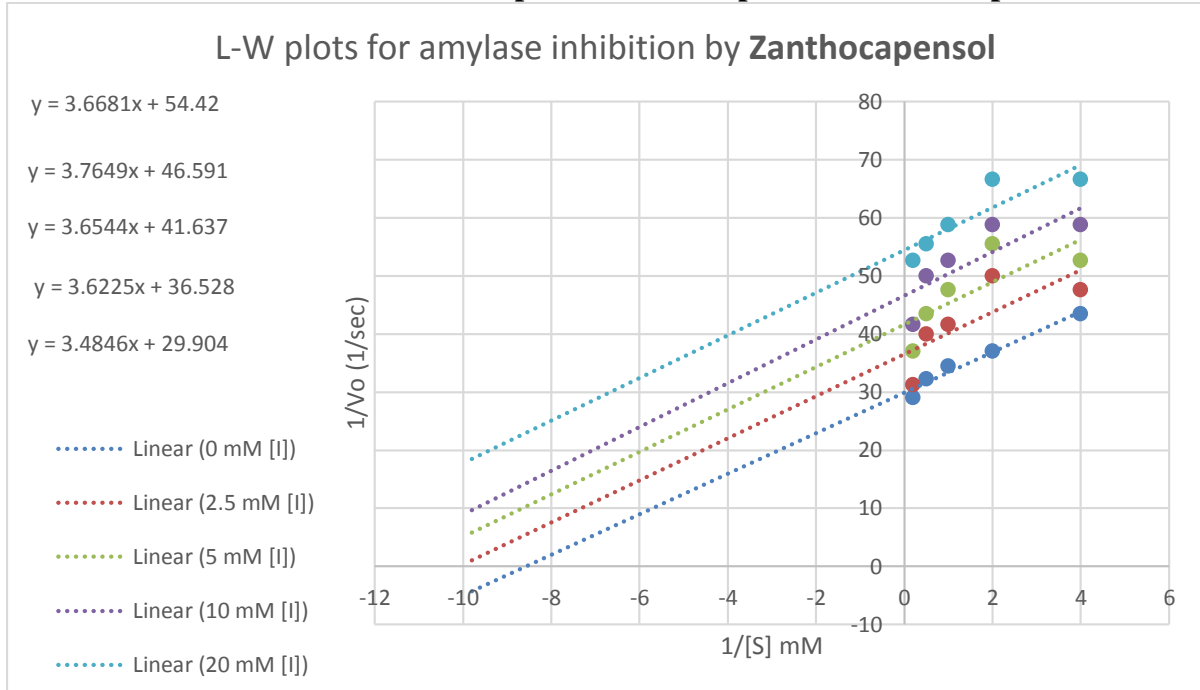
$$y = 0.0189x + 0.065$$

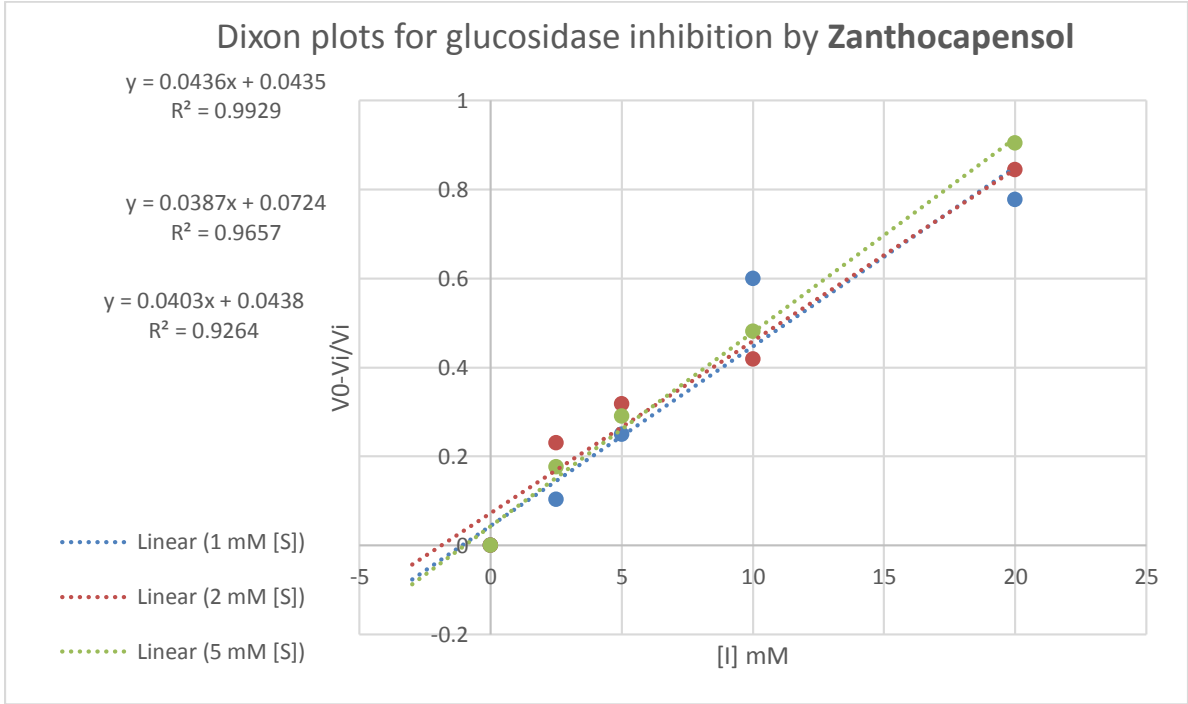
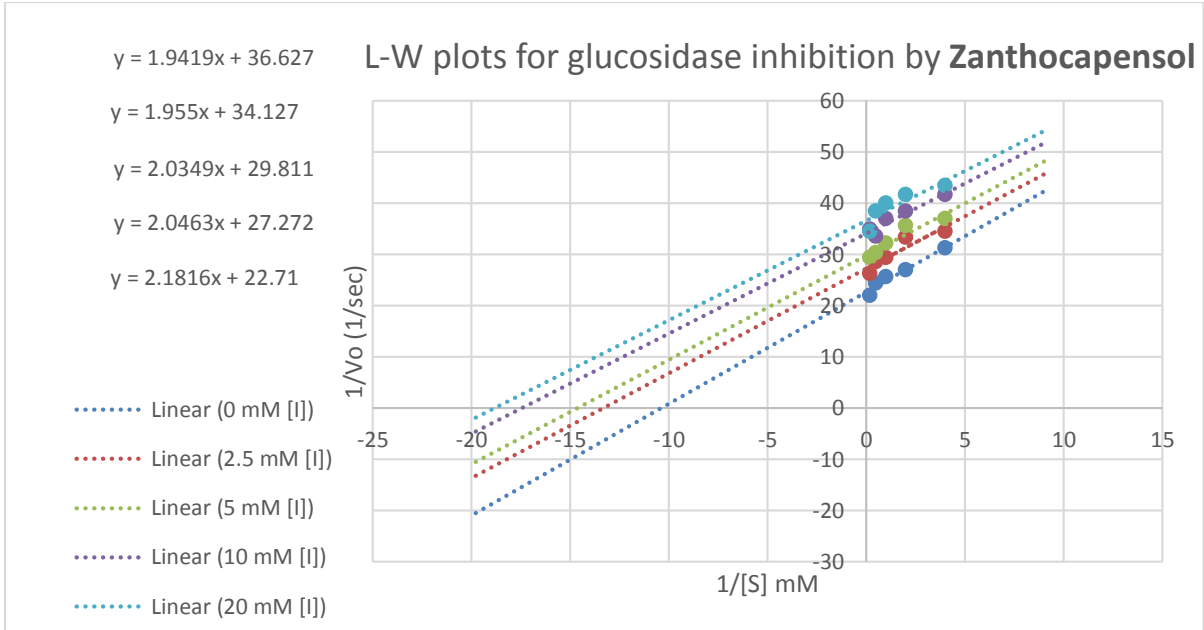
$$R^2 = 0.8825$$

- ..... Linear (1 mM [S])
- ..... Linear (2 mM [S])
- ..... Linear (5 mM [S])

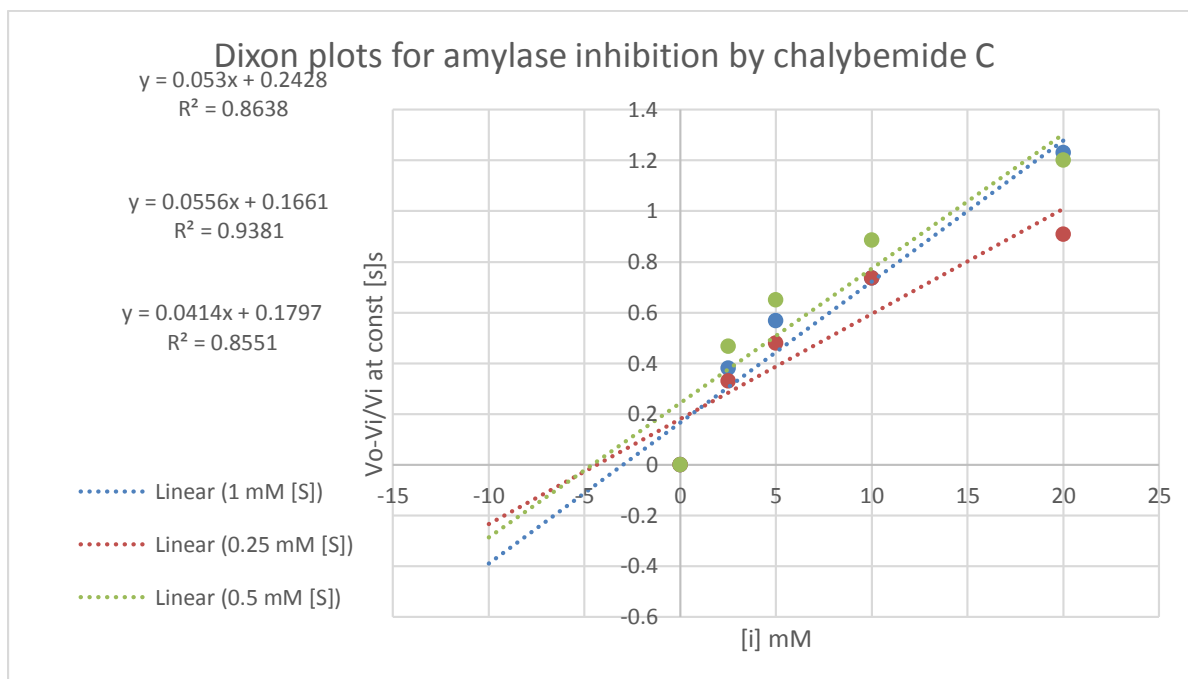
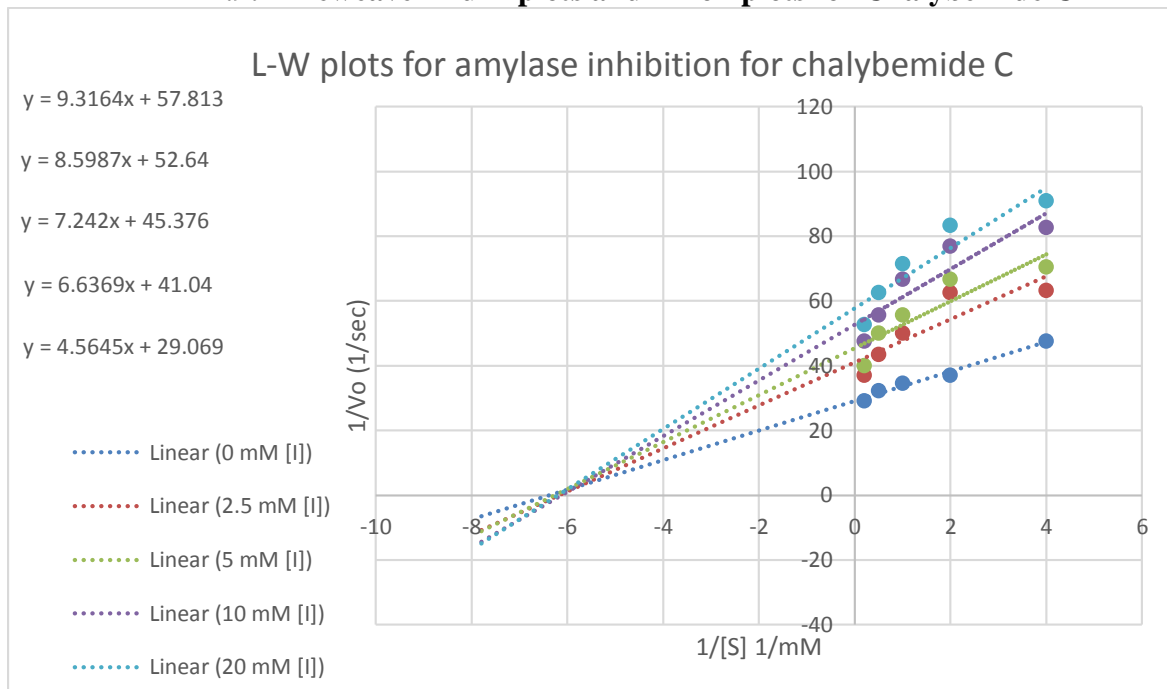


## 12.8: Lineweaver-Burk plots and Dixon plots for Zanthocapsol





## 12.9: Lineweaver-Burk plots and Dixon plots for Chalybemide C





### L-W plots for glucosidase inhibition by chalybemide C

$$y = 5.4495x + 37.366$$

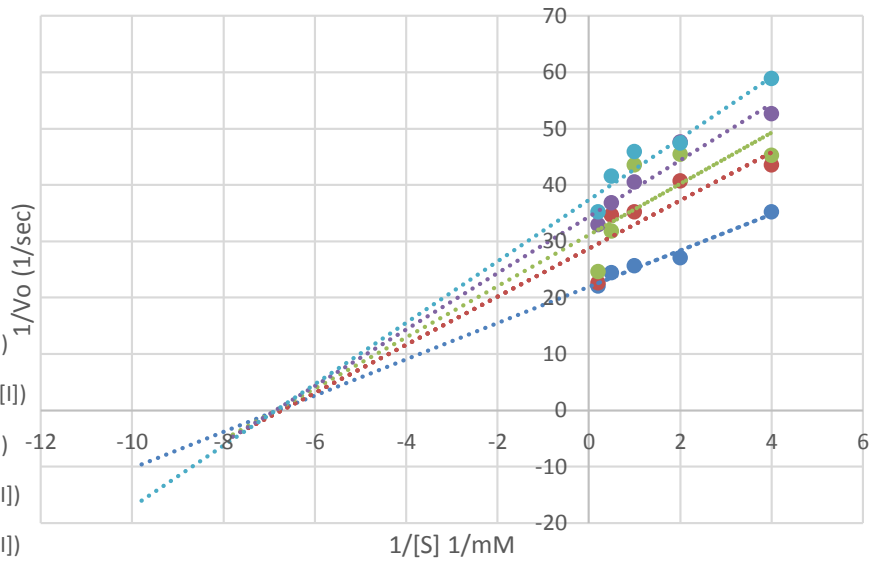
$$y = 5.0044x + 34.372$$

$$y = 4.5381x + 31.111$$

$$y = 4.2768x + 28.727$$

$$y = 3.2148x + 21.911$$

- ..... Linear (0 mM [I])
- ..... Linear (2.5 mM [I])
- ..... Linear (5 mM [I])
- ..... Linear (10 mM [I])
- ..... Linear (20 mM [I])



### Dixon Plots for glucosidases inhibitions by chalybemide C

$$y = 0.0499x + 0.0704$$

$$R^2 = 0.9824$$

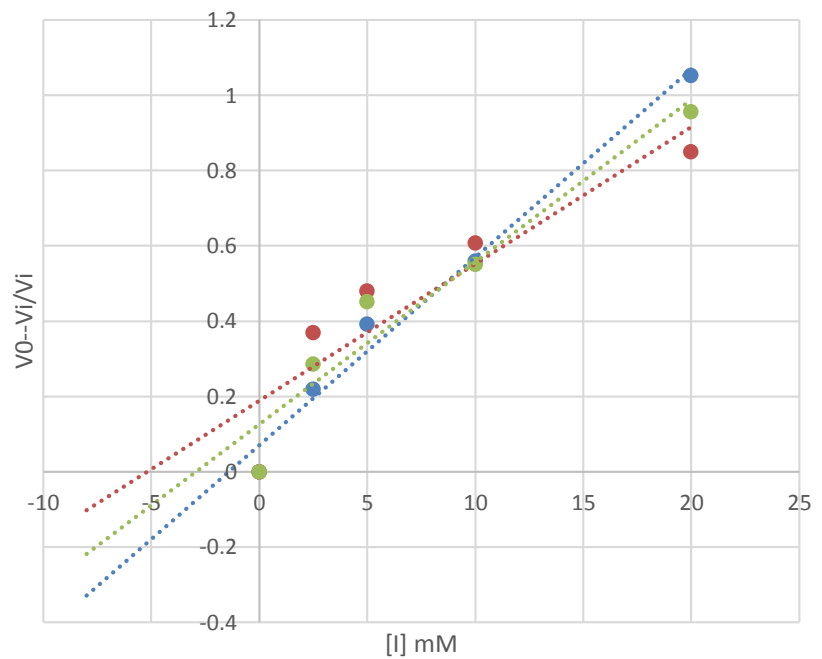
$$y = 0.0364x + 0.1888$$

$$R^2 = 0.8394$$

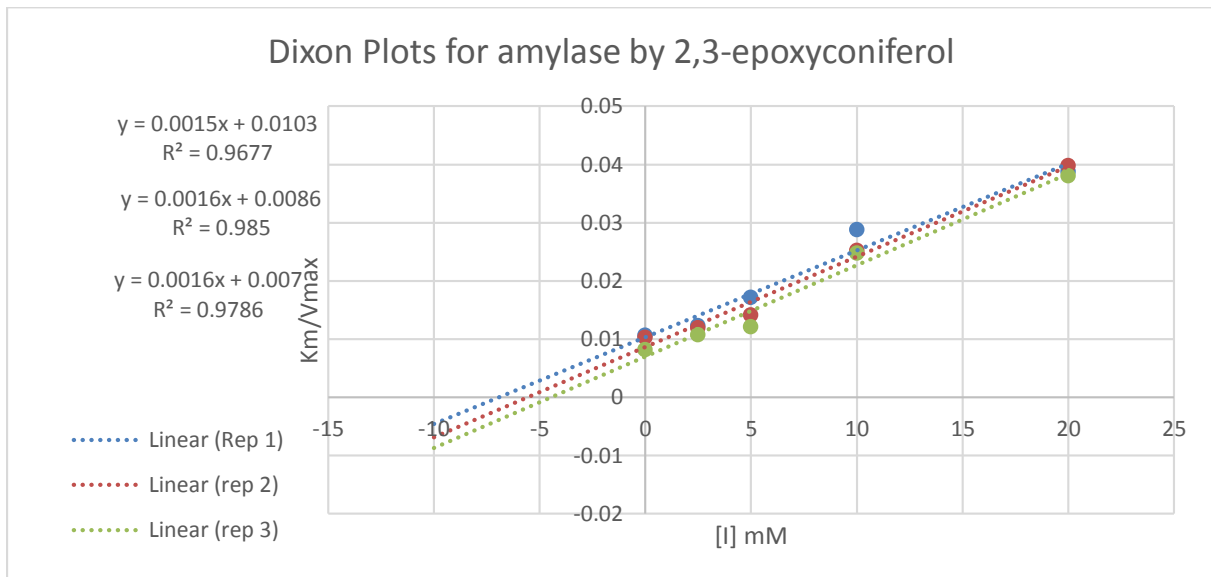
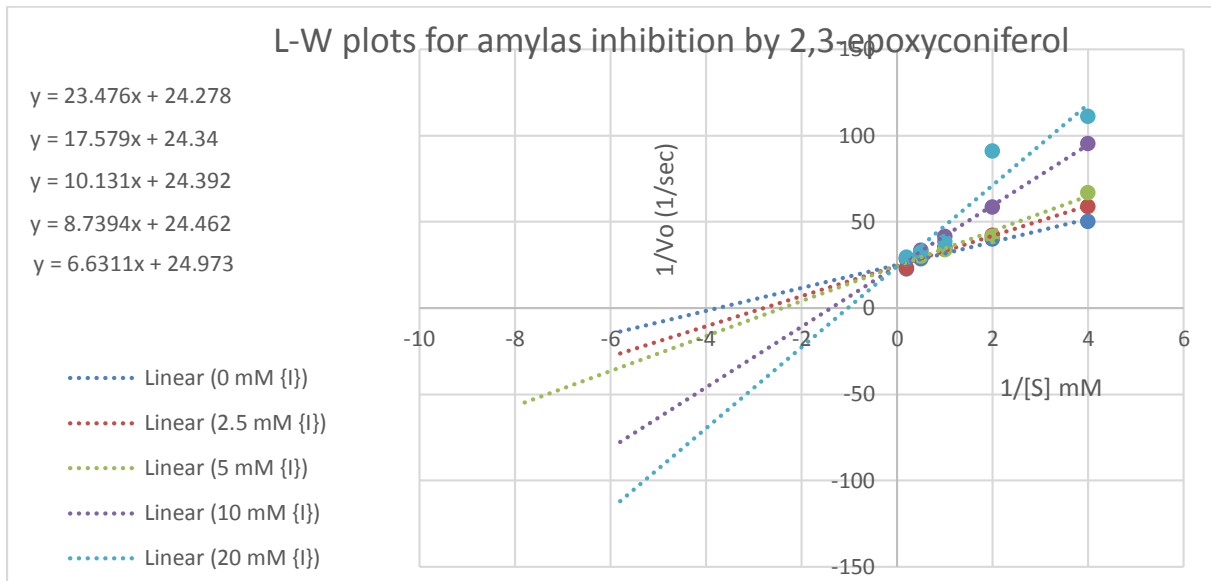
$$y = 0.0431x + 0.1258$$

$$R^2 = 0.9361$$

- ..... Linear ([S] at 0.25 mM)
- ..... Linear ([S] at 0.5 mM)
- ..... Linear ([S] at 1 mM)



### 12.10: Lineweaver-Burk plots and Dixon plots for 2,3-epoxy-6,7-methlenedioxy coniferyl alcohol



### L-W plots for glucosidase inhibition by 2,3-epoxyconiferol

$$y = 15.004x + 20.107$$

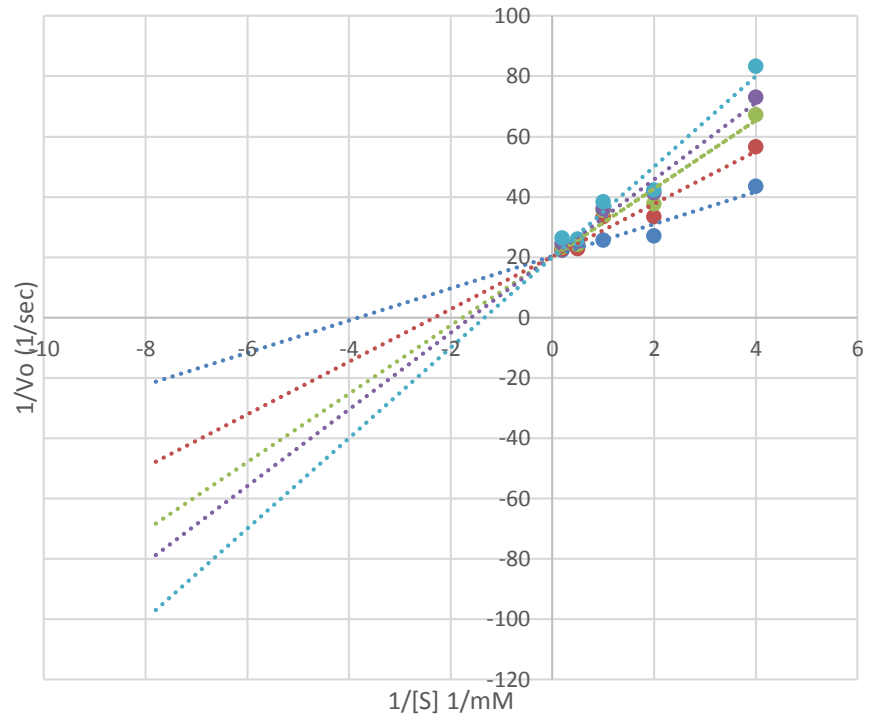
$$y = 12.726x + 20.424$$

$$y = 11.333x + 20.088$$

$$y = 8.7305x + 20.291$$

$$y = 5.3431x + 20.326$$

- ..... Linear (0 mM [I])
- ..... Linear (2.5 mM [I])
- ..... Linear (5 mM [I])
- ..... Linear (10 mM [I])
- ..... Linear (20 mM [I])



### Dixon plots for glucosidase by 2,3-epoxyconiferol

$$y = 0.4326x + 7.371$$

$$R^2 = 0.8316$$

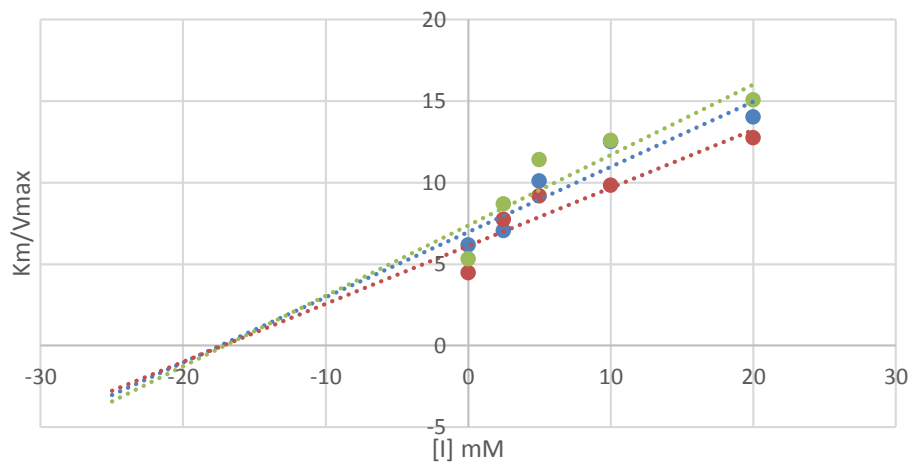
$$y = 0.4003x + 6.9613$$

$$R^2 = 0.8692$$

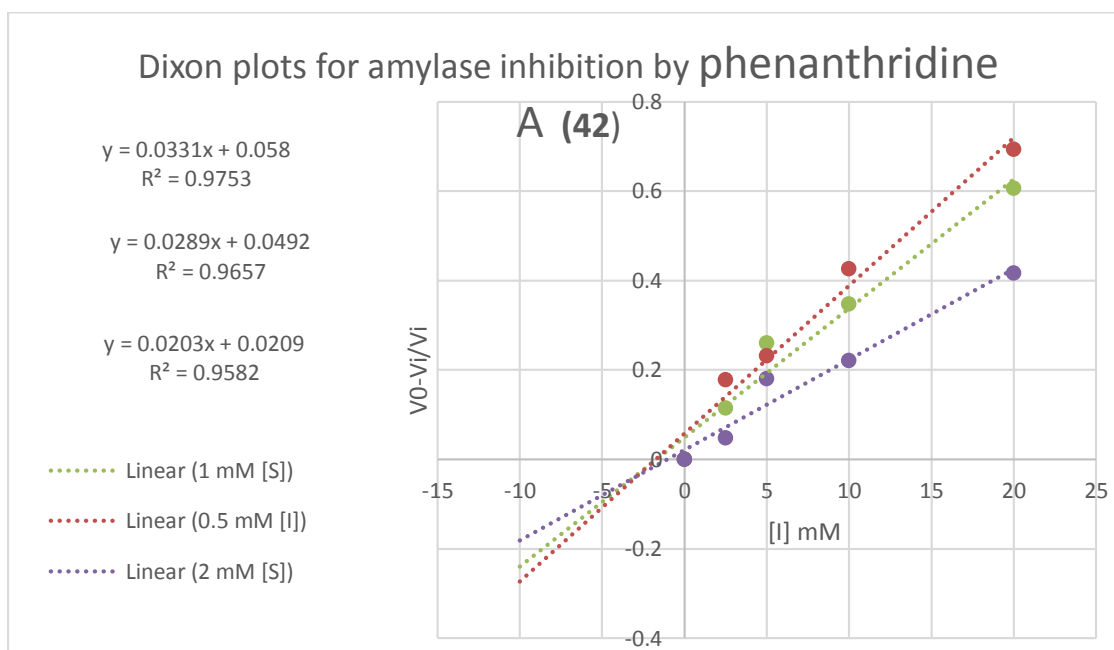
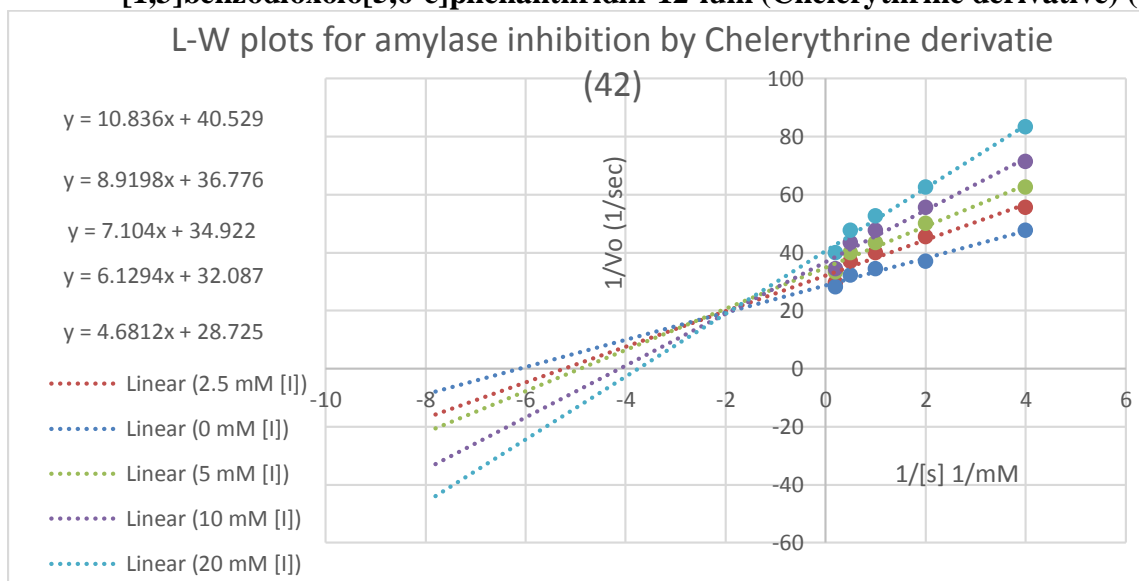
$$y = 0.3557x + 6.1124$$

$$R^2 = 0.8597$$

- ..... Linear (rep 1)
- ..... Linear (rep 2)
- ..... Linear (rep 3)



**12.11: Lineweaver-Burk plots and Dixon plots for 1,3-dimethoxy-12-methyl-9H-[1,3]benzodioxolo[5,6-c]phenanthridin-12-ium (Chelerythrine derivative) (42)**



### L-W plots for glucosidase inhibition by phenanthridine A (42)

$$y = 4.0214x + 29.821$$

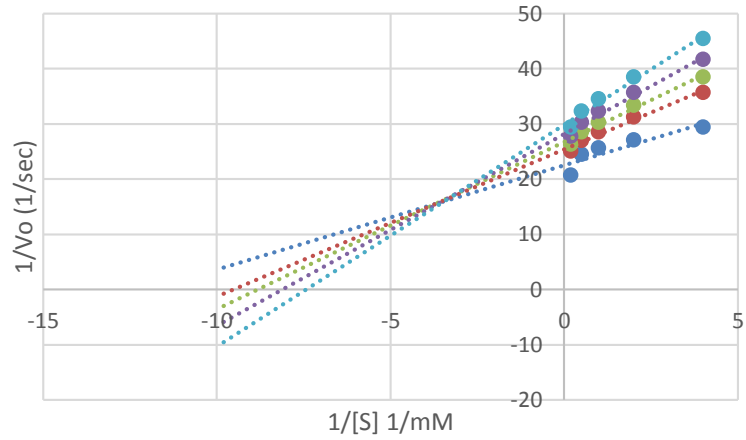
$$y = 3.4745x + 28.193$$

$$y = 3.033x + 26.726$$

$$y = 2.6713x + 25.399$$

$$y = 1.8902x + 22.529$$

- ..... Linear (0 mM [I])
- ..... Linear (2.5 mM [I])
- ..... Linear (5 mM [I])
- ..... Linear (10 mM [I])
- ..... Linear (20 mM [I])



### Dixon plots for glucosidase inhibition by phenanthridine A (42)

$$y = 0.0274x + 0.1281$$

$$R^2 = 0.8604$$

$$y = 0.0216x + 0.0733$$

$$R^2 = 0.9205$$

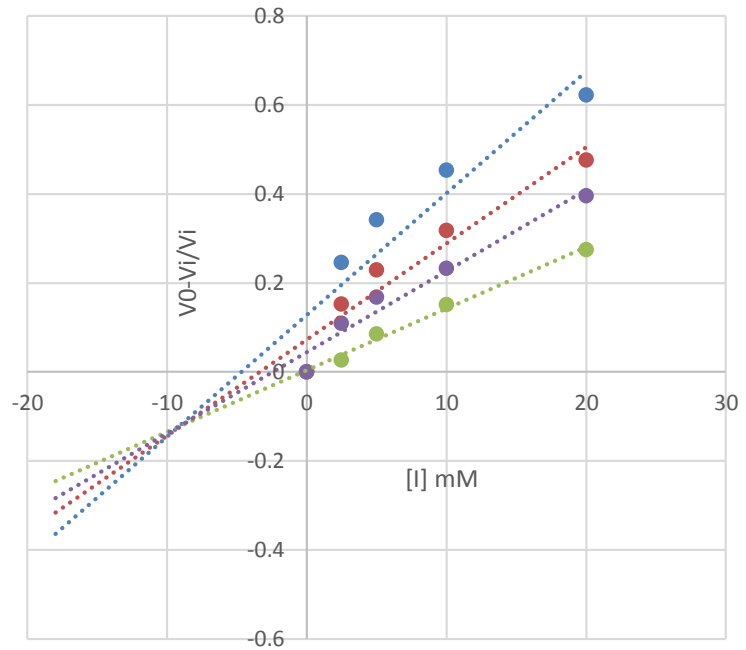
$$y = 0.0183x + 0.0446$$

$$R^2 = 0.958$$

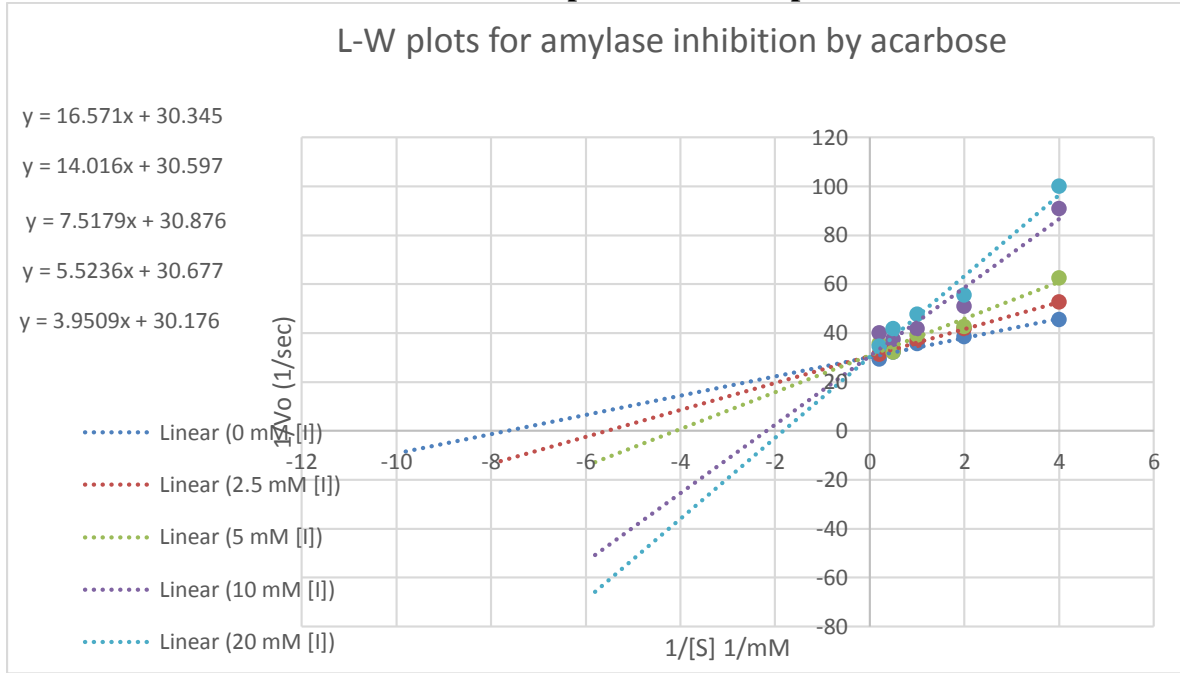
$$y = 0.0139x + 0.0038$$

$$R^2 = 0.9913$$

- ..... Linear (0.5 mM [S])
- ..... Linear (0.5 mM [S])
- ..... Linear (1 mM [S])
- ..... Linear (2 mM [S])



### 12.12: Lineweaver-Burk plots and Dixon plots for Acarbose



### L-W plots for glucosidase inhibition by acarbose

$$y = 12.954x + 23.396$$

$$y = 11.577x + 23.441$$

$$y = 11.426x + 23.423$$

$$y = 8.1322x + 23.077$$

$$y = 5.0929x + 23.246$$

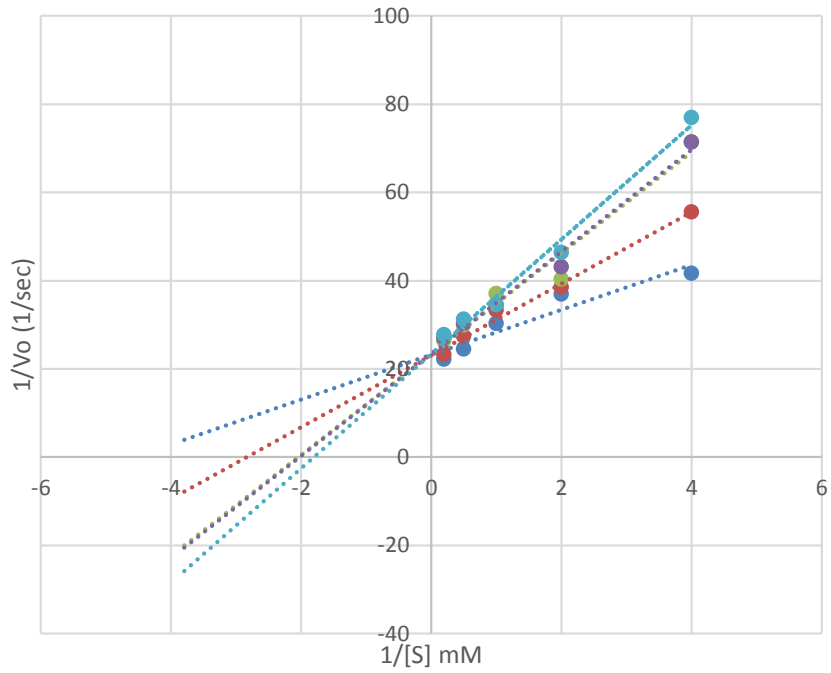
..... Linear (0 mM [I])

..... Linear (2.5 mM [I])

..... Linear (5 mM [I])

..... Linear (10 mM [I])

..... Linear (20 mM [I])



### Dixon Plots for glucosidases inhibitions by chalybemide A

$$y = 0.0499x + 0.0704$$

$$R^2 = 0.9824$$

$$y = 0.0364x + 0.1888$$

$$R^2 = 0.8394$$

$$y = 0.0431x + 0.1258$$

$$R^2 = 0.9361$$

..... Linear (1 mM)

..... Linear (2 mM)

..... Linear (5 mM)

