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**Potential of South African medicinal plants as treatments
for Alzheimer's disease through A β 42 protein reduction
and identification of their active compounds**

by

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Submitted in partial fulfilment of the requirements for the degree

PHILOSOPHIAE DOCTOR CHEMISTRY

In the Faculty of Natural & Agricultural Sciences

UNIVERSITY OF PRETORIA

PRETORIA

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December 2019

Submission declaration

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Acknowledgements

I would like to express my sincere gratitude to my supervisors Prof. V.J.Maharaj (main supervisor) and Dr N.October (co-supervisor) for their support, guidance, time, ideas and significant scientific contribution to this research project.

I would also like to thank Dr H.O.Yang and Dr Y.S.Chun, A.Park and Dr J.Kim of Korean Institute of Science and Technology (KIST) for the training provided me on A β 42 and bioassaying at KIST.

Special thanks to Dr. M.Selepe for her help with NMR and LC-MS SPE-NMR and providing me with the necessary training on the use of equipment. I would also like to thank M.Wooding for running samples on UPLC-QTOF-MS.

I am grateful for the help from J.Sampson the curator at University of Pretoria for the collection of plant material from University of Pretoria gardens. I also thank M.Nel for identifying plant species at plant herbarium at University of Pretoria and providing me voucher specimen numbers.

I express my gratitude for the support from bioprospecting group members (S. Muyisa, M.Sephora, B. Tembeni, B. Mzondo, N.K Khorommbi, L. Kruger, C. Ezeofor, F. Katele) in the department of chemistry at University of Pretoria.

This work would not have been successful without funding from University of Pretoria as well as Korean Institute of Science and Technology (KIST).

I am also thankful for emotional support, love and encouragement from my husband P. Thakur, my little boy Yohan and my parents as well. Above all I want to thank God for the opportunity to pursue my PhD studies and the strength to bring this work to completion.

Summary

Alzheimer's disease is a slow and progressive neurodegenerative disease that destroys memory and other important mental functions. APP (Amyloid precursor protein) is an integral membrane protein expressed in higher levels in brain and its abnormal cleavage produces beta amyloid (A β 42). Accumulation of A β 42 in the brain produces amyloid plaques, the hallmark of the Alzheimer's disease.

Most of the research for the treatment of Alzheimer's disease on medicinal plants in South Africa, focuses on acetyl cholinesterase inhibition, however this does not play a significant role against amyloid plaques. The goal of this study is to identify, characterize and develop new natural ingredients from the South African medicinal plants for the treatment of Alzheimer's disease by targeting the reduction of A β 42 resulting in decreased levels of beta amyloid plaques.

A literature survey was done for the selection of plants by using the keywords memory loss, aging, epilepsy, depression, mental health, dementia, forgetfulness, Alzheimer's disease, African medicinal plants, acetylcholinesterase. The plants were prioritized by a scoring system following the key criteria- plant part use, strength of the traditional use in relative to memory loss, plant toxicity, plant availability, published information against acetylcholinesterase inhibition. A total of 21 plant species were identified for collection, extraction and screening.

Different plant parts (leaves, bark, stem, roots and fruits) of the selected plants were collected from the University of Pretoria gardens and extracted with DCM:MeOH (1:1) after drying and grinding. The 34 plant extracts were screened to determine their inhibitory properties for A β 42 production in HeLa cells stably transfected with APP^{sw}. We found that of 34 extracts, 11 extracts (32.3%) were found significantly decrease the A β 42 level. Amongst them, the extracts of the leaves of *Xysmalobium undulatum*, leaves of *Cussonia paniculata*, leaves of *Schotia brachypetala* and stems of *Heteromorpha arborescens* potently reduced the secreted level of A β 42 by $77.3 \pm 0.5\%$, $57.5 \pm 1.3\%$, $44.8 \pm 0.1\%$ and $68.6 \pm 0.2\%$, respectively at 50 $\mu\text{g/ml}$.

These extracts were selected for further studies and UPLC- QTOF-MS was used to identify the active ingredients in the extracts. A total of five compounds were tentatively identified including four flavonoids, isoquercetin (**25**), myricetin-3-O-alpha-L-rhamnopyranoside (**26**), quercetin-3-O-rhamnoside (**27**), quercetin (**28**) from leaf extract of *S. brachypetala*. The quercetin derivatives were previously studied for their potential to treat Alzheimer's disease through A β reduction in different bioassays, in all likelihood the compounds would be responsible for the A β 42 reduction in *S. brachypetala*.

Using ESI negative mode of UPLC-QTOF-MS, a total of nine compounds were tentatively identified as quinic acid (**29**), 3,5-dicaffeoylquinic acid (**30**), rutin (**31**), valeriananoid E (**32**), acuminoside (**33**), dictamnoid D (**34**), 2''-O- β -D-glucopyranosylsaikosaponin B2 (**35**), clinoposaponin C (**36**), spinasaponin C (**37**) from the leaf extract of *C. paniculata*. Rutin is one of the most researched compound for the Alzheimer's disease in previous studies. 3,5-dicaffeoylquinic acid was previously reported for AChE inhibition. Rutin or 3,5-dicaffeoylquinic acid are in all likelihood the active ingredients in *C. paniculata* leaf extract.

Using ESI negative mode, a total of four compounds were tentatively identified named cynarin (1,5-dicaffeoylquinic acid) (**38**) alternoside IX (**39**), alternoside I (**40**) and saikogenin B₄ (**41**) the stem extract of *H. arborescens*. Cynarin was previously reported for the reduction of A β cytotoxicity while the other identified compounds, triterpenoid saponins were not examined for Alzheimer's disease, but found to exhibit neuroprotective properties. These compounds may in all likelihood contribute as the active ingredients from *H. arborescens* leaf extract for the reduction of A β 42.

The DCM:MeOH leaf extract of *X. undulatum* was fractionated using silica gel column chromatography to produce 15 fractions which were screened for A β 42 production. Of these two fractions, were found to be active as they significantly reduced A β 42 levels. Five pure compounds were successfully isolated from active fractions using preparative HPLC. Using NMR and UPLC-QTOF-MS, the structure of these compounds were elucidated as acetylated glycosydated crotoxinigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxinigenin-3-O-

β -digitalopyranosyl-(1-4)-*O*- β -digitoxopyranoside (**58**), crotoxinigenin 3-*O*-glucopyranoside (**59**) and desglucouzarin (**60**) isolated from the leaves of *X. undulatum*. The present study reports for the first time, the evaluation of the compounds for their potential to reduce A β 42 levels. The four cardenolide glycosides, acetylated glycosylated crotoxinigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxinigenin-3-*O*- β -digitalopyranosyl-(1-4)-*O*- β -digitoxopyranoside (**58**), desglucouzarin (**60**) were found to significantly decreased the A β 42 levels at 10 μ M by $75.04 \pm 0.85\%$, $74.66 \pm 0.1\%$, $65.55 \pm 0.19\%$ and $69.15 \pm 0.25\%$, respectively as compared to the negative control. These results revealed that all the pure compounds including the inactive compound (crotoxinigenin 3-*O*-glucopyranoside (**59**)) isolated from *X. undulatum* leaf extract were cardiac glycosides and contain the characteristic lactone ring which is known to be responsible for the cardiotoxicity. Since this was present in both the inactive and active cardenolides in all probability the lactone ring is the active site of the molecules contributing towards activity as such any analogues synthetically prepared lacking the lactone ring may still retain A β 42 reduction while also reducing cardiotoxicity. Further the results of dose response studies of the 4 cardenolides indicated that crotoxinigenin-3-*O*- β -digitalopyranosyl-(1-4)-*O*- β -digitoxopyranoside (**58**) was the most potent compound as it significantly reduced A β 42 levels at the low concentration of 0.025 μ M. These results revealed that this is the most active compound responsible for A β 42 reduction in the plant. The bioassay-guided approach used for identifying the active ingredient in *X. undulatum* responsible for the reduction of A β 42 resulted in identification of 4 cardenolides active ingredient, with one being the most active at nanomolar levels, which could be developed further into pharmaceutical drugs.

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Abbreviations

^1H NMR	Proton NMR
^{13}C NMR	Carbon-13 nuclear magnetic resonance
5-HT	5- hydroxytryptamine
6-OHDA	6-hydroxydopamine
ABTS	2,2'-azino-bis-3-ethylbenzotiazolin-6-sulfonic acid
AChE	Acetylcholinesterase
AChEI	Acetylcholinesterase inhibition
AD	Alzheimer's disease
ANOVA	Analysis of variance
APOE	Apolipoprotein E
APP	Amyloid precursor protein
APP ^{sw}	APP carrying Swedish mutation
A β	Amyloid beta
<i>B. natalensis</i>	<i>Bulbine natalensis</i>
<i>B. salvifolia</i>	<i>Buddleja salvifolia</i>
BPI	Base peak ion
BuChE	Butyrylcholinesterase
<i>C. africana</i>	<i>Comelina Africana</i>
<i>C. asiatica</i>	<i>Centella asiatica</i>
<i>C. edulis</i>	<i>Catha edulis</i>
<i>C. orbiculata</i>	<i>Cotyledon orbiculata</i>
<i>C. paniculata</i>	<i>Cussonia paniculate</i>
<i>C. spicata</i>	<i>Cussonia spicata</i>
ChEBI	Chemical Entities of Biological Interest
CNS	Central nervous system
COSY	Correlation spectroscopy
COX	Cyclooxygenase
DCM	Dichloromethane
DEPT	Distortionless enhancement by polarisation transfer
DMEM	Dulbecco's Modified Eagle Medium
DMSO	Dimethyl sulphoxide
DPPH	2,2-diphenyl-1-picrylhydrazyl
EC ₅₀	Effective concentration 50
EDTA	Ethylenediaminetetraacetic acid
ELISA	Enzyme-linked immunosorbent assay
ESI	Electron spray ionisation
FDA	Food and drug administration
FTIR	Fourier-transform infrared spectroscopy
GLIDE	Grid based Ligand Docking with Energetics
<i>H. arborescens</i>	<i>Heteromorpha arborescens</i>
<i>H. trifoliata</i>	<i>Heteromorpha trifoliata</i>
HEK	Human embryonic kidney
Hep2	Human epithelial type 2
HMBC	Heteronuclear multiple bond correlation
HSQC	Heteronuclear single quantum correlation
IC ₅₀	Inhibitory concentration 50
iFit	Isotopic fit value

KIST	Korean Institute of Science and Technology
LPS	Lipopolysaccharides
<i>M. whitei</i>	<i>Mondia whitei</i>
<i>m/z</i>	Mass to charge ratio
MAO	Monoamine oxidase
MAP	Microtubule-associated protein
MeOD	Deuterated methanol
MeOH	Methanol
min	Minute
ml	Millilitre
MS	Mass spectrophotometry
MTT	3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide
ng	Nanogram
NMDA	N-methyl-D-aspartate receptor
NMDA	<i>N</i> -methyl-D-aspartate
NMR	Nuclear magnetic resonance
Nrf2	Nuclear factor erythroid 2-related factor 2
OGD/R	oxygen-glucose deprivation followed by reoxygenation
<i>P. auriculata</i>	<i>Plumbago auriculata</i>
PBS	Phosphate-buffered saline
PDE-4	phosphodiesterase-4
PHF	Paired-helical filament
Prep HPLC-MS	Preparatory High-performance- liquid- chromatography
<i>R. graveolens</i>	<i>Ruta graveolens</i>
<i>S. brachypetala</i>	<i>Schotia brachypetala</i>
SANBI	South African National Biodiversity Institute
sAPP α	soluble alpha-amyloid precursor protein
sAPP β	Soluble beta-amyloid precursor protein
SASH	South Africa Stress and Health
SEM	Scanning electron microscope
SERT	Serotonin transporter
SSRI	Serotonin-specific re-uptake inhibitor
<i>T. dregeana</i>	<i>Trichilia dregeana</i>
<i>T. riparia</i>	<i>Tetradenia riparia</i>
TLC	Thin layer chromatography
TMT	Treadmill Stress Test
TPA	Tetradecanoylphorbol
UPLC-QTOF-MS	Ultra-high performance liquid chromatography-quadrupole time-of-flight mass spectrometry
UPLC-QTOF-MS	The ultra-high performance liquid chromatography-quadrupole time-of-flight mass spectrometry
VCR	Vincristine
VLB	Vinblastine
WHO	World Health Organization
<i>X. undulatum</i>	<i>Xysmalobium undulatum</i>
<i>Z. capense</i>	<i>Zanthoxylum capense</i>
<i>Z. mucronata</i>	<i>Ziziphus mucronata</i>

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Chapter 1

General introduction

1.1. Natural products and their value in drug discovery

Natural products have historically demonstrated their value as a powerful source of bioactive compounds with therapeutic potential, and nowadays still represent an important tool for the identification of novel drug leads. This field of study forms a vital part of drug discovery as there is a consistent need to identify and develop novel pharmacophores to fight diseases that include Alzheimer's, cancer, HIV-AIDS and many infectious diseases. In the developing countries, medicinal plants are continued to be used as the primary source of medicine by about 80% of the population for health care, due to their easy accessibility and low costing. The analysis of secondary metabolites derived from these medicinal plants contain several bioactive compounds which may potentially lead to the development of novel drugs for the pharmaceuticals [1]. In addition, particularly in developing countries, there is a long history of the use of herbal medicine because of its cultural and spiritual point of views [2]. In the early 19th century, the isolation of active compounds from medicinal plants began with the isolation of morphine from opium. Furthermore, the isolation and identification of many earlier discovered potent drugs such as cocaine, codeine, digitoxin, and quinine from medicinal plants pathed the way for continuous interest in the field [3].

The rational use of traditional medicines was encouraged by the member states of the World Health Assembly (WHO, 1978), aiming for providing affordable and accessible health care to the global population. In order to facilitate this process, WHO in 1993 and 1998 has designed some guidelines for the assessment of the quality, safety and efficacy of herbal medicines [4]. An analysis based on the *Annual Reports of Medicinal Chemistry* from 1983 to 1994, indicated that 60-70% of anti-cancer and anti-infective approved drugs were shown to be of natural origin [5]. There are a number of plant derived compounds which include vinblastine, vincristine, topotecan and irinotecan, etoposide, camptothecin derivatives, and paclitaxel

(taxol®) have been used clinically as anticancer agents. While some other natural products including flavopiridol and combretastin A4 phosphate are considered as promising potent anticancer agents [6]. Over 60% of currently using anticancer agents are derived from natural sources including plant species, micro-organisms and marine organisms. Vinca alkaloids, vinblastine (VLB) and vincristine (VCR) are the first anticancer agents isolated from Madagascar periwinkle, *Catharanthus roseus* G. Don. (Apocynaceae). These agents are used in combination with other cancer chemotherapeutic drugs for the treatment of various types of cancers and also diabetes [6].

It was found historically that quinoline based antimalarials and artemisinin derivatives derived from the medicinal plants are most commonly used for the treatment of malaria and its symptoms [7]. Artemisinin isolated from an ancient Chinese herb named Qing Hao (*Artemisia annua*) is used as remedy in traditional Chinese medicine for the treatment of malaria. [8].

A variety of secondary metabolites derived from plant species including alkaloids, coumarins, flavonoids, lignans, phenolics, quinones, saponins, terpenes, xanthenes and some other natural products showed moderate to good anti- HIV activity. Currently, there is no plant-derived drug in clinical use for the treatment of HIV-AIDS. But three natural products, calanolide A, 3-hydroxymethyl-4-methyl DCK (PA-334B) and DSB (PA-457) have shown promising activities to develop into a potent drug for HIV-AIDS [9].

Salvia officinalis and *Melissa officinalis* plant extract showed efficacy towards the treatment of mild to moderate Alzheimer's. Both plant extracts exhibited CNS acetylcholine receptor activity [10].

Natural products and their derivatives contribute more than one-third of all FDA-approved new molecular entities (NMEs). It is also revealed that one-quarter of these NMEs are derived from plants [11].

1.2. Background and history of Alzheimer's disease (AD)

Alzheimer's disease, is a multi-factorial neurodegenerative disorder that progresses slowly and destroys human memory, important mental functions and interrupt the daily activities in the ageing population [12]. It is a heterogenous group of dementia, also known as senile dementia that share common clinical symptoms, involving progressive cognitive impairment [13]. Alzheimer's is one of the most common form of dementia and primarily affects the people after age 65. The disease was named after the German Doctor Alois Alzheimer who was the first person to describe Alzheimer's in a patient named Auguste D. The patient, 51 years old woman who had shown some symptoms of progressive cognitive impairment, reduced comprehension and memory, aphasia, disorientation, unpredictable behavior, paranoia, hallucinations, delusions and psychosocial incompetence [14].

1.2.1. Epidemiology, facts and figures and risk factors of Alzheimer's disease

Alzheimer's patients have been predicted to grow from 47 million to 130 million by [15]. Global prevalence of dementia, a Delphi consensus study, in 2001, estimated that 24.3 million people have dementia with 4.6 million new cases reported every year and is expected to be doubled every 20 years and will reach up to 81.1 million by 2040 [16].

1.2.2. Causes and risk factors of Alzheimer's disease

Like all types of dementia, Alzheimer's is caused by brain cell death. It is a neurodegenerative disease which means there is progressive brain cell death that happens over a course of time. The total brain size shrinks with Alzheimer's – the tissue has progressively fewer nerve cells and connections [17].

1.2.2.1. Structural abnormalities

Although cerebral atrophy is a typical manifestation of Alzheimer's disease, it does not distinguish normal aging from Alzheimer's accurately enough to be diagnostic; this applies to

neuroimaging as well as gross inspection at post-mortem. However, microscopic examination reveals the critical features of the disease – a cerebral cortex peppered with the deposits of amyloid plaques or senile plaques and neurofibrillary tangles [18]. Alzheimer's disease is characterised by two deposits of abnormalities, senile plaques and neurofibrillary tangles and neural cell death [13]. Amyloid plaques contain mainly amyloid beta ($A\beta$) peptide generated by the proteolysis of a large β -amyloid precursor protein (APP) [19]. Accumulation of amyloid β in the brain is the primary factor causing AD [20]. Neurofibrillary tangles consist of an abnormal form of hyperphosphorylated tau aggregates [21]. Tau is one of the major microtubule-associated protein (MAP) in the brain, abundant in neurons present in any part of the body and performs as a stabilizer of microtubules in neurons and other cells, and also plays an important role in cell differentiation and polarization [22]. These neurofibrillary tangles make neuron unable to properly maintain its cytoskeleton and also to perform numerous processes [18]. The microscopic examination reveals the comparison of a healthy brain and Alzheimer's brain and also reveals the presence of senile plaques and neurofibrillary tangles in Alzheimer's brain as shown in Figure 1.1.

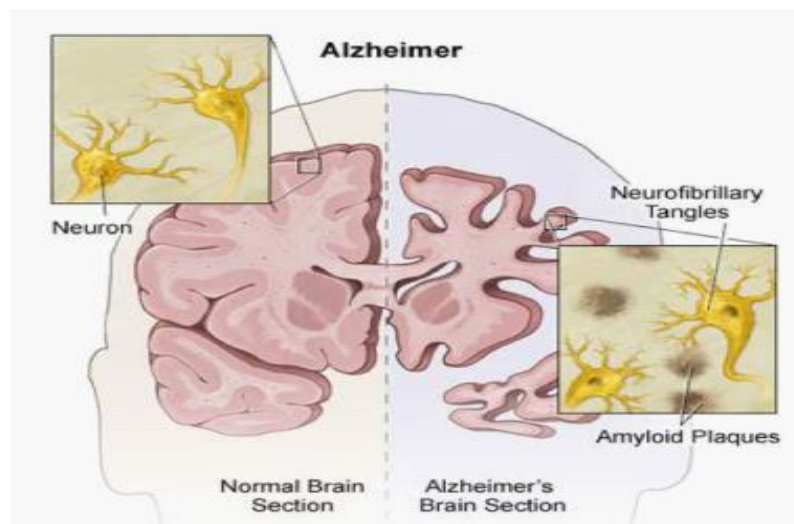


Figure 1.1: Comparison of normal brain and Alzheimer's brain¹

¹ The picture of normal brain and Alzheimer's brain was adapted from an online resource:
https://www.hopkinsmedicine.org/healthlibrary/conditions/adult/nervous_system_disorders/alzheimers_disease_85.p00772

1.2.2.2. The role of the genes

Genetic studies identified specific mutations in three different genes which are responsible for the early - onset of Alzheimer's disease, although these genetic mutations account for less than 1% of cases [23]. These gene mutations are APP (on chromosome 21), PSEN-1 (on chromosome 14), PSEN-2 (on chromosome 1). These are proposed to increase the production of less soluble and more toxic A β 42 [24]. The gene responsible for the late-onset Alzheimer's disease has been identified as *APOE* located on chromosome 19. *APOE* is involved in the process of synaptic repair and also plays an important role in the maintenance of neuronal structure and cholinergic functions [18]. *APOE* has three allelic variants (ϵ 2, ϵ 3 and ϵ 4) with *APOE*- ϵ 4 is the one which is generally associated with the late-onset Alzheimer's disease while *APOE*- ϵ 2/*APOE*- ϵ 3 appears to lower the Alzheimer's risk [25]. Everyone inherits one of these genes from each parent and the people who inherit one copy of *APOE*- ϵ 4 develop four-fold risk as compared with the people having two copies of *APOE*- ϵ 3 who have 12-fold risk of developing Alzheimer's. Also *APOE*- ϵ 4 is also responsible for developing Alzheimer's at a younger age than those with the other two allelic variants (ϵ 2, ϵ 3) of *APOE* gene [26].

1.2.2.3. Family history and Aging

Family history, age, environmental factors, cardiovascular risk factors and education are some other risk factors which can contribute to the Alzheimer's disease. 25% of people with age 55 or older suffering from dementia have a family history of dementia involving a first-degree relative. However a family history does not necessarily means that an individual will develop the disease [27]. Aging is one of the most important risk factor for Alzheimer's disease and thought to affect 4-8% of the population over 65 years [28].

1.2.3. ²Signs and Symptoms of Alzheimer's disease

The common symptoms include difficulty in retaining information and expressing thoughts. Patient may start forgetting recently learned information, important dates or events and relying on the memory aids or on the family members for the daily activities. The Alzheimer's patients become furious, confused, depressed or anxious and may become angry for no apparent reason. Alzheimer's patients may struggle with the loss of speech and language. Patient may start showing some personality changes. A previously sociable person may withdraw from conversation and activities. People with Alzheimer's may have confusion with time and place. They can lose the track of dates, seasons, and passage of time.

1.3. Effective Alzheimer targets

1.3.1. Amyloid beta

Amyloid beta peptide is a sticky peptide generated by the cleavage of APP by β -secretase followed by an accumulation of γ -secretase in the brain, which finally leads to the formation of amyloid plaques, a key peptide responsible for Alzheimer's disease [28]. The entire process of production of A β 42 is explained by the amyloid cascade hypothesis [20] shown in Fig.1.3.

² https://www.alz.org/alzheimers-dementia/10_signs

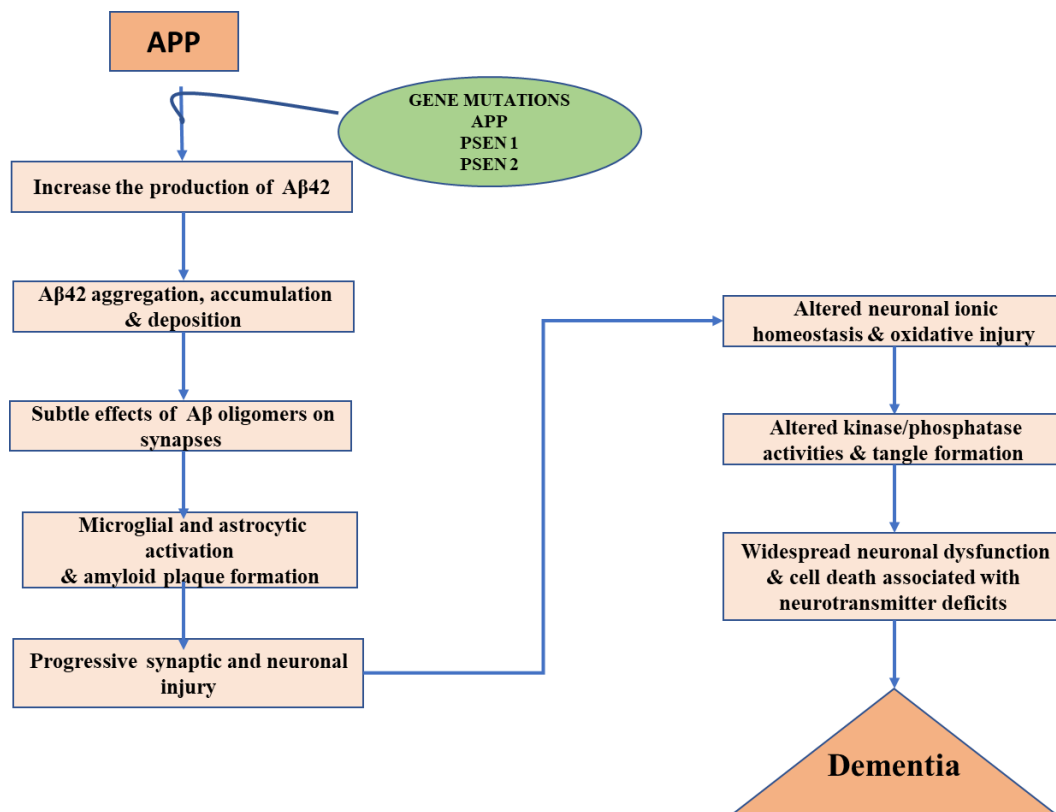


Figure 1.2: Amyloid Cascade Hypothesis

The A β protein is the main target and the core focus of Alzheimer's disease drug development research, and it can be achieved by reducing the A β production which involves either the inhibition of β - or γ - secretase activities, or by enhancing the α - secretase activity [29]. As mentioned earlier, amyloid plaques comprise mainly of amyloid β (A β) peptide generated by the proteolysis of a large β -amyloid precursor protein (APP) [19]. APP can be processed by the proteolytic enzymes called secretases (α , β and γ) in two different pathways [20]. The processing of APP by α -secretase in a non-amyloidogenic pathway produces a soluble fragment sAPP α which has neuroprotective activities. The other pathway is an amyloidogenic pathway in which APP is processed by β -secretase and γ -secretase, resulting in the production of neurotoxic A β peptide (39-43 amino acids) [30], [31]. A β 40 (amyloid peptide containing 40 amino acids) and A β 42 (amyloid peptide containing 42 amino acids) are the major isoforms, but A β 42 is the major contributor to the Alzheimer's disease [32]. Certain gene mutations increase the production of A β 42, and thus its accumulation in the brain induces aggregation to form amyloid plaques [33].

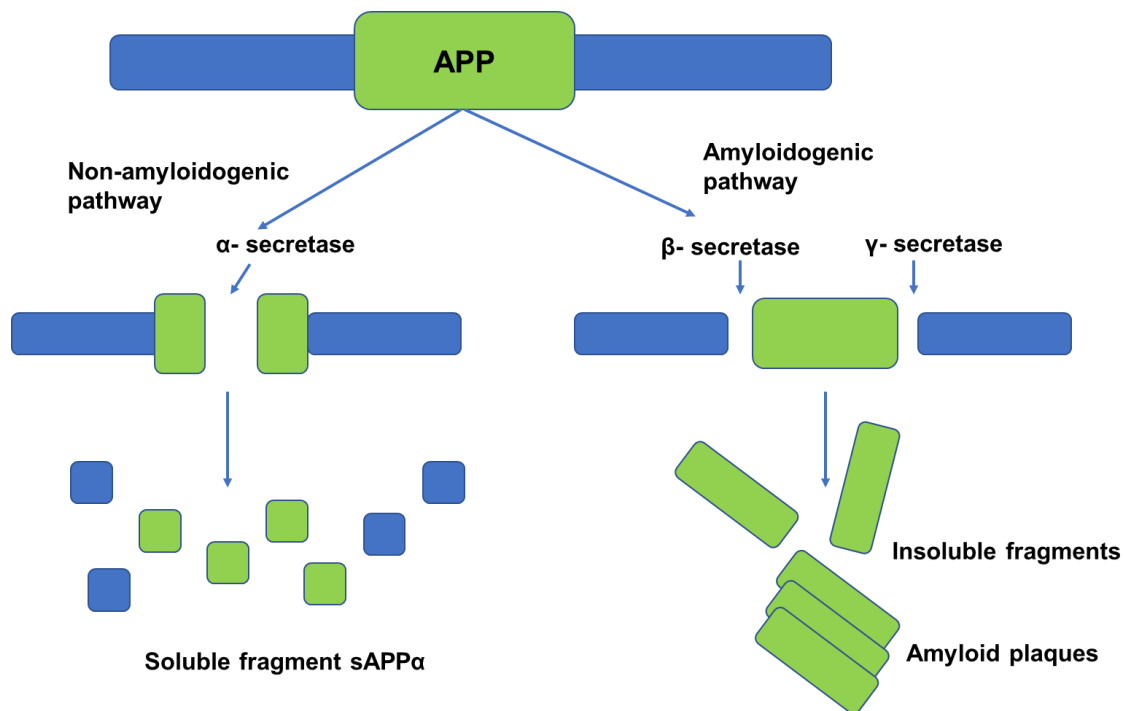


Figure 1.3: Non-amyloidogenic and amyloidogenic pathways for the cleavage of APP

1.3.2. Neurofibrillary tangles

One of the most characteristic neuropathological lesion and the main component of neurofibrillary tangles is the accumulation of paired-helical filaments (PHF) [34]. The PHF consists of hypophosphorylated tau protein, one of the microtubule-associated proteins present in the neurons. The main physiological role of tau protein in the normal neurons is promoting the assembly and stability of axonal microtubules, which play an important role in intraneuronal transport. Tau protein also controls the polymerisation of microtubules [13], [35].

1.3.3. Cholinergic system in AD

The cholinergic system involves the neural pathways in which acetylcholine (ACh) is released for the purpose of neurotransmission. Previous studies identified a pathophysiological significance between cholinergic neurotransmission and amyloidogenesis as well as tau-phosphorylation. This present research supports the hypothesis that the loss of cholinergic innervation is one of the cause of amyloid plaques and neurofibrillary tangles [36].

1.4. Current Alzheimer's treatments

To date there is no cure for Alzheimer's disease, however the drugs currently available in the market have the ability to temporarily slow down and alleviate the symptoms of Alzheimer's and as a result can improve the quality of life for Alzheimer sufferers. The U.S. Food and Drug Administration (FDA) has approved five medications to treat the symptoms of Alzheimer's disease, Table 1.

Table 1.1: FDA-approved drugs

Drug name	Brand name	Approved for	FDA Approved
1. Donepezil	Aricept	All stages	1996
2. Glantamine	Razadyne	Mild to moderate	2001
3. Memantine	Namenda	Moderate to severe	2003
4. Rivastigmine	Exelon	All stages	2000
5. Donepezil and Memantine	Namzaric	Moderate to severe	2014

Unfortunately, the anti-Alzheimer effects of these drugs persist for a limited time, on average 6-12 months. Common side effects are usually mild for these medications and include diarrhea, vomiting, nausea, fatigue, insomnia, loss of appetite, and weight loss.

1.5. Natural products as Potential inhibitors for Alzheimer's disease

Various natural products have shown promise to treat Alzheimer's disease and have contributed towards the search of potential Alzheimer's inhibitors. The phytochemicals derived from these studies could serve as a basis for future research [37]. Some of these promising natural anti-Alzheimer's plants include *Ginkgo biloba*, *Withania somnifera*, *Poncirus trifoliata*, *Hedyotis diffusa*, *Salvia officinalis*, *Huperzia serrata*, *Scrophularia buergeriana*, *Moringa*

olifera, *Bacopa monnieri*, *Rosmarinus officinalis*, *Paeonia suffruticosa*, *uncaria rhynchophylla*, *Galanthus woronowii*, *Panax ginseng*, *Magnolia officinalis*, *Phyllanthus emblica*, *Coriandrum sativum* [38].

Previous studies showed that phenolic rich extracts from *Schotia brachypetala*, *Camelia sinensis*, *Markhamia platycalyx* exhibited notable anti-Alzheimer's activity, when bioassay tests were conducted. These findings led to the conclusion of a possible relation between the intake of polyphenolic rich foods and the prevention of Alzheimer's disease [39].

Flavonoids containing natural products have also demonstrated the ability to improve memory, learning capacity and cognitive function due to their antioxidant properties [40]. Natural flavonoids were also shown to potently reduce the levels of A β deposits either by promoting the α -secretase activity or by inhibiting the β -secretase activity. Another notable inhibitory activity displayed by flavonoids was the reduction of heparin- induced tau formation which is also considered as one of main causes of Alzheimer's disease [40]. The phytochemistry of flavonoid derivatives such as quercetin, myricetin, catechin, epigallocatechin have been extensively studied in the search and hope of finding potential Alzheimer's inhibitors [41].

The *in vitro* and *in vivo* bio-evaluation studies conducted on selected terpenoids such as ginsenosides, ginkgolides, cannabinoids, cornel iridoid glycoside, oleanolic acid, tenuifolin, cryptotanshinone and ursolic acid have suggested that these compounds possess excellent neuroprotective effects [42].

It has been shown that a diet rich in certain edible fruits like blue berry, mulberry, strawberry, grape seed, pomegranate, papaya, apple, green tea, walnut, coffee, saffron, curcumin, pepper, cinnamon, garlic, ginger also play an important role in controlling the Alzheimer's disease [38].

Many studies have been conducted involving the isolation of the active substance responsible for the protective effects against Alzheimer's disease. There are some phytochemicals which have been mostly investigated with a potential of developing drug. These phytochemicals include galanthamine, berberine, curcumin, ginsenoside Rg1, puerarin, silibinin, bilobalide,

quercetin, (-)-epigallocatechin-3-gallate, resveratrol, huperzine A, rosmarinic acid, luteolin, apigenin and chitosan [43], [37].

Galanthamine **(1)**, a natural alkaloid belonging to amaryllidaceae derived from *Galanthus woronowii*, is one of the FDA approved AChE inhibitor drugs currently prescribed for the treatment of Alzheimer's disease. However, this drug is associated with many side effects like frequent nausea, vomiting irregular agitation and sleep disturbance and deprivation [44]. In the early 1960s and 1964, galanthamine was identified as a cholinergic booster and as a reversible inhibitor of acetylcholinesterase (AChE). It was evidently proved to enable its activity in the CNS by penetrating through the blood-brain barrier. [45]

Berberine **(2)** is an isoquinoline alkaloid isolated from *Coptidis rhizoma*. *Coptidis rhizoma* is a major natural product used by the Chinese as a traditional herbal medicine. Previous studies supported that Berberine treatment reduces the reduction of A β 42 in the brain and stimulates the α -secretase activity [46], [47]. Due to its anti-diarrheal, anti-microbial and anti-inflammatory effects, burberine has been used for the treatment of diarrhea, liver diseases, skin and other disorders [48]. The contribution of berberine's multiple activities such as antioxidant activity, AChE and BChE Inhibitory Activity, MAO inhibitory activity, A β level-reducing activity, cholesterol lowering activity and other activities support the fact that berberine may be considered as a potential multipotent natural product to combat Alzheimer's disease [46].

Curcumin **(3)** has many medicinal properties which includes, anti-inflammatory, anti-oxidant, anti-tumour such as chemopreventive and chemotherapeutic properties. Curcumin has been reported for the inhibition of β -secretase and is a promising candidate for the treatment of Alzheimer's disease [49]. Literature has also proposed that curcumin has the ability to inhibit A β -aggregation and A β -induced inflammation in *in vitro* studies [50]. An *in vitro* study revealed the inhibitory effects of curcumin against fibrillar A β aggregation, extension and destabilization at low micro molar concentrations (EC_{50}) between 0.1 and 1.0 μ M [51]. The outcome of an *in vivo* study, highlighted the inhibition of A β -formtion and extension in the brains of AD model mice, when 160-500 ppm (0.43-1.36 μ mol/g) of curcumin was administered orally [50]. These

findings propose that curcumin could be a potential phytochemical candidate in the development of Alzheimer's disease drugs therapy.

Ginsenoside Rg1 (**4**), a pharmacological bioactive component purified from Ginseng. Pre-treatment with the Rg1 helps in the reduction of A β 42 and also in the improvement of cell viability. It also reduced cerebral A β in Tg 2576 mouse model [52], Figure 1.7.

Puerarin (**5**), is one of the major isoflavonoid isolated from the roots of *Pueraria lobata* (Wild.), a Chinese medicine known as Gegen. It has various medicinal properties for the treatment of cardiovascular diseases, cancer, Alzheimer's disease, Parkinson's disease, cancer, cerebrovascular disorders, hyperlipidemia, osteonecrosis etc [53]. Puerarin treatment decreased A β -induced cell death and have a neuroprotective effect in rat hippocampal neurons via various mechanisms [48], Figure 1.8.

Silibinin (**6**), is a flavonoid isolated from the plant *Silybum marianum* commonly known as milk thistle and is known to exhibit anti-oxidative and anti-inflammatory properties. Silibinin was reported to be an inhibitor of A β aggregation and formation in PC12 cells (neuroendocrine in origin), protective against the oxidative damage caused by A β 42 in SH-SY5Y cells (neuroblastoma cells) by decreasing the H₂O₂ levels and also help in the improvement of behavioural abnormalities in mouse model of Alzheimer's disease [54], Figure 1.9.

Bilobalide (**7**), is a sesquiterpenoid lactone derived from *Ginkgo biloba*, a plant which exhibits antioxidant, antiplatelet and hemorrheological properties [55], [56]. Earlier research has shown the significant neuroprotective effects of bilobalide against learning and memory impairments induced by A β ₂₅₋₃₅ in a Morris water maze test, and it also prevented cerebral alterations of oxidative stress and neuronal apoptosis caused in rats [57].

Quercetin (**8**), is a natural occurring flavonoid found in various foods, beverages and plants. It has been reported to play an integral role in the prevention of neurodegenerative diseases [58]. Studies suggested that the pre-treatment of quercetin with neurons for 1 h has notably reduced the A β 42 induced cytotoxicity at lower doses of 5 and 10 μ M (showed the best effects

at 5 μ M), however high doses at 20 and 40 μ M did not show any significant reduction in A β 42 and induced neurotoxicity. Quercetin protected HT-22 murine neuroblastoma cells from A β ₂₅₋₃₅ oxidative attack by strongly inhibiting the A β fibril formation and aggregation [59]. The findings from this work showed that quercetin could be a therapeutic potential for Alzheimer's disease, Figure 1.11.

(-)-Epigallocatechin-3-gallate (**9**), has been reported to activate α -secretase cleavage which promotes non-amyloidogenic APP processing and prevent the A β production [60], Figure 1.12.

Resveratrol (**10**), is a non-flavonoid polyphenol which occurs in grapes and red wine. Previous reports have demonstrated their neuroprotective effects and their potential to reduce amyloid protein concentration due to its allosteric inhibitory activity against the sirtuin proteins [38]. Reports also revealed that resveratrol has the ability to prevent learning impairment, and also reduced the hippocampus neurodegeneration in the p25 transgenic AD mouse model [61], Figure 1.13.

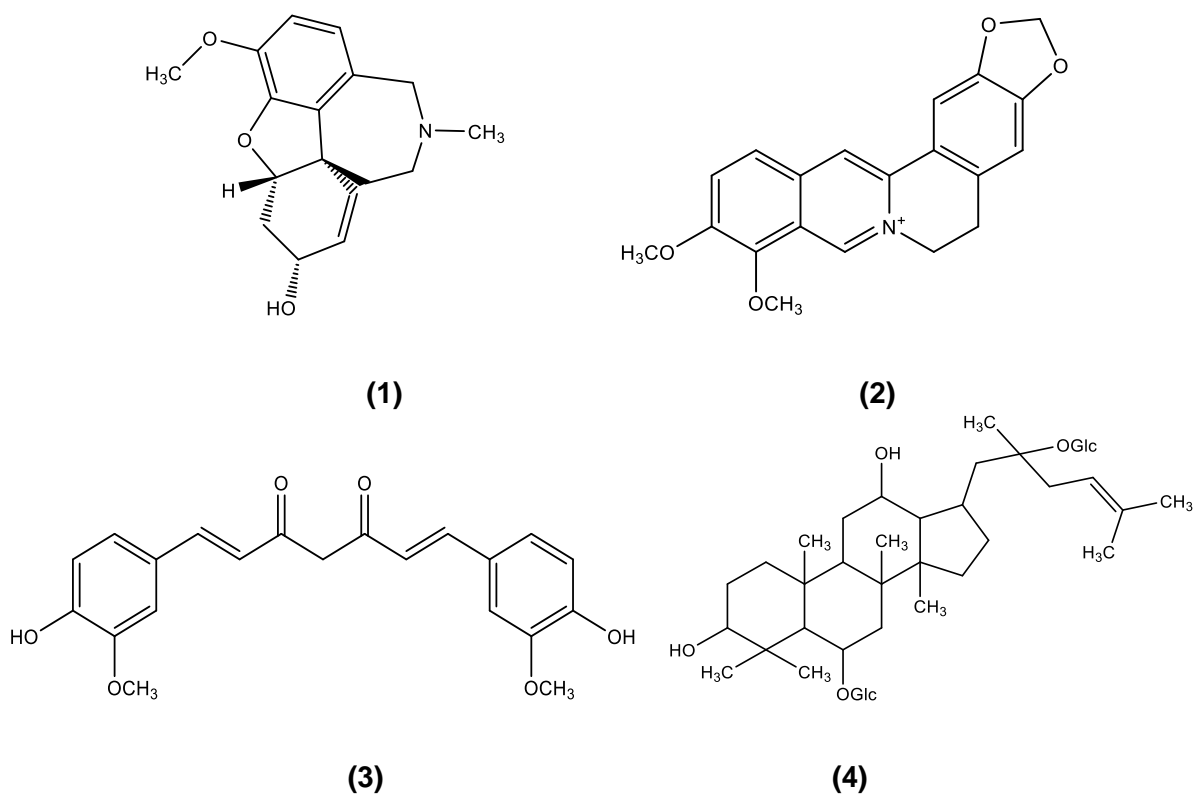
Huperzine A (**11**), the lycopodium alkaloid derived from the Chinese club moss, *Huperzia serrata* is a potent AChE inhibitor. It is used for the treatment of schizophrenia, swelling and strain. Research has shown that it regulates the amyloid precursor protein (APP) and protein kinase C in human embryonic kidney 293 (HEK-293) cells [38]. In China, Huperzine A is widely used as AD therapy enhance the memory, cognitive skills and daily life activities of AD patients [62], Figure 1.14.

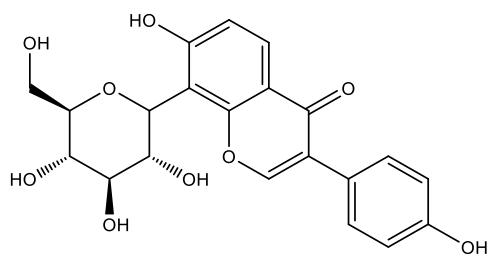
Rosmarinic acid (**12**), a natural scaeanger of peroxyinitrite (ONOO⁻) has been shown to exhibit protective effects on the memory impairment in a mouse model induced by acute i.c.v. injection of A β ₂₅₋₃₅ [63], Figure 1.15.

Luteolin (**13**), a flavonoid derivative is one of the major active ingredients isolated from an edible medicinal plant, *Flos chrysanthemi*. This compound is known to display many pharmacological activities such as anti-oxidant potential, anti-tumorigenic effects and anti-

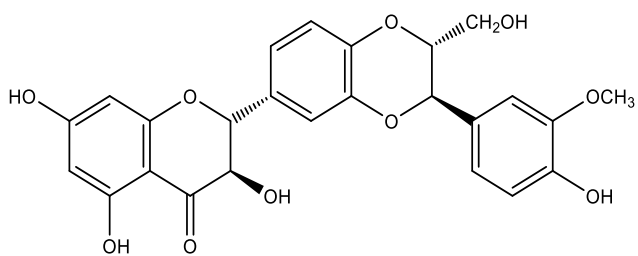
inflammatory activities [64]. Luteolin has revealed to prevent AD-like symptoms after traumatic brain injury in an amyloid- β depositing mouse model [65], Figure 1.16.

Apigenin (4',5,7-trihydroxyflavone) (**14**), is a non-mutagenic flavone, found in many fruits, vegetables and medicinal plants including *Elsholtzia Rugulosa* and *Carduus Crispus*. It has been shown to exhibit anxiolytic, sedative and antidepressant effects and also neuroprotective effects against A β -induced toxicity in rat cerebral microvascular endothelial cells. Results from studies conducted independently have proven the ability of apigenin to reduce learning and memory deficits by decreasing oxidative damage and improving cholinergic neuronal transmission [66], Figure 1.17.

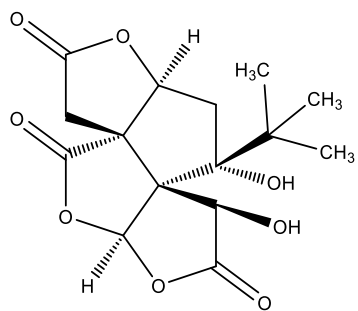




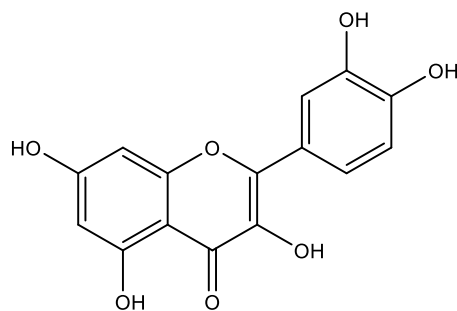
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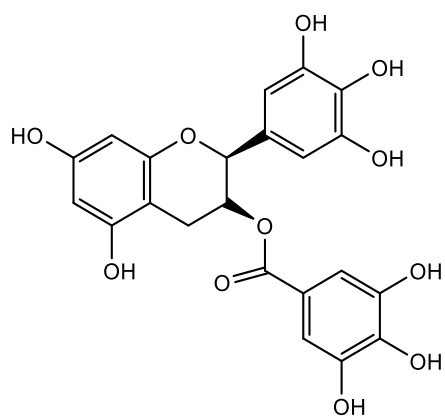
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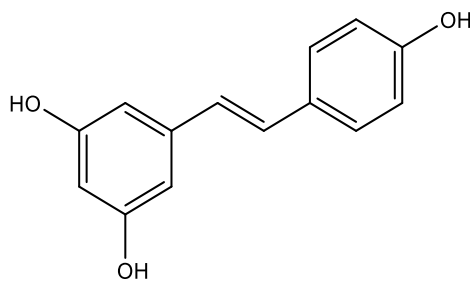
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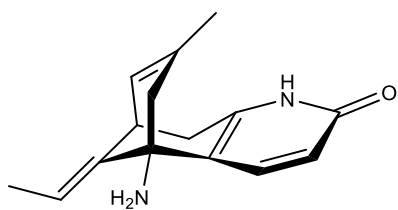
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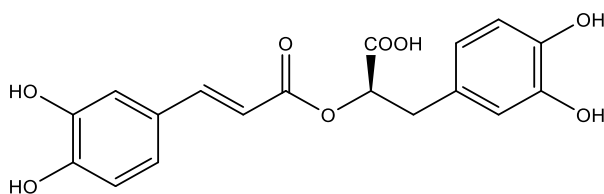
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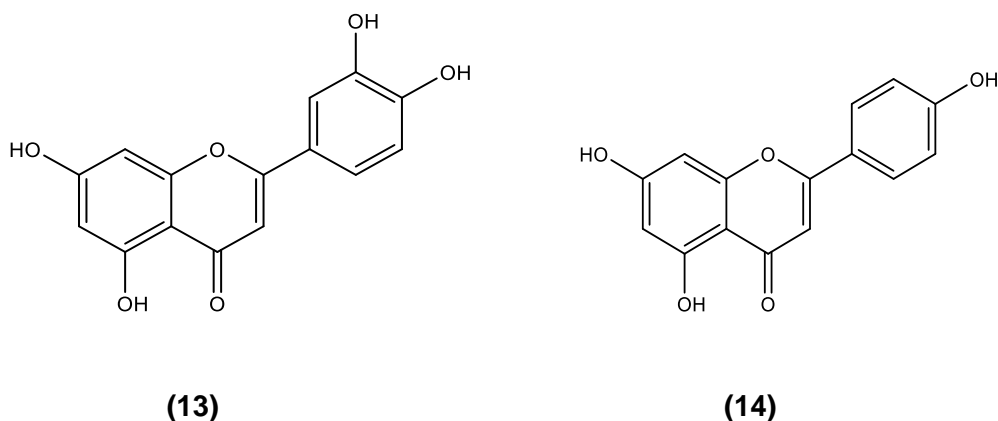


Figure 1.4: Structure of galanthamine (1), berberine (2), curcumin (3), ginsenoside Rg1 (4), puerarin (5), silibinin (6), bilobalide (7), quercetin (8), (-)-epigallocatechin-3-gallate (9), resveratrol (10), huperzine A (11), rosmarinic acid (12), luteolin (13), apigenin (14)

1.6. Problem statement and justification

Regardless of all the research efforts conducted towards understanding the pathology, treatment, and cure of the Alzheimer's disease, to date no cure for this disease has been reported. Consequently, the disease is incurable and irreversible, once an individual start showing distinctive key symptoms related to Alzheimer's disease.

In conclusion, no single treatment has been proven to be significantly effective for the treatment of the Alzheimer's disease. Current FDA drugs used for the treatment of Alzheimer's are overpriced and unaffordable to many or may display undesirable side effects which may be detrimental to human health and well-being.

In South Africa, most of the research is aimed at the diseases such as malaria and HIV-AIDS. However, Alzheimer's disease is also growing slowly and progressively in South Africa and therefore need more attention as such there is a need for developing new drugs in the field of Alzheimer's disease. The outcome of extensive research and studies done in this field, has shown that acetylcholinesterase inhibitors are effective in treatment of cognitive symptoms in

mild to moderate stages of Alzheimer's disease. However these are ineffective against the vascular dementia, suggesting that more attention should be given to other inhibitors, targeting the amyloid plaques and neurofibrillary tangles [67]. The FDA approved acetylcholinesterase drugs also have many side effects. All these approved drugs are temporary treatments for Alzheimer's disease. Hence there is a need for undertaking research for the discovery of new drugs in the field of Alzheimer's disease.

1.7. Aim and objectives

1.7.1. Aim

The aim of this study is to isolate, identify, characterize and develop novel plant-derived anti-Alzheimer's agents from South African's rich, traditional medicinal plant varieties, targeting the amyloid plaques as the precursor enzyme.

1.7.2. Objectives

The overall objectives of the present research are:

- Selection of plants with the help of electronic databases Google scholar, Scopus and Scifinder using the keywords, African traditional medicinal plants in combination with the memory loss, mental illness, depression, Alzheimer's disease, dementia, anxiety, epilepsy, forgetfulness, convulsion, hysteria, sedative effects, acetylcholinesterase inhibition.
- Collection and extraction of different parts of the plant (leaf, stem, root, bark and flowers).
- Fractionation of extracts using silica gel column.
- Identification, isolation and purification of active ingredients using UPLC-MS-QTOF, preparative HPLC.
- A β 42 bioactivity screening of extracts, fractions and pure compounds.
- Structure elucidation of pure compound using 1D, 2D NMR and mass spectrometry.

Schematic flow chart (Figure 1.18) detailing the overall objectives is as described in Figure 1.5.

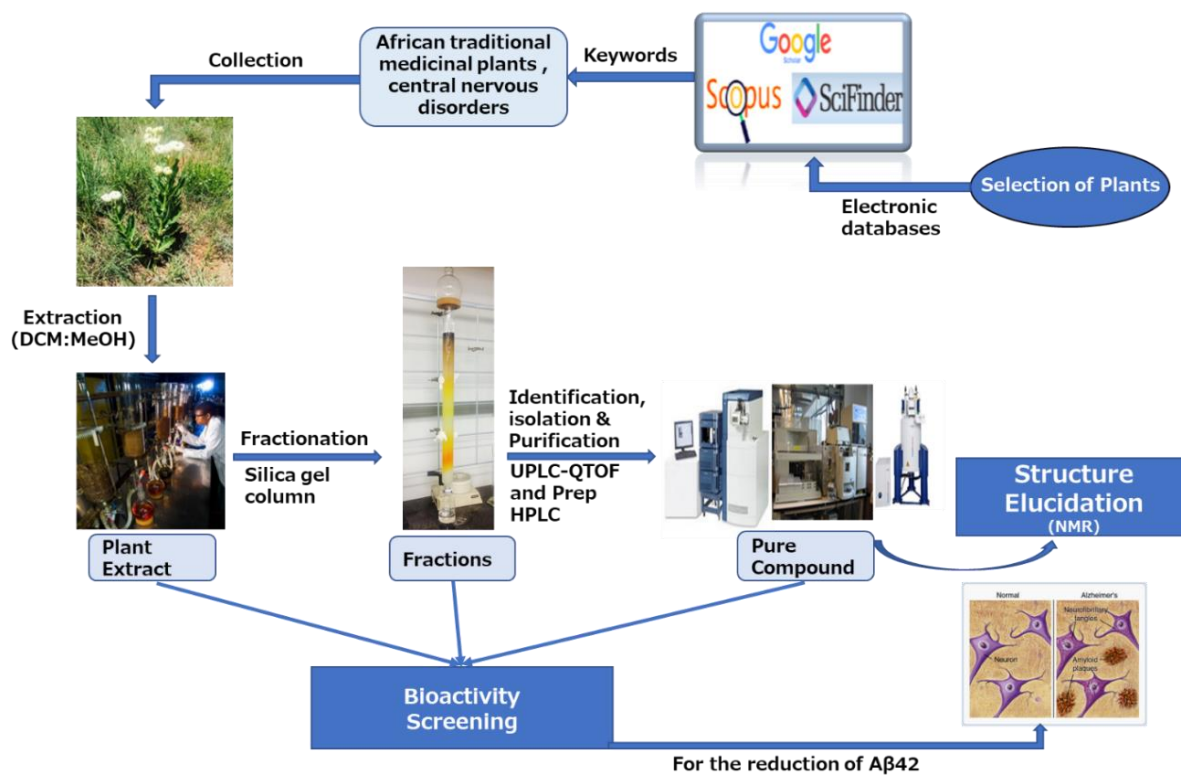


Figure 1.5: Flow chart showing the objectives of present research

The specific objectives of **Chapter 2**:

- Literature survey and selection of plants.
- Plant collection and extraction.
- Bioassay screening of plant extracts for the reduction of A β 42.
- Selection of the active plant extracts for the further investigation.

The specific objectives for **Chapter 3**:

- Collection and extraction of leaves of *Schotia brachypetala*, *Cussonia paniculata* and *Heteromorpha arborescens*.
- Develop chemical profiles of the leaf extracts using UPLC-QTOF-MS for the identification of the major classes of compounds.
- Confirm the presence of compounds by using standards.

The specific objectives for **Chapter 4**:

- Collection and extraction of leaves of *Xysmalobium undulatum*.
- Develop chemical profile of the leaf extract using UPLC-QTOF-MS for the identification of the class of compounds.
- Bioassay-guided fractionation to identify the active ingredients from *X.undulatum* leaf extract based on the A β 42 bioassays.
- Structure elucidation of the active pure compounds using NMR.
- Bioassaying of the pure compounds

The specific objectives for **Chapter 5** include the detailed account of the materials and methods used in the study.

Chapter 6 gives the concluding remarks of the study.

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Chapter 2

In vitro Anti-Alzheimer's activity of medicinal plants native to or naturalised to South Africa

2.1. Introduction

2.1.1. The potential of South African medicinal plants in drug discovery

According to South African National Biodiversity Institute (SANBI), South Africa is the third most diverse country in the world. South Africa owns such a rich plant diversity because of variety of climate and altitude. It exhibits 24,000 plant taxa reported to belong to 368 families. About 15% of these plant species are used for medicinal purposes. Approximately 3000 plant species are used as the traditional medicines [1] and about 80% of South African people use traditional medicines for their health care [2]. Of 3000 plant species, 350 species are traded as medicinal plants and 132 plant species are included on the basis of well-known and fully understood herbal medicines [3].

According to WHO, African traditional medicine is the sum total of all knowledge and practices, whether explicable or not, used in diagnosis, prevention and elimination of physical, mental, or societal imbalance, and relying exclusively on practical experience and observation handed down from generation to generation, whether verbally or in writing [4]. African traditional medicine also known as 'Cradle of Mankind', is the oldest medicinal system and South Africa has a cultural diversity marked by traditional healing practices [5]. Traditional healers are the major source of health services for a large population in South Africa. In South Africa, the traditional doctor is usually a man also known as inyanga who uses herbal medicinal preparations for treating disease [4]. They are also called herbalists with a great knowledge of herbs and medicines of animal origin [6].

In South Africa, approximately 200 000 traditional healers practice herbal medicines [7]. The use of the medicinal plants by South African people has a long history. For example, native

hottentots used the willow species for the treatment of rheumatic fever in the Cape of Good Hope [8]. In addition a Basuto tribal healer used a root decoction prepared from *Pelargonium sidoides* for the cure of tuberculosis of an Englishman, named Charles Stevens [8] which is now commonly traded in Europe as a herbal medicine. In Sub-Saharan Africa, traditional healer practitioners together with the modern health practitioners play a significant role in the treatment of sexually transmitted diseases including HIV/AIDS by integrating the biomedical information of HIV/AIDS in their practices [9]. Bapedi traditional healers from Limpopo province, South Africa used 35 exotic plant species belonging to 21 different families for the treatment of different ailments such as hypertension, diabetes mellitus, erectile dysfunction and gonorrhoea, diarrhoea and tuberculosis. The families, Fabaceae, Solanaceae, Apocynaceae and Asteraceae contain largest number of exotic plant species with medicinal value in traditional healing [10]. In 2002, Steenkamp conducted a survey in which he found that about 10% of the 156 plant species were used for treating gynaecological problems including infertility, menopausal complaints, breast and uterus cancer and menstrual disorders; amenorrhoea, dysmenorrhoea, menorrhagia and irregular menstruation [8], [11]. Two alkaloids, vincristine and vinblastine isolated from *Catharanthus roseus* (Apocynaceae) are well known for their antitumor activity. This plant has been reported for its use as a treatment of breast and uterine cancer [11].

An ethnobotanical survey conducted in Eastern Cape Province; South Africa revealed that 17 plant species from 13 different families were used by traditional healers for the treatment of cancer [12]. From the 13 families, two named Hyacinthaceae and Hypoxidaceae were used prominently and *Solanum aculeastrum*, belonging to family Solanaceae is the most commonly used plant species for cancer treatment [12]. The combretastatin, isolated from *Combretum caffrum*, commonly known as bush willow act as anti-angiogenic agents and played an important role in the treatment of cancer. Six South African plants *Bidens pilosa*, *Centella asiatica*, *Cnicus benedictus*, *Dicoma capensis*, *Hypoxis hemerocallidea* and *Sutherlandia frutescens* were found to exhibit cytotoxicity against two cancer cell line MCF-7 and a non-

malignant breast cell line MCF-12A [13]. *Pelargonium reniforme*, a South African plant species was reported to exhibit antioxidant properties and used for the treatment of liver disorders [14].

The anti-plasmodial activity of 134 plant taxa native to or naturalised in South Africa, belonging to 54 families were tested *in vitro* the results of which showed that 17% were found to exhibit high anti-plasmodial activity [15].

The acetone and aqueous extracts of *Cryptocarya latifolia*, *Helichrysum melanacme*, *Euclea natalensis*, *Nidorella anomala* and *Thymus vulgaris* have been reported to inhibit the growth of *M. tuberculosis* when screened against a drug-sensitive strain, H37Rv by using the agar plate method [16].

The *Sceletium* species are the well-researched and commercially promising plant species among the South African plants from a pharmaceutical point of view [17]. *Sceletium tortuosum* has been reported to reduce anxiety, stress or depression and has also been reported to elevate mood in a human clinical trial [18]. The standardised *S. tortuosum* extract, Zembrin®, has been shown to inhibit phosphodiesterase-4 (PDE-4) with an IC₅₀ value of 8.5 µg/ml and also act as a potent blocker in 5-HT transporter binding assay with an IC₅₀ value of 4.3 µg/ml [19]. Three major alkaloids mesembrine, mesembrenone and mesembrenol isolated from *S. tortuosum* are responsible for the potent activity of the plant and affect many CNS disorder targets [18].

There are few African plants which have been recognised for their commercialization and role in drug discovery on a global scale. These plants are *Xysmalobium undulatum*, *Agathosma betulina*, *Aloe ferox*, *Aspalathus linearis*, *Artemisia afra*, *Cyclopia genistoides*, *Hoodia gordonii*, *Harpagophytum procumbens*, *Hypoxis hemerocallidea*, *Lippia javanica*, *Warburgia salutaris*, *Pelargonium sidoides* and *Sutherlandia frutescens* [20].

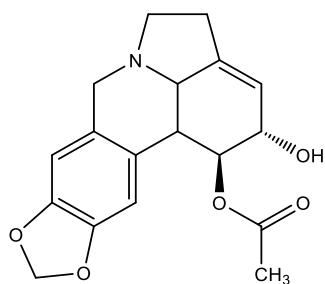
South Africa with a rich biodiversity has resulted in the isolation and discovery of several potent bioactive compounds. Thus, in this dissertation, finding novel potential plant-derived anti-Alzheimer compounds from the South African medicinal plants, through conducting bioassay

guided fractionation, employing standard isolation and modern spectroscopic techniques, underpins the main focus and aim of this study.

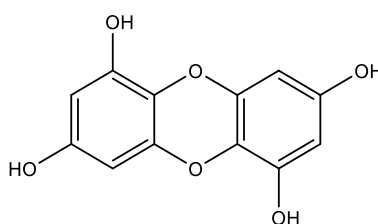
2.1.2. South African medicinal plants with acetylcholinesterase inhibition and reduction of A β 42 for treating Alzheimer's disease

Acetylcholinesterase inhibitors are considered as having one of the promising therapeutic potential in the field of Alzheimer's disease [21]. The majority of research done on South African medicinal plants for treating Alzheimer's disease has focused on the inhibition of acetylcholinesterase. Alkaloids isolated plant extracts *Boophone disticha*, *Crinum campanulatum*, *Crinum moorei*, *Crinum macowanii*, *Crinum bulbispermum* and *Cyrtanthus falcatus* belonging to family *Amaryllidaceae* have been reported for AChE inhibition activity [22]. One of the alkaloid, 1-O-acetyllycorine (**15**) was found to be potently active against AChE inhibitory with an IC₅₀ value of 0.96 ± 0.04 μM while galanthamine exhibited IC₅₀ value 1.9 ± 0.16 μM in the same assay indicating that the compound was twice as potent as galanthamine [23]. Withanoloids isolated from *Withania somnifera* (L.) Dun belonging to family Solanaceae, inhibited AChE and BuChE dose dependently. It also significantly regenerated axons and dendrites in memory- deficient mice [24]. The DCM:MeOH root extracts of *Chamaecrista mimosoides* potently inhibited the AChE activity with IC₅₀ value 0.03 ± 0.08 μg/ml [25]. Aqueous and ethanol extracts of *Crinum moorei* and *Boophone disticha* showed promising results for the treatment of Alzheimer's disease by potently inhibiting AChE activity in the TLC bio-autography assay. Both species belong to *Amaryllidaceae* family, known for containing alkaloids which are responsible for the inhibition of AChE [26]. Phenolic compounds isolated from *Croton gratissimus* exhibited good acetylcholinesterase inhibitory activity [27]. Phlorotannins, dibenzo [1,4] dioxine-2,4,7,9-tetraol (**16**) and eckol (**17**) isolated from *Ecklonia maxima* have been reported to exhibit AChE inhibitory activity with an IC₅₀ value of 84.48 ± 0.26 and 76.70 ± 0.35, respectively as compared to the galanthamine [28]. A crinine alkaloid, 6-hydroxycrinamine (**18**) isolated from methanol extract of the bulbs of *B. disticha* was found to exhibit AChE inhibitory activity with an IC₅₀ value of 445 μM. Although the

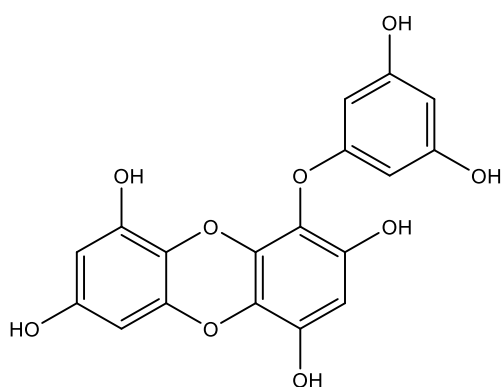
compound was active, it showed cytotoxicity towards neuroblastoma cells [29]. The three compounds, hydroxyphenyl ethyl vanillate (**19**), acteoside (**20**) and quercetin (**21**) isolated from *Buddleja salviifolia* showed AChE inhibition activity with an IC_{50} values of 259.5 ± 0.00 , 123.1 ± 1.23 , 66.8 ± 0.09 , respectively as compared to galanthamine [30]. Assoanion (**22**), one of the alkaloids isolated from *Cyrtanthus contractus* showed very good AChE activity with an IC_{50} value of $3.87 \pm 0.24 \mu\text{M}$ as compared to the control galanthamine with an IC_{50} value of $1.9 \pm 0.16 \mu\text{M}$ [31].



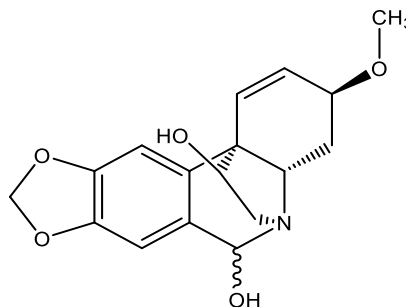
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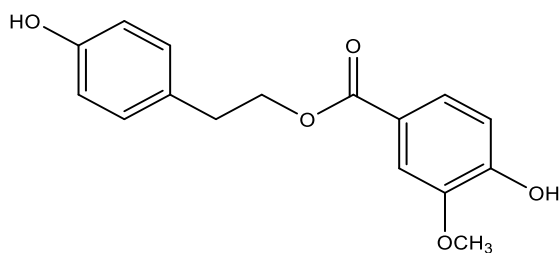
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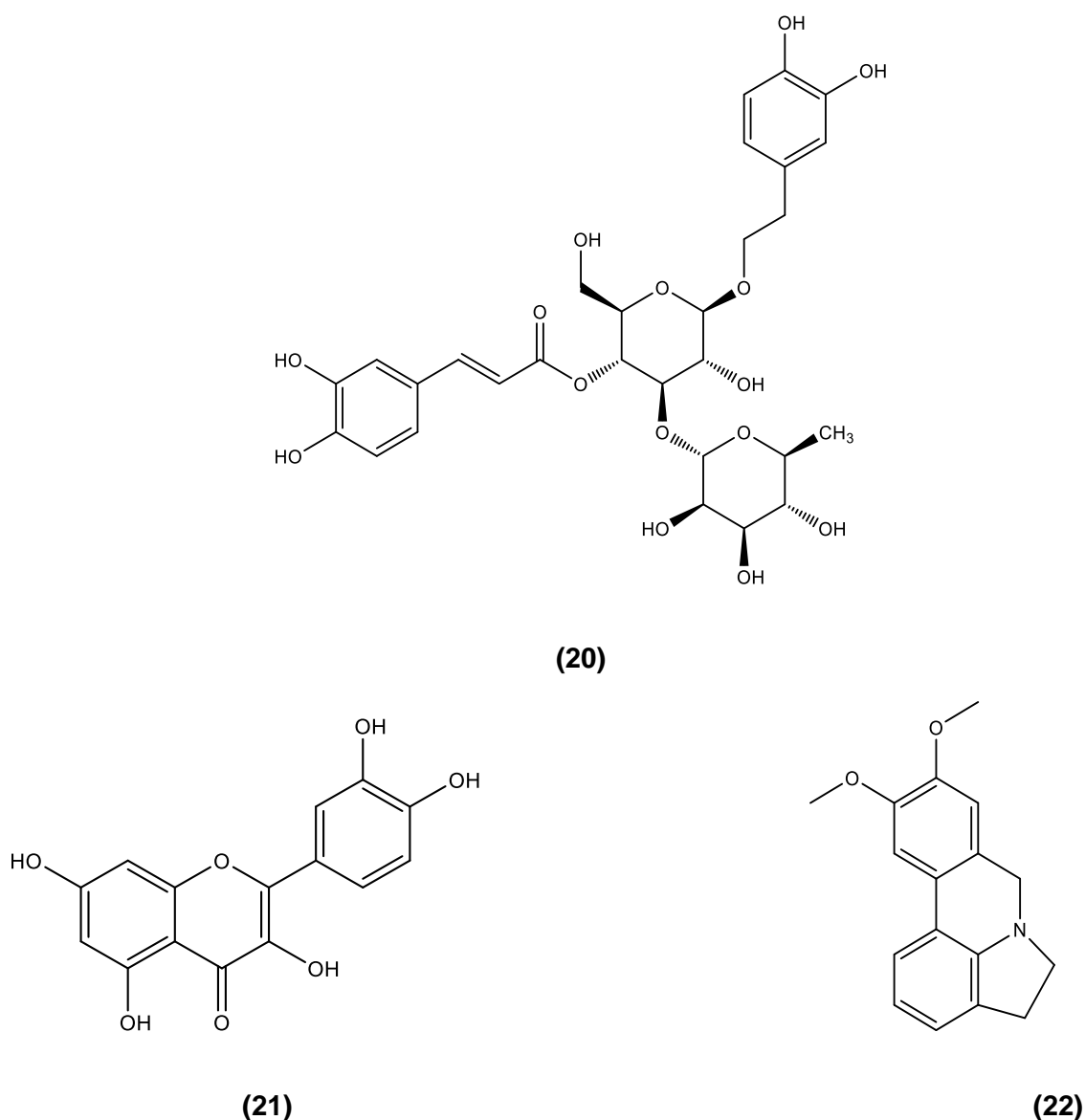


Figure 2.1: Structures of the compounds exhibited acetylcholinesterase inhibition activity, 1-O-acetyllycorine (**15**), dibenzo [1,4] dioxine-2,4,7,9-tetraol (**16**), eckol (**17**), 6-hydroxycrinamine (**18**), hydroxyphenyl ethyl vanillate (**19**), acteoside (**20**), quercetin (**21**) and assoanion (**22**)

One of the previous studies showed that phenolic rich extracts of three plants *Schotia brachypetala*, *Camellia sinensis*, *Markhamia platycalyx* were able to significantly reduce A β 42 and could lead to the discovery of anti-Alzheimer's drugs [32]. The ethyl acetate extracts of *Xysmalobium undulatum* also exhibited acetylcholinesterase inhibition activity in *in vitro* studies [33]. Lycorine (**23**) isolated from *Euclea crispa* and 3-Oxo-oleanolic acid (**24**) isolated

from *Crinum macowanii* exhibited the significant A β 42 reduction from APPsw-transfected HeLa cells in a dose dependent manner [34]. The extracts of *Monsonia* species belong to family Geraniaceae were found very active in suppressing the A β 42 production in the brain [35].

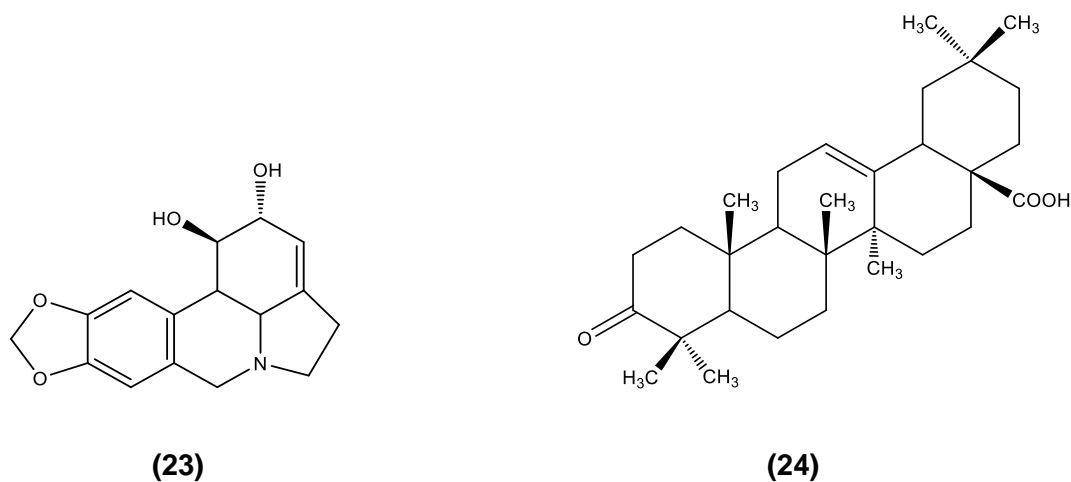


Figure 2.2: Structure of the compounds exhibited A β 42 reduction, Lycorine **(23)**, 3-oxo-oleanolic acid **(24)**.

Many plants native to or naturalised to South Africa were researched for mental disorders and Alzheimer's disease through A β 42 and AChE inhibition in *in vitro* and *in vivo* studies. These studies provided evidence that South African plant species could lead to new drug leads for the pharmaceutical industry in the field of Alzheimer's disease.

2.1.3. Need for research on Alzheimer's disease in South Africa

In 2003/4, the South Africa Stress and Health (SASH) conducted a study which estimated that approximately one in three South Africans suffer from some form of mental disorders [36]. In 2014, the Universities of Cape Town and Stellenbosch, further ratified this study and came to conclusion that mental disorders is probably higher than one in three of the South African population [36].

Another report on “sick state of mental health” in South Africa published by the country’s Sunday Times Newspaper stated that “More than 17-million people are dealing with depression, substance abuse, anxiety, bipolar disorder and schizophrenia” [37]. These surveys showed that mental illnesses is becoming a serious threat to South Africa and needs special attention. In South Africa, a database analysis was conducted on the prescription pattern and cost of drugs for Alzheimer’s disease. This study revealed that a total of 588 patients were found who received 2623 medicine items with an average cost per item R596.15 (South African Rand). This cost may be unaffordable for the South African people. This data analysis indicated that either the Alzheimer’s disease is under diagnosed or the medicine for Alzheimer’s disease is under-prescribed in South Africa because of the cost effectiveness [38]. Later it was found that approximately 750 000 people in South Africa are suffering from Alzheimer’s disease [39]. Due to the rapidly growing problem of mental disorders such as Alzheimer’s disease in South Africa, there is a need to focus on it. South African plants have great therapeutic potentials and can also lead to the discovery of novel and affordable drugs in the field of Alzheimer’s disease. This study thus focuses on the medicinal plants of South Africa with traditional uses associated with mental disorders to determine their efficacy by screening the different plant part extracts against A β 42 as a potential treatment of Alzheimer’s disease.

2.2. Materials and Methods

2.2.1. Selection and collection of plant material

A literature study was undertaken using Google scholar, SciFinder and Scopus for the selection of the plants by using the keywords; African medicinal plants and traditional medicinal plants in combination with the memory loss, mental illness, depression, Alzheimer’s disease, dementia, anxiety, epilepsy, forgetfulness, convulsion, hysteria, sedative effects, acetylcholinesterase inhibition. The identified plants were then prioritized by a scoring system

of 1, 2 and 3 where 1 represents strong, 2 medium and 3 weak. The criteria used for scoring was as follows

a) Strength of the traditional uses for Alzheimer's disease, in which the plants with direct relation with Alzheimer's disease either through A β 42 reduction or by AChE inhibition were given a score of 1, plants with traditional uses such as memory loss were given a score of 2 and the plants with other traditional uses such as epilepsy, hysteria, headache and sedative effects were given a score of 3.

b) Strength of published use for Alzheimer's disease (peer reviewed) - plants with significant published information for Alzheimer's disease were given a score of 1, plants with less and no published information for Alzheimer's disease were given scores 2 and 3, respectively.

c) Plant toxicity - plants with no or limited toxicity were given a score of 1 and plants with medium and high toxicity were given scores of 2 and 3, respectively.

d) Plant part used - leaves given a score of 1, stems and stalks given a score of 2 and roots given a score of 3. The schematic process of selection of plants is shown in Figure 2.3.

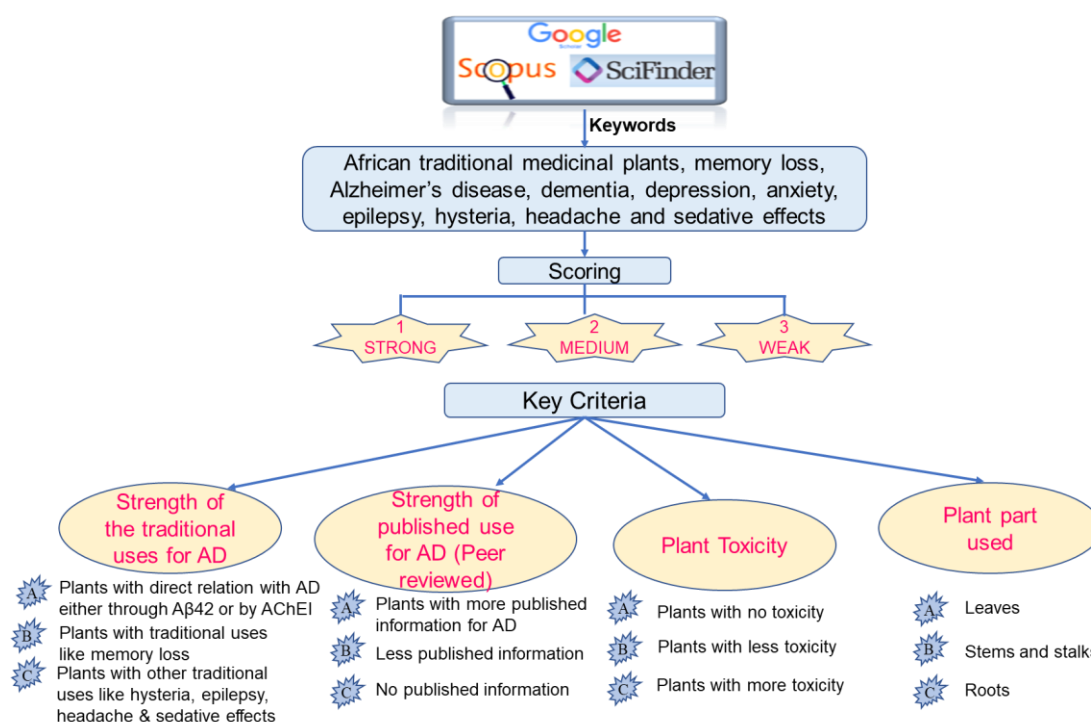


Figure 2.3: Criteria for the selection of plants species

2.2.2. Collection and extraction of plants

Plants with their different parts were collected from *Maine vander Schijff* Botanical Garden and experimental farms of University of Pretoria, South Africa with the help of Curator Jason Samuels. Plant identification was established at the H.G.W.J. Schweickerdt Plant Herbarium at of Pretoria University. Plant parts were separately cut into small pieces and then dried in an oven at 60°C. The drying time varied depending on the nature of plant and the plant part. Dried plant material was ground to a coarse or fine powder using a grinder. Each of the powdered plant materials (2g) were extracted repeatedly three times for 8 hours with dichloromethane/methanol (DCM/MeOH) (1:1) with the help of a magnetic stirrer. Extracts were filtered using a Buchner funnel. Plant extracts obtained from three different extractions were combined and then concentrated using a rotary vacuum evaporator and dried under high

vacuum. All the extracts were stored in a cold room prior to screening prior to bioassaying in collaboration with the Korean Institute of Science and Technology (KIST).

2.2.3. Cell culture

HeLa cells stably transfected with APP carrying Swedish mutation (APP^{sw}) were cultured at 37°C, 5% CO₂, in Dulbecco's Modified Eagle Medium (DMEM) supplemented with 10% heat inactivated fetal bovine serum containing 100 units/ml penicillin, 100 µg/ml streptomycin, 260 µg/ml Zeocin, 400 µg/ml G418.

2.2.4. Aβ₄₂ inhibition bioassay

APP^{sw}-transfected HeLa cells at 80% confluence in a 35mm dish were cultured with sample solubilized in dimethyl sulphoxide (DMSO) for 8 h in serum free medium. The conditioned medium was analyzed by a sandwich ELISA (Invitrogen) for detection of Aβ₄₂ according to the supplier's instruction.

2.2.5. Statistical analysis

Data was expressed as mean ± SEM. Statistical comparisons between controls and treated experimental groups were performed using the Student's *t*-test. *P* < 0.05 was considered statistically significant.

2.3. Results and discussion

2.3.1. Selection of plant material

Using the selection criteria and the databases searches, at least 52 plant species were identified as candidates for screening their potential for Aβ₄₂ reduction. As this was a broad general search, the additional criteria and scoring system was applied which resulted in 21 plant species being identified for the collection, extraction and screening. A scoring system

which was shown to be useful was adapted from the previous studies [40], [15]. In instances when the criteria could not be applied for the plant species as the information may not have been publicly available, these were excluded from the scoring system. This resulted in the identification of 21 species which had scores between 1 and 9. Table 2.1 shows the plants species identified through the scoring system with the scores assigned for each of the criteria and the total scores per plant. These species represented 15 different families.

Plant species with the lowest scores representing the highest priority based on the criteria used where *Mondia whitei* and *Schotia brachypetala* with scores of 4. Two plants species *Centella asiatica* and *Trichilia dregeana* scored 5; five plants species, *Commelina africana*, *Cussonia paniculata*, *Cussonia spicata*, *Tetradenia riparia* and *Zanthoxylum capense* had a score of 6; seven plants, *Buddleja salvifolia*, *Bulbine natalensis*, *Catha edulis*, *Heteromorpha arborescens*, *Ruta graveolens*, *Xysmalobium undulatum* and *Ziziphus mucronata* had a score of 7; three plants *Cissus quadrangularis*, *Cotyledon orbiculata*, and *Tabernaemontana elegans* scored 8 and two plants *Plumbago auriculata* and *Stapelia gigantea* with the lowest score of 9. The published phytochemistry of the plants was also established through literature searches and this was done to guide any further fractionation process to identify the active ingredients and to provide any information on whether some of these compounds may have been published for the treatment of Alzheimer's disease.

Table 2.1: Scoring for selection of plant species

Plant species and Family	Traditional use (A)	Previously reported bioassays (B)	Toxicity (C)	Phytochemistry	Total Score (T) (A)+(B)+(C) = (T)
<i>Buddleja salvifolia</i>	Traditionally used in the treatment of neurodegenerative disorders. <i>Buddleja</i> species together with <i>Heteromorpha trifoliata</i> & <i>Cussonia</i>	<i>B. salvifolia</i> leaf extracts showed significant antimicrobial and acetylcholinesterase inhibitory properties [30]. The whole plant showed good antioxidant	Highly toxic, DCM and MeOH leaf extract WSI Cell line, IC ₅₀ >200µg/ml [41], [42]	4 hydroxyphenyl ethyl vanillate, Acteoside and quercetin [30].	

	<p><i>paniculata</i> are used for the treatment of early nervous and mental disorders in South Africa [25].</p> <p>Score- 2</p>	<p>activity with ABTS; IC₅₀ = 0.14 ± 0.08 mg/ml and DPPH; IC₅₀ = 0.23 ± 0.01 mg/ml and DCM:MeOH (1:1) showed IC₅₀ value of 0.05 ± 0.02 mg/ml as compared to the positive control, galanthamine with IC₅₀ value of 5.3 × 10⁻⁴ mg/ml [25].</p> <p>Score- 2</p>			
<p><i>Bulbine natalensis</i></p>	<p>Tuber decoctions are used for the treatment of convulsions in South Africa [43].</p>	<p>The biochemical analysis of the excisional wound tissue of <i>B. natalensis</i> treated wounds showed a significant increase in the total collagen content compared to that of the untreated wounds [44]. The <i>in vitro</i> antimicrobial activity of ethanolic extract, n-butanol and ethyl acetate fractions of 4 Gram positive and 12 Gram negative bacteria as well as 3 fungal species were using agar dilution inhibited 75, 87.5 and 100% respectively and produced growth inhibition at MIC range of 1-10, 3-10 as well as 1 and 5</p>	<p>Low toxic, Leaf extract, Brine shrimp Cell line, IC₅₀= 2210.0µg/ml. [46];</p> <p>Mixed cytotoxicity at different concentration, Cell line, HEp-2</p>	<p>Tannins, anthraquinones, cardiac glycosides, saponins, and alkaloids [47].</p>	<p>7</p>

	Score- 3	mg/ml respectively [45]. Score- 3	Score- 1		7
<i>Catha edulis</i>	Fresh leaves reduce fatigue and relieve sleepiness. It increases mental power and communication skills but also causes hypertension and affects male fertility [48].	The khat extracts showed selective effect on spatial learning and memory in CBA mice. The high dose (360 mg/kg body weight) of khat extract improved the memory but impaired learning while the moderate (120 mg/kg body weight) and low dose (40 mg/kg body weight) impaired accuracy for spatial memory of the platform location [49].	Highly toxic, leukemia cell line [50].	The major alkaloids found in Khat is phenylalkylamines and cathedulins. Phenylalkylamines contain (+)-Cathine; (-)-Cathinone; 3,6-Dimethyl-2,5-diphenyl pyrazine; (-)-Norephedrine; (-)-Norephedrine N-formyl; Merucathine; Merucathinone; 1-Phenylpropane-1,2-dione; Pseudomerucathine. Cathedulins contain Cathedulin E1–E6; Cathedulin K1, 2, 6, 12, 15, 17, 19, 20; Cathedulin Y7–Y10; Cathidine A; Cathidine B; Cathidine D [51].	
	Score- 2	Score- 2	Score- 3		7
<i>Centella asiatica</i>	Dried powdered leaves reportedly produce a calming and sedative effect and in some parts of India it is given with milk to improve memory against dementia [24].	The extracts of <i>C. asiatica</i> have shown effective to increase the antioxidant enzymes in corpus striatum and hippocampus in the aged rats and rats with Parkinson's disease and also prevented the	Low cytotoxicity, Whole plant, Cell line L6, IC ₅₀ = 82.6 µg/ml [42], [53].	Triterpenoid saponins including asiaticosides, madecassoside and madsiatric acid. brahmoside and brahminode, isothankuniside and thankuniside, plant sterols,	

		oxidation of proteins, lipid peroxidation and prooxidant processes in them [52].		flavonoids, centilioside and its derivatives [54]	
	Score- 2	Score- 2	Score- 1		5
<i>Cissus quadrangularis</i>	Boiled with water and used for the treatment of epilepsy in Dar es Salaam, Tanzania [55].	Bone fracture healing activity, analgesic activity, antiosteoporotic activity, antioxidant activity, antiulcer activity, Parasympathomimetic activity, Anabolic and Androgenic activity, Antihemorrhoidal activity, Anti-inflammatory activity and Gastroprotective Activity [56].	Moderately toxic MeOH stem extract Cell line- Brine shrimp IC ₅₀ = 1300.0µg/ml [57], [42].	Tetracyclic triterpenoids- stilbenes resveratrol, piceatannol, pallidol; stilbene derivatives- quadrangularins A, B, and C, parthenocissin A and the known flavonols- quercetin and kaempferol [58]. Alkaloids, resveratrol, piceatannol, pallidol, parthenocissin, quadrangularins, ascorbic acid, carotene, phytosterol substances, calcium, flavinoids, vitamins, enzymes, nicotinic acid, tyrosin, and triterpenoids [59]	
	Score- 3	Score- 3	Score-2		8
<i>Commelina Africana</i>	In Lesotho plant decoctions with <i>Tephrosia capensis</i> are used for heart	<i>C. africana</i> reduced fasting blood glucose of experimental diabetic rats by	Low toxic, Vero cell line, IC ₅₀ = 441.1µg/ml [61].	Alkaloids, cardenolides, saponins and tannins [60].	

	complaints and nervous disorders [24]. Score- 2	78.0%. Thus showed a comparable activity with the reference hypoglycaemic agent [60]. Score- 3	Score- 1		6
<i>Cotyledon orbiculata</i>	Leaf juice has been used for the treatment of epilepsy [48]. Score- 3	<i>C. orbiculata</i> has shown the anticonvulsant activity which supports the fact of its traditional use for the treatment of epilepsy [62]. Water extract of <i>C. orbiculata</i> at 7.5 mg/ml prevented nematode eggs from hatching with 82.63% success rate and suppressed nematode larval development with 85.32% at the concentration of 2.5 mg/ml which showed that it possess anthelmintic activity [63]. Score- 3	Moderate toxic, Cell line, HEK 293 IC ₅₀ = 44.63µg/ml [64]. Score- 2	Alkaloids, saponins, tanins, cardiac glycosides, triterpene steroids, reducing sugar, flavonoids and anthraquinones [62]. Four bufadienolides, namely orbicusides A,B and C and tyledoside C [65].	8
<i>Cussonia paniculata</i>	Used for the treatment of rheumatism, dysmenorrhea, colic and diseases associated with the nervous system [66]. it is used as a traditional medicine against	The stem bark of <i>C. paniculata</i> showed anti-inflammatory and analgesic activities. DCM:MeOH leaf extract of <i>C. paniculata</i> screened for the in vitro anticancer activity and showed	Low toxic, APPsw-transfected HeLa cells Toxic only at high concentration 100µg/ml [69].	Triterpenoid glycosides [70], sterol/triterpene, flavonoids such as rutin, quercetin and luteolin [71].	

	<p>pain, inflammation, infections and malaria. The species is also used for the treatment of mental illness, diarrhea and sexually transmitted diseases [67]. The thick tuberous roots are peeled and eaten raw as an emergency food or as a source of water. In South Africa, the mixture of <i>Buddleja</i> species with <i>Heteromorpha trifoliata</i> and <i>Cussonia paniculata</i> is used for the treatment of early nervous and mental disorders [25].</p> <p>Score- 2</p>	<p>an inhibition of 100% cell growth at a concentration of 5.25 (renalTK10), 15.0 (melanoma UACC62) and 55.69 g/ml (breast MCF7) [68] .</p> <p>Score- 3</p>	<p>Score- 1</p>		<p>6</p>
<i>Cussonia spicata</i>	<p>The succulent and edible mashed roots of <i>C. spicata</i> are used for the treatment of Malaria. The root decoction is used to treat fever and mental illness. The leaves are used for the treatment of indigestion. It is also used for the</p>	<p>Disc diffusion bioassays for <i>C. spicata</i>, the ethanolic and ethyl acetate extracts of the bark and roots showed inhibitory activity with the MIC'S value of 12.5 mg/ml against <i>Staphylococcus aureus</i> [67].</p>	<p>Low toxic, Water extract of root, Brine shrimp Cell line IC₅₀= 2600.0µg/ml [72].</p>	<p>Molluscicidal saponins, [α -L- arabinofuranosyl-(1 \rightarrow 4)- β -D- glucuronopyranosyl-(1 \rightarrow 3)]-3 β - hydroxyolean-12-en-28-oic acid and [(α -L- arabinofuranosyl-(1 \rightarrow 4)- β -D-</p>	

	treatment of epilepsy in Dar es Salaam, Tanzania [55].			galactopyranosyl-(1 → 2)- β -D-glucuronopyranosyl-(1 → 3)]-3 β -hydroxyolean-12-en-28-oic acid were isolated from <i>C. spicata</i> [73].	
	Score- 2	Score- 3	Score- 1		6
<i>Heteromorpha arborescens</i>	Used to treat headache, pain, fever, dysmenorrhea, nervous and mental disorders [74], [75].	The organic extracts of root showed a good COX-1 inhibition activity [74]. The methanolic and the aqueous root extracts showed significant pain inhibition of carrageenan induced inflammatory pain and also the significant reduction of edema induced by albumin and carrageenan [75]. The leaf extract of <i>H. arborescens</i> inhibited nitric oxide production by 89.98% at a concentration of 30µg/ml and also showed a good <i>in vitro</i> anti-arthritic activity with IC ₅₀ value of 53.78 µg/ml [76].	Highly toxic, Leaf extract, Vero monkey kidney Cell line, IC ₅₀ = 0.04mg/ml [77]	Saponins, saikogenin B4 and saikogenin F; Falcarindiol, a polyacetylene and sarisan. [74], [78] .	
	Score- 2	Score- 2	Score- 3		7
<i>Mondia whitei</i>	Root infusions are taken by unspecified groups in South	The plant showed a strong acetylcholinesterase inhibition	Low toxic, Breast cancer cell lines, MDA-MB-231,	Alkaloids, flavonoids, saponins, tanins, polyphenols,	

	<p>Africa to treat stress in adults [24]. It is used as a traditional medicine to treat nervous disorders. Pulverized root is used to treat schistosomiasis and abdominal pain and also taken in water for headache [79].</p> <p>Score- 2</p>	<p>bioactivity and acted as an antibacterial agent. [80]. A compound monoterpene–lactone loliolide isolated from <i>M. whitei</i> leaves have shown affinity to SERT in a binding assay. The activity of compound with an IC₅₀ value of 1.3 nM (Ki=0.6nM) than the activity of Citalopram [81].</p> <p>Score- 1</p>	<p>MCF-7 and MCF-12A [82].</p> <p>Score- 1</p>	<p>moderate anthranal glycosides and terpenoids [83]. Chlorinated coumarinolignan, 5-chloropropacin [84].</p>	<p>4</p>
<p><i>Plumbago auriculata</i></p>	<p>Used for headache in Zulu, Xhosa and Sotho medicine [85].</p> <p>Score- 3</p>	<p><i>P. auriculata</i> showed a significant <i>in vitro</i> antimalarial activity with different solvent extracts. n-hexane extract showed IC₅₀ 45.9 ± 3.3µg/ml, DMC extract showed 40.2 ± 1.6 µg/ml; 53.8 ± 3.1 µg/ml; 80.0 ± 15.1µg/ml [86]. Plumbagin isolated from <i>P. auriculata</i> inhibited the seed germination of lettuce and wheat [87].</p> <p>Score- 3</p>	<p>High toxic, HGE-17 cell line, MTT assay, IC₅₀= 278.59µg/ml [88].</p> <p>Score- 3</p>	<p>Plumbagin, episinoshinanolone, plumbagic acid, palmitic acid, sitosterol and 3-Oglycosylsitosterol [89].</p>	<p>9</p>
<p><i>Ruta graveolens</i></p>	<p>Leaf infusions are taken for epilepsy, hysteria, convulsions and fits in children [48]. It has been used as a traditional medicine for the</p>	<p>It has been shown to exhibit antibacterial, analgesic, anti-inflammatory, antidiabetic, insecticidal, anticancer, xanthine oxidase inhibition,</p>	<p>High toxic, Hep2 cell line, IC₅₀= 62µg/ml [92].</p>	<p>Rutin, quercetin, psoralen, methoxypsoralen, rutacridone, rutacridone epoxide and gravacridondiol and the main</p>	

	treatment of pain relief, eye problems, rheumatism and dermatitis. It has been used as an antidote for the snake toxins and snake venoms [90]. Score- 3	antihyperlipidemic and antiandrogenic activities [90]. The acridone alkaloid arborinine isolated from <i>R. graveolens</i> exhibited the strong AChE inhibition activity with an IC ₅₀ < 50 [91]. Score- 1	Score- 3	monoterpenes namely, α-Pinene, limonene and 1,8-cineole [90].	7
<i>Schotia brachypetala</i>	Roots and barks are used for nervous conditions [93]. Bark decoction is used for the treatment of heartburn and to calm down the effects of too much drinking [48]. The plant is traditionally used in the treatment of neurodegenerative diseases [25]. Score- 2	The DCM:MeOH (1:1) root extract of <i>S. brachypetala</i> represented the IC ₅₀ 0.89±0.01 and water extract represented IC ₅₀ value 3.40±0.50mg/ml for AChEI activity. The DCM:MeOH bark extract displayed IC ₅₀ 0.27±0.07 mg/ml and 0.49±0.04mg/ml for water extract. The presence of Stilbenes and phenolics have been indicated the plant to have good radical scavenging activity [25]. Score- 1	Low toxicity, hexane extract of leaf, Brine shrimp Cell line IC ₅₀ = 3300.0µg/ml [72]. Score- 1	3,3',4',5,5'-pentahydroxystilbene [94], Linolenic acid (9,12,15-octadecatrienoic) and methyl-5,11,14,17-eicosatetraenoate, two antibacterial fatty acids [95], Flavonolacyl glucosides [96].	4
<i>Stapelia gigantea</i>	Zulu healers use hot stem infusions to treat hysteria [24]. Score- 3	Not known Score- 3	Not known Score- 3	Steroidal glycosides [97].	9
<i>Tabernaemontana elegans</i>	Used for the treatment of heart diseases, cancer, tuberculosis, stomach ache,	Monoterpene indole alkaloid and β-carboline alkaloids isolated from <i>T. elegans</i> reported	High toxic, THP-1 Cell line IC ₅₀ < 4µg/ml [72].	Three β-carboline indole alkaloids and a monoterpene	

	infertility[98]. Treat various neurological disorders [99]. Score- 2	apoptosis induction activities in human hepatoma HuH-7 cells [100]. Score- 3		indole alkaloid [101].	8
<i>Tetradenia riparia</i>	Leaf infusions are taken for the treatment of respiratory ailments, mouth ulcers, fever, influenza, diarrhea, stomach ache, swollen legs and crushed leaves are used for headache [48]. Score- 3	The diterpene diol isolated from <i>T. riparia</i> exhibited significant antimicrobial activity against several bacteria and fungi with a bacterial activity of 12.5 µg/ml against <i>Bacillus subtilis</i> and <i>Streptococcus pyogenes</i> [102]. The ethanolic extract of <i>T. riparia</i> showed the AChE inhibition activity with IC50 value of 750 µg/ml [103]. Score- 1	Moderate toxic, Vero cell line, IC ₅₀ = Leaf extract, 124.455µg/ml; Stem extract, 186.2µg/ml; Root extract, 153.1µg/ml [104]. Score- 2	Tannins, Saponins, Terpenoids, Flavonoids, Cardiac glycosides, Phenols and Alkaloids [105]. A diterpene diol 8(14),15-Sandaracopimara diene-7α,18-diol [102]. Ibozol, 7α-hydroxyroyleanone, umuravumbolide, deacetylumuravumbolide, deacetylboronolide and 1',2'dideacetylboronolide [102]. 6	
<i>Trichilia dregeana</i>	Used as wound healing, for the treatment of dysentery, fever, stomach and intestinal complaints [106]. Traditionally it is used for the treatment of inflammatory diseases such as kidney pain, a sore back and backache,	Cycloart-23-ene-3,25-diol isolated from the ethyl acetate leaf extract showed a strong inhibitory activity of 80% against COX-2 at a concentration of 100µM with an IC50 value of 40µM. The compound also showed the weak inhibitory effect against acetylcholinesterase in an anti-	No toxicity, MTT cellular viability assay, Cellular viability (% cell growth at 100 µg/ml ± s.d.) Aqueous extract= 100 ± 0.78. Organic extract= 99±0.87 [109].	Limonoids from which a known crystalline compound rohituka-7 and the other compounds designated dregeana-1 to 4 were isolated from the seeds of <i>T. dregeana</i> [110]. A compound named cycloart-23-ene-3,25-diol was isolated from ethyl acetate leaf	

	bronchial inflammation, fever and rheumatism [107]. Score- 3	cholinesterase test at a concentration of 0.4 mM [107]. Stored plant material showed strong AChE inhibitory activity which confirm its therapeutic value towards Alzheimer's disease [108]. Score- 1		extract of <i>T. dregeana</i> [107].	5
<i>Xysmalobium undulatum</i>	Plant is used for headache and abdominal pains in Botswana [85]. Traditionally, it is also used for the treatment of diarrhoea, dysentery, stomach cramps, indigestion [48]. Score- 3	<i>X. undulatum</i> has been reported to exhibit acetylcholinesterase inhibitory activity (IC ₅₀ = 0.05–0.5 µg/ml) [111]. It has been reported to exhibit various <i>in vitro</i> activities such as antioxidant activity, antibacterial, antifungal, antiplasmodial, antidiarrhoeal and serotonin reuptake inhibitors (SSRI) antidepressant activity [112]. Score- 1	High toxic [33] Score- 3	Uzarin, xysmalorin and their isomers allouzarin and alloxysmalorin, allouzarigenin, alloxysmalogenin and other compounds ascleposide, coroglaucigenin, coroglaucigenin-3-O-glucoside, pachygenol, pachygenol-3β-O-glucoside, desglucouzarin, smalogenin, desglucoxysmalorin, uzaroside, pregnenolone and β-sitosterol [33].	7
<i>Zanthoxylum capense</i>	A species of <i>Zanthoxylum</i> is used for the treatment of mental illness in Gabon. It is also used as an epilepsy remedy among Europeans [24].	A compound named decarine isolated from <i>Z. capense</i> showed a strong antimycobacterial activity found, both <i>in vitro</i> and <i>ex vivo</i> mycobacteria [113].	Low toxic, THP-1 Cell line IC ₅₀ = 45.7 µg/ml [42], [72]	Benzophenanthridine alkaloid. 2-arylbenzofuran neolignans [113], [114].	

	Score- 2	Score- 3	Score- 1		6
<i>Ziziphus mucronata</i>	Decoctions are taken for pneumonia. It is also used for the treatment of the swellings of skin, wounds [115].	The roots of <i>Z. mucronata</i> showed good antioxidant and acetylcholinesterase inhibitory activity [99].	High toxic, Brine shrimp Cell line IC ₅₀ = 900µg/ml [57], [42].	Phenolic compounds, flavonoids and proanthocyanidins [116]. Two cyclopeptide alkaloids were isolated from the roots of <i>Z. mucronata</i> [117]. Mucronine D and sanjoinine [118], [3].	
	Score- 3	Score- 1	Score- 3		7

2.3.2. Collection and extraction of plant material

The selected 21 plants species and where possible their different plant parts (e.g. leaves, stems, stalks, flowers) were collected in the spring season during the months October to December 2015. Two plants species, *Commelina africana* and *Ziziphus mucronata* were collected from the Dinokeng Nature Reserve Pretoria; one species, *Xysmalobium undulatum* was collected from experimental farm of University of Pretoria and remaining 18 species were collected from Mainie *vanderSchijff* Botanical Garden, University of Pretoria. Voucher specimens were deposited and identified at the H.G.W.J. Schweickerdt Plant Herbarium at University of Pretoria. A total of 34 extracts were prepared for the different plants species and their plant parts after extraction with dichloromethane/methanol (DCM:MeOH) (1:1). The extraction yield of 34 plant extracts ranged from 3% to 25%, labelled 1 to 34. The variation in the extraction yields depended on the plants species and the plant part collected. The voucher specimen of the plants and extraction yields are shown in Table 2.2.

Table 2.2: Plant species selected and collected with voucher specimen numbers³

Sample No.	Family	Plant species	Plant part extracted	Voucher specimen	Extraction yield (calculated from dry plant material) as w/w%
1	Apiaceae	<i>Centella asiatica</i> (L.) Urb.	Leaves + Stalks	124298	13.9
2	Apiaceae	<i>Heteromorpha arborescens</i> (Spreng.) Cham. & Schltldl.	Leaves	124318	8.7
3	Apocynaceae	<i>Mondia whitei</i> (Hook.f.) Skeels	Leaves	124299	10.3
4			Stems		7.6
5	Apocynaceae	<i>Stapelia gigantea</i> N.E.Br.	Stems	124308	12.5
6			Seeds		23.3
7			Seed pods		3.1
8	Apocynaceae	<i>Tabernaemontana elegans</i> Stapf	Leaves	124300	13.0
9	Apocynaceae	<i>Xysmalobium undulatum</i> (L.) W.T.Aiton	Stalks + Flowers	124301	7.6
10	Araliaceae	<i>Cussonia spicata</i> Thunb.	Leaves	124302	20.9
11			Stems		6.0
12	Araliaceae	<i>Cussonia paniculate</i> Eckl. & Zeyh.	Leaves	124309	5.7
13			Stems		7.7
14	Asphodelaceae	<i>Bulbine natalensis</i> Baker	Leaves	124310	24.5
15	Celastraceae	<i>Catha edulis</i> (Vahl) Endl.	Leaves	124303	13
16			Stems		6.9
17	Commelinaceae	<i>Commelina africana</i> L.	Leaves	124311	11.2
18	Crassulaceae	<i>Cotyledon orbiculata</i> L.	Leaves	124312	2.2
19			Stems		6.1
20	Fabaceae	<i>Schotia brachypetala</i> Sond.	Leaves	124313	9.2
21	Lamiaceae	<i>Tetradenia riparia</i> (Hochst.) Codd	Leaves	124304	12.5
22			Stems		5.4

³ All plant species name have been confirmed as accepted on <http://www.theplantlist.org>.

23	Maliaceae	<i>Trichilia dregeana</i> Sond.	Leaves	124305	9.9
24			Stems		6.8
25	Plumbaginaceae	<i>Plumbago auriculata</i> Lam.	Leaves	124306	19.3
26			Stems		7.1
27	Rhamnaceae	<i>Ziziphus mucronata</i> Willd.	Leaves + Stalks	124314	10.5
28	Rutaceae	<i>Ruta graveolens</i> L.	Leaves	124315	22.8
29			Stems		10.5
30	Rutaceae	<i>Zanthoxylum capense</i> (Thunb.) Harv.	Leaves	124316	7.2
31			Stems + Thorns		9.1
32	Scrophulariaceae	<i>Buddleja salviifolia</i> (L.) Lam.	Leaves	124307	10.5
33			Stems		6.0
34	Vitaceae	<i>Cissus quadrangularis</i> L.	Stems	124317	9.6

2.3.3. The effect of the plant extracts on A β 42 production

The bioassaying was made possible with the support of Korean institute of Science and Technology (KIST). Special thanks to the strong collaboration of Dr Yoon Sun Chun, Ansun Park in Dr Hyun OK Yang's laboratory at KIST. I also acknowledge the work done by KIST in the training on running the assays and interpreting of data during my research visit to their laboratories. Cells were incubated with 50 μ g/ml extracts for 8 h, and the level of A β 42 was measured from the conditioned media by using specific ELISA method. Galanthamine, which was reported to inhibit amyloid aggregation [119] and release of A β 42 was decreased by 37% with 10 μ M galanthamine in SH-SY5Y cells [120] was used as a positive control (Figure 1). The A β 42 production of galanthamine was measured by treating cells with varying concentrations for 8 h. However, galanthamine at a high concentration of 10 μ M did not significantly affect the A β 42 production under the experimental conditions, thus, APPsw transfected HeLa cells with same concentrated DMSO (vehicle) was used for negative control.

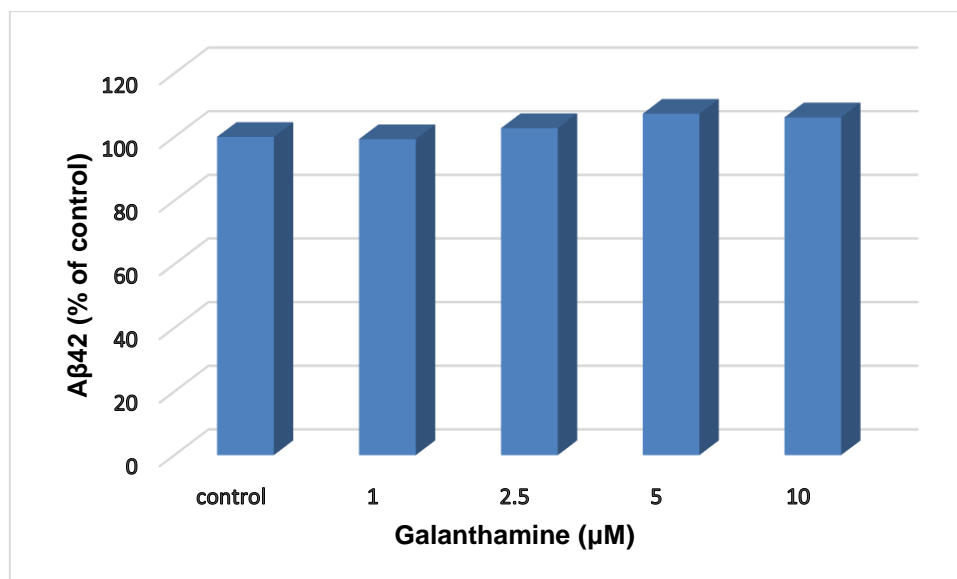


Figure 2.4: Galantamine did not change the level of Aβ42. APPsw-transfected HeLa cells were incubated with indicated concentrations of galantamine for 8 h. The level of Aβ42 was measured from the conditioned media by using specific ELISA methods. The level of Aβ42 was not changed by galantamine (n = 6)

The 34 plant extracts were screened to determine their inhibitory properties for Aβ42 production in HeLa cells stably transfected with APPsw. The results showed that 11 extracts (32.3%) (Extract numbers: 1, 2, 3, 4, 9, 12, 15, 17, 18, 20 and 32) significantly decreased the Aβ42 level as compared to the negative control as shown in Figure 2.5 to Figure 2.8.

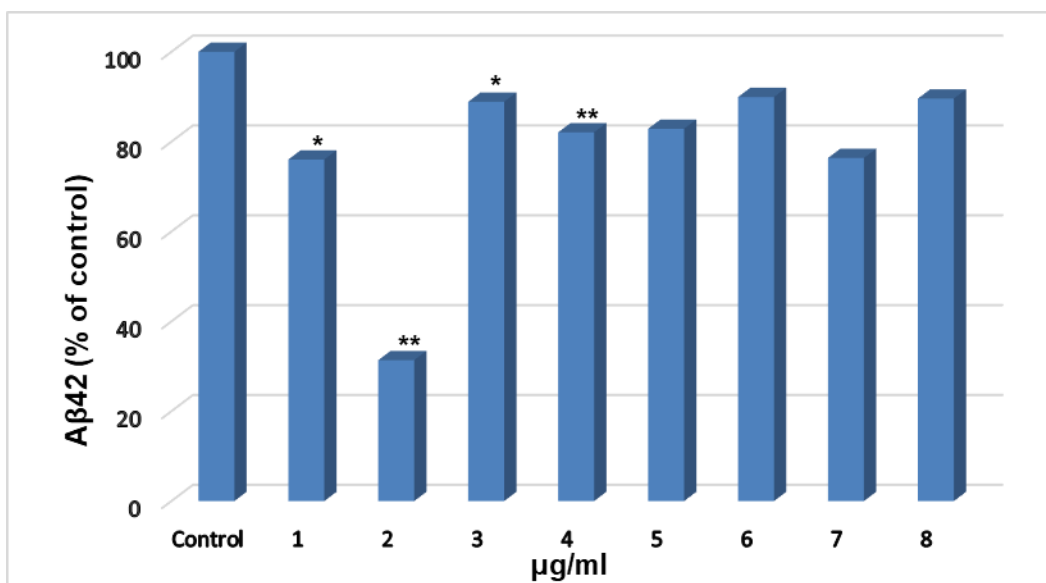


Figure 2.5: Effect of plant extracts on the level of Aβ42 in APPsw-transfected HeLa cells. APPsw-transfected HeLa cells were incubated with 50 µg/ml extracts for 8 h, and the level of Aβ42 was measured from the conditioned media by using specific ELISA methods. *, P<0.05; **, P<0.01. *Centella asiatica* (1), *Heteromorpha arborescens* (2), *Mondia whitei* (3 and 4), *Stapelia gigantea* (5, 6 and 7), *Tabernaemontana elegans* (8)

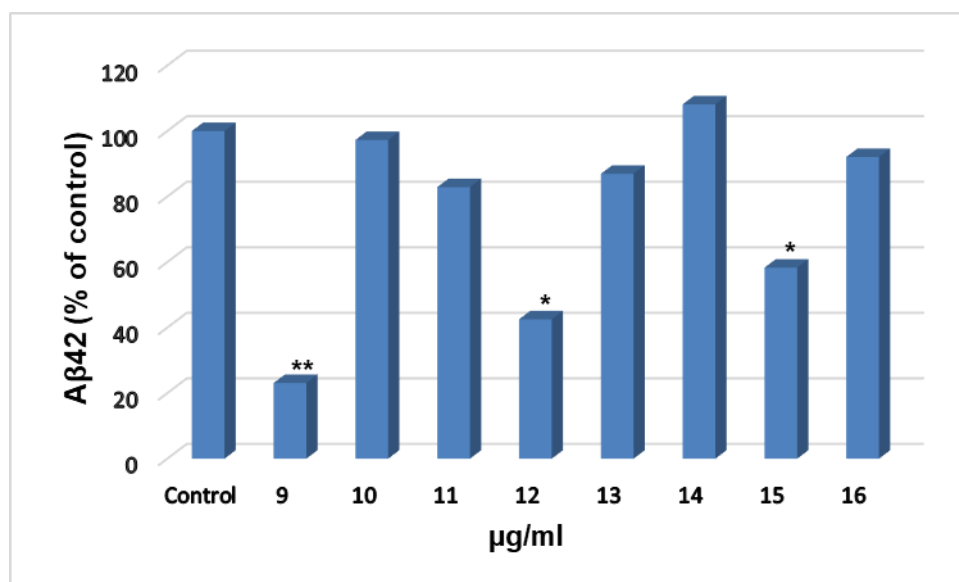


Figure 2.6: Effect of plant extracts on the level of Aβ42 in APPsw-transfected HeLa cells. APPsw-transfected HeLa cells were incubated with 50 µg/ml extracts for 8 h, and the level of Aβ42 was measured from the conditioned media by using specific ELISA methods. *, P<0.05; **, P<0.01. *Xysmalobium undulatum* (9), *Cussonia spicata* (10 and 11), *Cussonia paniculata* (12 and 13), *Bulbine natalensis* (14), *Catha edulis* (15 and 16)

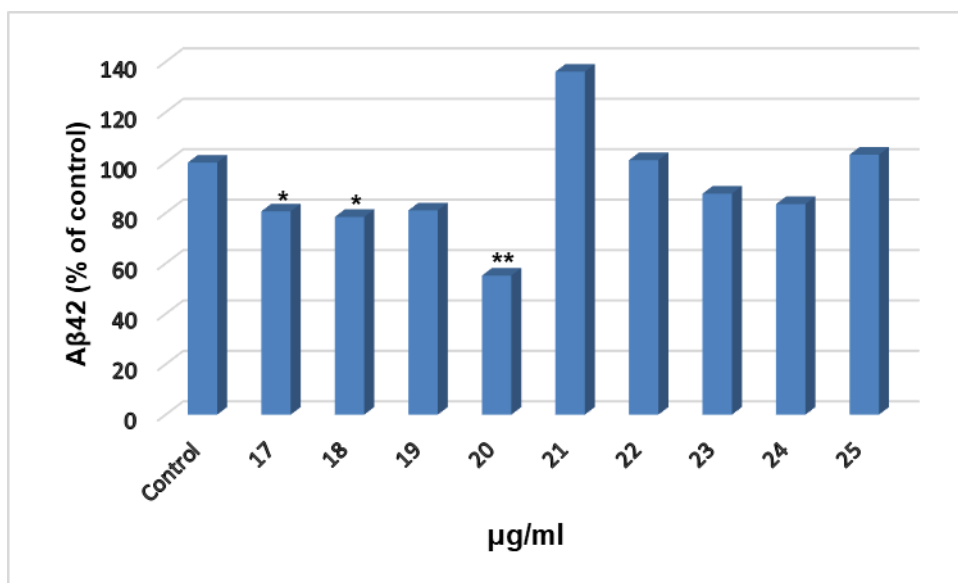


Figure 2.7: Effect of plant extracts on the level of Aβ42 in APPsw-transfected HeLa cells. APPsw-transfected HeLa cells were incubated with 50 μg/ml extracts for 8 h, and the level of Aβ42 was measured from the conditioned media by using specific ELISA methods. *, P<0.05; **, P<0.01. *Comelina africana* (17), *Cotyledon orbiculata* (18 and 19), *Schotia brachypetala* (20), *Tetradenia riperia* (21 and 22), *Trichilia dregeana* (23 and 24), *Plumbago auriculata* (25)

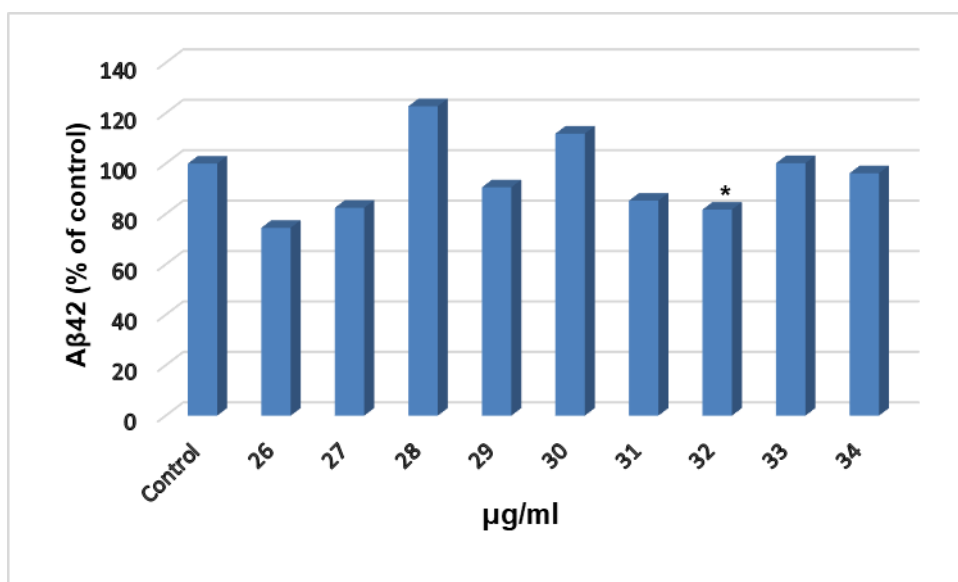


Figure 2.8: Effect of plant extracts on the level of Aβ42 in APPsw-transfected HeLa cells. APPsw-transfected HeLa cells were incubated with 50 μg/ml extracts for 8 h, and the level of Aβ42 was measured from the conditioned media by using specific ELISA methods. *, P<0.05; **, P<0.01. *Plumbago auriculata* (26), *Ziziphus mucronata* (27), *Ruta graveolens* (28 and 29), *Zanthoxylum capense* (30 and 31), *Buddleja salviifolia* (32 and 33), *Cissus quadrangularis* (34)

Among them, the extracts of stems of *Heteromorpha arborescens* (No. 2), leaves of *Xysmalobium undulatum* (No. 9), leaves of *Cussonia paniculata* (No. 12) and leaves of *Schotia brachypetala* (No. 20) potently reduced the secreted level of A β 42 compared to the control by 68.63 ± 0.2 , $76.9 \pm 1\%$, $57.5 \pm 1.3\%$ and $44.8 \pm 0.1\%$, respectively. These four extracts were selected for further studies due to their ability to inhibit A β 42 production by 44% and more and their limited toxicity at the same test concentrations. The cytotoxicity of these plants is further discussed in Chapter 3 and Chapter 4.

H. arborescens is traditionally used for the treatment of nervous disorders [121], [74]. *X. undulatum* has been reported to exhibit antidepressant, acetylcholinesterase inhibition activity and Serotonin reuptake inhibitory activity [33], [112], [111]. *C. paniculata* is also used traditionally for the treatment of nervous disorders [29]. In a previous study, phenolic compounds isolated from *S. brachypetala* were reported to inhibit A β 42 production and proved its efficacy to be the anti-Alzheimer's drug in future [32]. Either the traditional uses or the reported bioassays of these plants also supported their efficacy for the reduction of A β 42 in the present study.

The active plant extracts were also compared with their scoring system used for the selection of the plants. It was found that, of these four plants only *S. brachypetala* showed very good correlation with the scoring system. It was given a good score **4**, because of its traditional uses related to nervous disorders and a well-researched studies on A β 42 bioassays [32]. While *M. whitei* with the similar score showed a very low activity towards the reduction of A β 42, as the levels of A β 42 was reduced by only 11.1 ± 0.1 and 17.9 ± 0.6 with *M. whitei* leaf and *M. whitei* stem extract, respectively. Some other plant extracts, *C. asiatica*, *C. edulis*, *C. africana*, *C. orbiculata* and *B. salvifolia* also showed some reduction in A β 42 by 23.9 ± 1.5 , 41.7 ± 1.9 , 19.3 ± 0.5 , 21.5 ± 0.5 and 18.2 ± 0.8 , respectively however all these extracts showed low percentage for A β 42 reduction. Due to the low A β 42 reduction, these plant extracts were not selected for further studies. The scoring system helped to select the plants

for testing against A β 42 reduction, but all the selected plant extracts did not show significant reduction towards A β 42.

2.4. Conclusion

At least 52 plant species were selected using different database searches. Due to the broad search, an additional criterion and a scoring system was used. With the help of scoring system, 21 plant species which were selected for the further experiments including collection, extraction and screening for A β 42 reduction. A total of 34 extracts were prepared from different plant parts of the 21 plants. Among them, four plant extracts, stems of *H. arborescens*, leaves of *X. undulatum*, leaves of *C. paniculata* and leaves of *S. brachypetala*, potentially decreased the levels of A β 42. The stem extract of *H. arborescens* and leaf extracts of *X. undulatum* and *C. paniculata* were screened first time for A β 42 reduction. These extracts were scientifically shown to exhibit A β 42 inhibition activities thus providing some scientific evidence authenticating their traditional uses and also exhibiting their potential to be further developed into anti-Alzheimer's ingredients. The *S. brachypetala* showed its ability for the reduction of A β 42 but in other assays from the present study. *H. arborescens* and *C. paniculata* were found to be traditionally used for the treatment of mental disorders while *X. undulatum* has been previously reported for acetylcholinesterase inhibition. The traditional uses and preliminary bioassays of these plant extracts also showed their efficacy towards the treatment of mental disorders. These four extracts were selected for further study to identify and isolate the active ingredients responsible for the inhibition of A β 42 activity.

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Chapter 3

Chemical profiling of extracts of *Schotia brachypetala*, *Cussonia paniculata* and *Heteromorpha arborescens* using UPLC-QTOF-MS for the purpose of identifying the active compounds

3.1. *Schotia brachypetala*

3.1.1. Botany and geographical distribution

S. brachypetala belongs to family Fabaceae or Leguminosae and is a tree endemic to South Africa. It is an attractive, medium to large deciduous tree with a widely spreading, densely branched round canopy up to 10-16 m tall and 10-15 m wide and occasionally reach up to 22 m. It has a single trunk with branches low down. Leaves are bright red in spring and mature to dark glossy green. The flowers are deep red in color and bloom in spring from August to November. The flowers produce nectar and can also be chewed. The fruit is small, hard and woody pod. It gets split on the tree and releases the seed from inside. The seeds look like domestic broadbean and is the reason why the plant is named as boer-bean while the seeds are edible after roasting. It is commonly known as weeping boer-bean, African walnut in English; huilboerboon in Afrikaans; umfofofo in Xhosa; uvovovo in Zulu and mununzwa, mutanswa in Venda [1]. The genus *Schotia* is mainly confined to southern Africa, ranging from southern Zimbabwe and Mozambique to southern South Africa and Namibia. *S. brachypetala* is widely distributed in the Eastern Cape, through KwaZulu-Natal, Swaziland, Mpumalanga, Northern Province and into Mozambique and Zimbabwe. It is found mainly in the sub-tropical regions of the country, growing in warm, bushveld, scrub forests and deciduous woodlands, often found on the banks of rivers [2], [3], [4].



(A)



(B)

Figure 3.1: Picture (A) showing *S. brachypetala* tree and (B) shows the *S. brachypetala* leaves (image taken by Anuradha Thakur at the time of collection).

3.1.2. Traditional medicinal uses

The decoction of barks and sometimes the mixture with roots is used for the treatment of heartburn and hangovers, flu symptoms, nervous disorders, heart conditions and diarrhoea. The bark and the root mixtures are also used as a blood purifier and to strengthen the body. The roots and bark are used in South Africa as a traditional remedy for dysentery and diarrhoea [5]. The mixture is also used for the facial saunas. The plant is traditionally used in the treatment of neurodegenerative diseases [6]. Traditionally, it is used to wash body, steam the face as a remedy for pimples and to reduce body swelling. It is also used for the treatment of tropical ulcers and wounds [7]. The seeds are edible and can be eaten after roasting [8].

3.1.3 Phytochemistry

The phytochemistry of *S. brachypetala* is well studied. A 3,3',4',5,5'-pentahydroxystilbene has been isolated in high yields (7.2% on wood weight) from the methanolic extract of the dark red heartwood of *S. brachypetala*. This compound was for the first time obtained as a mixture with 3,3',4',5 tetrahydroxystilbene from the wood of *Vouacapoua macropetala* which was further isolated as the pure pentaacetate and hydrolysed to give free phenol [9]. Stilbene based compounds are known for their remarkable pharmacological activities such as for the treatment of hyperlipidemia, arteriosclerosis and inflammatory diseases [10]. In nature the stilbenes are classified as phytoalexins, a small part of phenylpropanoids, the secondary metabolites synthesized by plants to protect themselves from biotic and abiotic stress. Due to the great potential of stilbene based hybrid compounds in the field of cancer, Alzheimer's and other relevant diseases, these compounds have become a great research interest for the chemists and biologists [11]. Two antibacterial fatty acids named linolenic acid (9,12,15-octadecatrienoic) and methyl-5,11,14,17-eicosatetraenoate were isolated from the ethanol extracts *S. brachypetala* leaves [5]. Flavonol acyl glucosides were isolated from the aerial parts of the plant which exhibit antioxidant, antibacterial and antimalarial activity [8]. Seven quercetin glucoside derivatives, 3-O-methylquercetin 7-O- β -glucopyranoside, 3,4'-di-O-methylquercetin 7-O- β -glucopyranoside, 3-O-methylquercetin 7-O-[β -D-6''(E-p-coumaroyl)glucopyranoside], 3,4'-di-O-methylquercetin 7-O-[β -D-6''(E-p-coumaroyl)glucopyranoside], quercetin 7-O- β -glucopyranoside (E), (2R, 3R)-dihydroquercetin 7-O- β -glucopyranoside and quercetin 3-O-[2-O- β -xylopyranosyl-6-O- α -rhamnopyranosyl]- β -glucopyranoside were isolated from the methanol extract of areal parts of *S. brachypetala* [12]. From the *S. brachypetala* leaf extract, three phenolic compounds gallic acid, myricetin-3-O- α -L-¹C₄-rhamnoside and quercetin-3-O-L-¹C₄-rhamnoside were isolated and identified [13]. Ten other phenolic compounds; daidzein, naringin, procyanidin isomers, procyanidin dimer gallate, quercetin 3-O-rhamnoside, quercetin 3-O-glucuronide, quercetin hexose gallic acid,

quercetin hexose protocatechuic acid, and ellagic acid were isolated from the aqueous alcohol extract of stalks of *S. brachypetala* [14].

3.1.4. Previously reported biological assaying

The wide traditional uses of different parts of *S. brachypetala* by traditional health practitioners have led researchers to seek and explore the opportunities in drug development from this plant. Two antibacterial fatty acids, 9,12,15-octadecatrienoic (linolenic) acid and methyl-5,11,14,17-eicosatetraenoate isolated from the activity-directed fractionation of ethanol extracts of the dry leaves showed strong antibacterial activity against two gram-positive bacteria, *Bacillus subtilis* and *Vibrio parahaemolyticus*, while to a lesser extent, against two gram-negative species, *Escherichia coli* and *Klebsiella pneumoniae* [5]. The presence of stilbenes and phenolics have indicated the plant to have good radical scavenging activity [6]. The *S. brachypetala* bark has been reported to exhibit anti-hyaluronidase activity with an IC_{50} value of $75.13 \pm 7.49\%$ which demonstrated its efficacy for anti-aging properties [7]. *S. brachypetala* (root and bark) were found to exhibit antioxidant activities. Antioxidant activity was measured by the DPPH and ABTS radical scavenging tests. The DCM:MeOH extract of *S. brachypetala* roots showed an IC_{50} value of 0.05 ± 0.02 mg/ml while the water extract showed an IC_{50} value of 0.05 ± 0.02 mg/ml measured by DPPH test. For ABTS test, the DCM:MeOH root extract showed an IC_{50} value of $3.26 \times 10^{-7} \pm 0.21 \times 10^{-9}$ mg/ml and the water extract showed an IC_{50} value of $3.7 \times 10^{-7} \pm 0.21 \times 10^{-9}$ mg/ml. The bark of *S. brachypetala* was also tested for DPPH and ABTS scavenging bioassays. It showed an IC_{50} value of 1.90 ± 0.50 mg/ml for the DCM:MeOH extract and an IC_{50} value of 0.13 ± 0.03 mg/ml for the water extract in DPPH test. For ABTS test, and IC_{50} was not determined for the DCM:MeOH extract and for water extract it showed an IC_{50} value of 0.15 ± 0.03 mg/ml. The organic root extract of *S. brachypetala* showed lowest IC_{50} values in both antioxidant assays indicating its good antioxidant potential [6].

3.1.5. Bioassaying of *S. brachyptala* as a potential for Alzheimer's disease

The plant extract was tested for acetylcholinesterase inhibition (AChEI) activity and compared to the positive control galanthamine [15]. The DCM:MeOH root extract of *S. brachyptala* gave an IC₅₀ value of 0.89 ± 0.01 mg/ml and the water extract had an IC₅₀ value of 3.40 ± 0.50 mg/ml for AChEI activity. The DCM:MeOH bark extract displayed an IC₅₀ 0.27 ± 0.07 mg/ml and 0.49 ± 0.04 mg/ml for the water extract [6]. *S. brachyptala* exhibited good monoamine oxidase B (MAO-B) inhibition activity. Recently, MAO-B inhibitors are also taken in consideration for the treatment of neurodegenerative diseases such as Parkinson's disease and Alzheimer's disease [16]. These bioassays provide evidence of the efficacy of *S. brachyptala* and substantiates further research into the potential of the plant as anti-Alzheimer's drugs.

3.2. *Cussonia paniculata*

3.2.1. Botany and geographical distribution

C. paniculata belongs to the family Araliaceae, commonly known as mountain cabbage tree. It is a short, slow growing tree or a large shrub reaching up to 5 m in height. It turns into an attractive plant in its juvenile stage and become an asset to small gardens. One of the most distinctive features of this plant is cabbage blue leaves which are arranged in whorl pattern. It is considered as succulent pachycaul because of its thick, squat and swollen stem. The roots are also thick and swollen. The plant bears yellowish to greenish colored flowers which blooms in January to April. It is an indigenous plant species which is well distributed in South Africa and very common in KwaZulu-Natal. It is often found in the mountains of Karoo and Eastern Cape through KwaZulu-Natal [17], [18], [19].



Figure 3.2: Picture showing *C. paniculata* tree (image taken by Anuradha Thakur at the time of collection).

3.2.2. Traditional medicinal uses

It is reportedly used as a traditional medicine against pain, inflammation, infections and malaria [17]. The species is also used for the treatment of mental illness, diarrhea and sexually transmitted diseases [20]. The thick tuberous roots are peeled and eaten raw as an emergency food or as a source of water. In South Africa, the mixture of *Buddleja* species with *Heteromorpha trifoliata* and *C. paniculata* is used for the treatment of early nervous and mental disorders [6].

3.2.3. Phytochemistry

Triterpenoid glycosides, 3-O- β -D-glucopyranoside of β -sitosterol, 3-O- α -L-arabinopyranosides of oleanolic acids, ursolic acids, hederagenin, 3-O- β -D-glucopyranosyl-(1 \rightarrow 2)- α -L-arabinopyranoside of oleanolic acid, 3-O- α -L-arabinopyranosyl-28-O- α -L-rhamnopyranosyl-(1 \rightarrow 4)-O- β -D-glucopyranosyl-(1 \rightarrow 6)-O- β -D-glucopyranosides of oleanolic and ursolic acids and 28-O- α -L-rhamnopyranosyl-(1 \rightarrow 4)-O- β -D-glucopyranosyl-(1 \rightarrow 6)-O- β -D-glucopyranoside of 23-hydroxyursolic acid were isolated from *C. paniculata* leaves [21]. Previous studies have shown that the plant contains high sterol/triterpene, flavonoids such as rutin, quercetin and luteolin [22].

3.2.4. Previously reported biological assaying

The stem bark of *C. paniculata* exhibited anti-inflammatory and analgesic activities. In a carrageenan-induced rat paw oedema experimental animal model, 50 mg/kg aqueous extract of *C. paniculata* bark showed potent anti-inflammatory properties at 2 h with percentage inhibition of 0.44 ± 3.1 (96.6) % as compared to 10 mg/kg indomethacin, 3.4 ± 2.1 (73.7) %. The analgesic effect was also recorded for the same extract by conducting a formalin test for 30 min. The extract at 50 mg/kg showed a significant licking time, $5.8 \pm 0.8^*$ sec and licking frequency $8.3 \pm 0.3^*/30$ min as compared to indomethacin which exhibited licking time $5.0 \pm 0.5^*$ sec and $14.3 \pm 1.8^*/30$ min (* = $P < 0.05$) [22]. The DCM:MeOH leaf extract of *C. paniculata* was screened for its *in vitro* anticancer activity and showed an inhibition of 100% cell growth at a concentration of 5.25 μ g/ml (renalTK10), 15.0 μ g/ml (melanoma UACC62) and 55.69 μ g/ml (breast MCF7) [23]. In another study different plant part extracts (leaves, bark, stem and roots) of *C. paniculata* using different solvents (water, ethanol and ethyl acetate) were screened for anti-inflammatory activity in the COX-1 assay as compared to control indomethacin. Indomethacin showed 78% inhibition at a concentration of 5 μ M. Ethanol and ethyl acetate extracts of *C. paniculata* were more active than water extracts as shown in Table 3.1 [24].

Table 3.1: Inhibition percentage of cyclooxygenase in COX-1 bioassay by different extracts of *C. paniculata*

<i>C. paniculata</i> (Plant part analysed)	% Inhibition at 0.2µg/µl		
	Water extract	Ethanol extract	Ethyl acetate extract
Leaf	29	80	58
Bark	0	42	78
Stem	27	85	80
Root	32	15	38

3.2.5. Bioassaying of *C. paniculata* as a potential for Alzheimer’s disease

Although traditionally the plant is used for the treatment of diseases associated with nervous system, there is no any scientific study to prove its activity for the reduction of Aβ42 and inhibition of acetylcholinesterase inhibition which are targets for treatment and management of Alzheimer’s disease [24].

3.1. *Heteromorpha arborescens*

3.3.1. Botany and geographical distribution

H. arborescens also known as *H. trifoliata* and belongs to the family Apiaceae. It is a medium-sized woody shrub or small tree approximately 15 metres high. The bark is papery, scale-like and is easily peels off. The leaves are green to grey green in colour and variable in shape and size. The leaves turn yellow and red in autumn. The flowers are small, green or yellow in colour with a strong smell and bloom in December and January. Fruits are flattened, heart shaped and pale brown. [2]. The plant is commonly known as “Parsley tree” in English, “wildepitersielie” in Afrikaans and “umBangandlala” in Xhosa. The plant is widely distributed in South Africa throughout the regions Gauteng, Free State, Mpumalanga, KwaZulu-Natal,

Midlands, Limpopo, North West. It is found in wooded grassland, bushveld and on forest margins [25], [2].



(A)



(B)

Figure 3.3: Picture (A) showing *H. arborescens* stem and leaves and (B) shows *H. arborescens* leaves (images taken by Anuradha Thakur at the time of collection).

3.3.2. Traditional medicinal uses

H. arborescens is one of the important medicinal plants of South Africa. It is used to treat abdominal pains, colic, fever, headaches, asthma, coughs, infertility, dysmenorrhea, dysentery, nervous and mental disorders as well as a vermicide in children [25]. In Eastern Cape province, South Africa, people take boiled leaves and roots of *H. arborescens* orally for the treatment of diabetes. It is also used traditionally by ethnic groups in central and southern Africa as a remedy for headache, fever and pain [26]. The other major medicinal uses of *H. arborescens*

include the treatment of respiratory problems, shortness of breath, intestinal worms, snakebite, skin infections, an aphrodisiac and malaria [27], [28].

3.3.3. Phytochemistry

Falcarindiol, a polyacetylene was isolated from the roots of the plant while the other compounds, germacrene D, sabinene and α -pinene were isolated from leaves of *H. arborescens* [29]. Two saikosaponins, 16 β ,23-dihydroxy-13,28-epoxyolean-11-en-3 β -yl- $[\beta$ -D-glucopyranosyl (1 \rightarrow 2)]- $[\beta$ -D-glucopyranosyl (1 \rightarrow 3)]- β -D-fucopyranoside named as saikogenin B₄ and 16 β ,23,28- trihydroxy-11 α -methoxyolean-12-en-3 β -yl- $[\beta$ -D-glucopyranosyl (1 \rightarrow 2)]- $[\beta$ -D-glucopyranosyl (1 \rightarrow 3)]- β -D-fucopyranoside named saikogenin F were identified from the leaves of *H. arborescens* [30].

3.3.4. Previously reported biological assying

H. arborescens showed antimicrobial, anti-inflammatory (COX-1) and anthelmintic activities [31]. The leaf and stem methanol extracts of *H. arborescens* has been reported to possess antibacterial activity against gram positive bacteria such as *Staphylococcus aureus*, *Bacillus subtilis*, *Micrococcus kristinae* and *Streptococcus faecalis*. The plant extract was found to exhibit antifungal property against *Penicillium notatum*, *Aspergillus flavas* and *Aspergillus niger* with an IC₅₀ value ranging from 0.07 to 1.07 mg/ml. [32]. In an *in vitro* α -glucosidase inhibition study, the ethyl acetate leaf extract and methanol bark extract of *H. arborescens* showed 50% α -glucosidase inhibition at concentrations of 3.12 \pm 0.62 and 3.10 \pm 1.73 mg/ml, respectively [26]. *H. arborescens* extracts also showed *in vitro* anti-arthritic activity with IC₅₀ values of 53.78 μ g/ml as compared to the positive control diclofenac sodium which had an IC₅₀ value of 32.37 μ g/ml [33]. Falcarindiol isolated from the roots of *H. arborescens* has been reported to exhibit antifungal, antibacterial and analgesic activity [2]. It has also been found to inhibit COX-1 [29]. The saikosaponins identified from *H. arborescens* leaf extract were reported to have inhibitory action on hepatitis [30]. The methanol leaf extract and ethanol

stem bark extract were observed to possess anti-inflammatory activities when tested against tetradecanoylphorbol acetate (TPA)-induced mouse ear edema with a percentage reduction in edema of 64% and 77%, respectively [34]. The ethanol and methanol extracts of the root bark extract of *H. arborescens* was reported to have antiviral activities against poliovirus at 10 µg/ml to 50 µg/ml concentration range while the DCM extract was active against the rhinovirus at a concentration range of 25 µg/ml to 50 µg/ml [35].

3.3.5. Bioassaying of *H. arborescens* for its potential against Alzheimer's disease

Although the plant is traditionally used for the treatment of nervous disorders, there are no previous studies reported on any bioassaying of *H. arborescens* for the reduction of A β 42 and acetylcholinesterase inhibition or any other mental disorders.

3.4. Materials and methods

3.4.1 Collection and extraction of plant material

S. brachypetala leaves, *C. paniculata* leaves and *H. arborescens* stems were collected from Maine vander Schijff Botanical Garden, University of Pretoria, South Africa with the help of Curator Jason Samuels. Plant identification was established at the H.G.W.J. Schweickerdt Plant Herbarium at University of Pretoria where voucher specimen were kept for further references. The collected leaves and stems were oven dried at 60°C and the dried plant material was ground to a fine coarse powder. The powdered leaves were extracted repeatedly (3 times) for 8 hours with DCM/MeOH (1:1) at room temperature using a magnetic stirrer. The solvents obtained after filtration of each extraction and were combined and concentrated using a rotary vacuum evaporator at 50-60°C and then dried under high vacuum for 24 hours. After drying, the extract was stored in the cold room.

3.4.2. Chemical profiling using UPLC-QTOF-MS

The crude extracts were analyzed using a Waters Acquity UPLC system (Waters corp., MAUSA), equipped with a binary solvent delivery system and an autosampler. The sample (1mg/ml) was prepared by dissolving dried extract, fraction, or pure compounds in 100% methanol. The solution was centrifuged at 10 000 g for 10 minutes to remove the particulates. The instrument was centrally operated by MassLynx 4.1 software (Waters Inc., Milford, Massachusetts, USA) for data acquisition. The Separation was achieved on Waters BEH C18 1.7 μm particle size (2.1 mm \times 100 mm). The mobile phase consisted of solvent A: water with 0.1% formic acid and solvent B: methanol with 0.1% formic acid). The gradient elution was optimized as follows: 3% B (0-0.1 min), 100% B (0.3-14.00 min), 100% B (14.00-16.00 min), 3% B (16.00-16.50 min), 3% B (16.50-20.00 min). The flow rate was 0.3 ml/min for the entire run, giving a total run time of 20 minute and the injection volume was 5 μl . The instrument was calibrated by direct infusion of 5 nM sodium formate solution at a flow rate of 10 $\mu\text{l}/\text{min}$ over a mass range of 50-1200 Da. The following MS source parameter were set for both positive and negative mode: Source temperature 100 $^{\circ}\text{C}$, sampling cone 15 V, extraction cone 4.0 V, desolvation temperature 400 $^{\circ}\text{C}$, cone gas flow 10.0 L/h, desolvation gas flow 700 L/h, capillary 2.0 kV. Both positive and negative modes were obtained, but the results were analyzed from the negative mode as the higher intensity peaks were obtained using this mode. Compounds were tentatively identified based on their accurate mass generated from MassLynx V 4.1, iFit values and by comparison of MS/MS fragmentation pattern with that of matching compounds from various search databases Reaxys, Scifinder, Scopus, Metlin, Metfusion, Pubchem, Chempider, Chemical Entities of Biological Interest (ChEBI) and Massbank. The iFit value is an indication of how well the measured isotopic ion ratio compares to the theoretical ion ratio [36]. The acquired accurate masses were also compared with those of known compounds in compound databases.

Compounds were also identified using a novel approach, integrated MassLynx and UNIFI (Waters) informatics software. Raw data was imported to UNIFI software to identify the

compounds which met the match criteria with scientific libraries incorporated in UNIFY software. The UNIFI data analysis automatically matched the compound information with the Chinese traditional medicinal library (Waters UNIFI Scientific Library Component Listing version: 01) and gave potential identifications. The traditional medicinal library contains more than 6000 natural compounds. The compounds with less than 5 mDa were shortlisted for further identification. Each shortlisted compound was provided with the detailed information including compound name, molecular formula, average molecular mass, monoisotopic retention time, exact mass error in mDa and ionization mode [37]. i-Fit confidence % of compounds with good match identified from UNIFI were compared i-Fit confidence % of MassLynx and thus confirmed a positive compound identification.

3.4.3. Cell culture

For cell culture methodology, see section 2.2.3 of chapter 2.

3.4.4. sAPP α , sAPP β , A β peptide assay

APP_{sw}-transfected HeLa cells at 80% confluence in a 35 mm dish were cultured with samples solubilized in dimethyl sulphoxide (DMSO) for 8 h in serum free medium. The conditioned medium was analyzed by a sandwich ELISA (Invitrogen) for detection of A β ₄₂ and A β ₄₀ according to the supplier's instruction. The levels of sAPP α and sAPP β -sw were measured from the conditioned medium using the specific ELISA (IBL). The APP_{sw}-transfected HeLa cells with DMSO (vehicle) as a negative control.

3.4.5. Cell viability measurement

Cell viability was measured by using an EZ-Cytox kit (Daeil Lab Co., Ltd., Republic of Korea) according to the manufacturer's instructions. APP_{sw}-transfected HeLa cells in a 96 well plate were cultured with sample solubilized in DMSO for 8 h, and then incubated with EZ-Cytox solution for 1 h at 37 °C. The absorbance was detected at 450 nm using a microplate reader.

3.4.6. Statistical analysis

For statistical analysis, see section 2.2.5 of chapter 2.

3.5. Results and discussion

3.5.1. Collection and extraction of plant material

S. brachypetala leaves, *C. paniculata* leaves and *H. arborescens* stems (50 g each) were collected from the Mainie vander Schijff Botanical Garden, University of Pretoria. Voucher specimen (PRU 124313) for *S. brachypetala*, (PRU 124309) for *C. paniculata* and (PRU 124318) for *H. arborescens* were deposited at the H.G.W.J. Schweickerdt Herbarium. After drying in oven at 60°C, 20 g powdered material of each plant was extracted with DCM:MeOH (1:1) solvent system. Three times repeated extraction using DCM:MeOH gave 9.2g of *S. brachypetala* leaf extract, 5.7g of *C. paniculata* leaf extract and 3.2 g of *H. arborescens* stem extract.

3.5.2. The effect of *S. brachypetala* on APP processing and cytotoxicity

The dose-dependent effect of *S. brachypetala* leaf extract on A β 42 production at 10, 25, 50, and 100 μ g/ml extract for 8 h, showed significant decrease in A β 42 production only at higher concentrations i.e. 29.69 \pm 4% at 50 μ g/ml and by 62.17 \pm 7% at 100 μ g/ml, while the lower concentrations of the leaf extract did not show any significant reduction in A β 42 i.e. 10 μ g/ml, A β 42 levels were reduced by 6.4 \pm 3.2% and by 12.37 \pm 4.3% at 25 μ g/ml. The decrease in secreted levels of A β 42 by *S. brachypetala* leaf extract is shown in Figure 3.4.

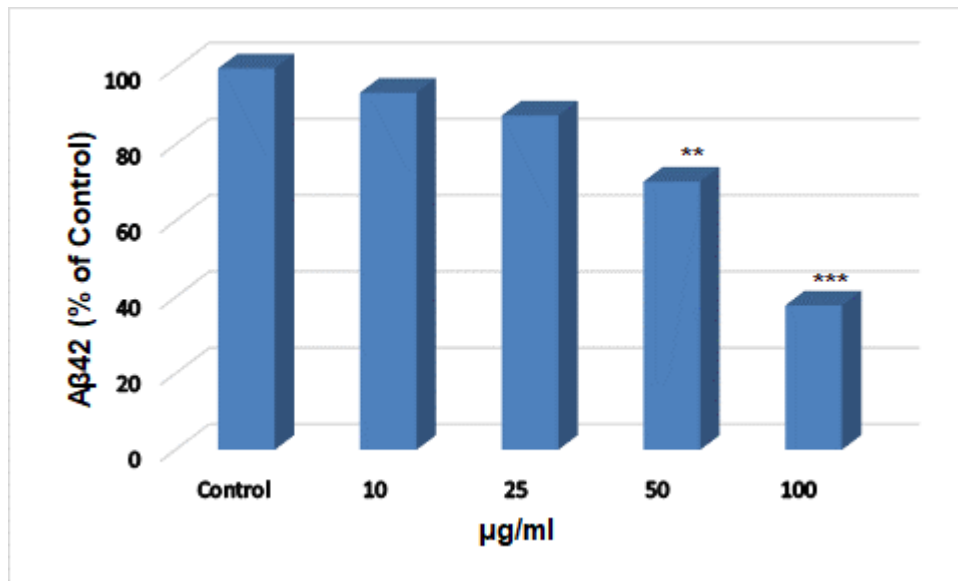


Figure 3.4: Change in the levels of Aβ42. Cells were incubated with indicated concentrations of leaf extract of *S. brachypetala* for 8 h, the levels of Aβ42 were measured from the conditioned media by using specific ELISA methods. The levels of Aβ42 (n = 4) were decreased by the extract at higher concentrations. **, P<0.01; ***, P<0.001.

Leaf extracts of *S. brachypetala* did not influence cell viability but high concentrations of 100 µg/ml induced cytotoxicity (Figure 3.5). At lower concentration, 10, 25 and 50 µg/ml the *S. brachypetala* leaf extract had a cell viability of 98.23, 97.84 and 98.34%, respectively. At the high concentration of 100 µg/ml, the leaf extract induced cytotoxicity with only 70.22% cell viability.

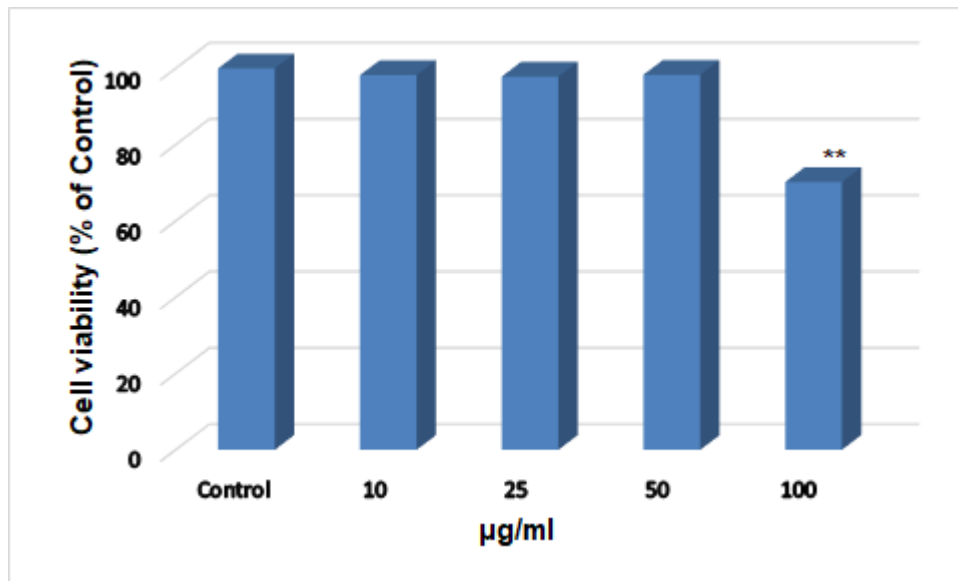


Figure 3.5: Cell viability measurement. Cells were incubated with indicated concentrations of extracts of leaves of *S. brachypetala* for 8 h, and then incubated with EZ-Cytox solution for 1 h. The leaf extract of *S. brachypetala* induced decreased cell viability at 100 µg/ml (n = 6). **, P<0.01; ***, P<0.001.

The levels of APP proteolytic products from the conditioned media using specific ELISA kits for Aβ40, sAPPβ-sw and sAPPα was then measured.

The leaf extract of *S. brachypetala* showed significant decrease in Aβ40 levels at the higher test concentrations of 50 µg/ml, for which the levels were reduced by $24.52 \pm 4.7\%$ and at 100 µg/ml, the percentage reduction of Aβ40 was $74.65 \pm 4.97\%$. At lower test concentrations, there was no significant reduction in Aβ40 levels. The decrease in secreted levels of Aβ40 by *S. brachypetala* leaf extract is shown in Figure 3.6.

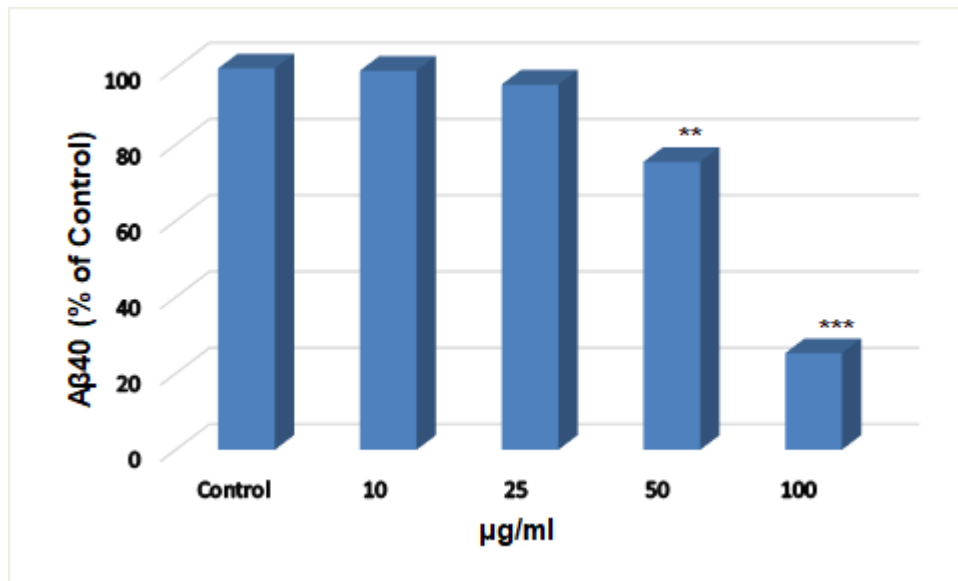


Figure 3.6: Change in the levels of Aβ40. Cells were incubated with indicated concentrations of leaf extract of *S. brachypetala* for 8 h, the levels of Aβ40 were measured from the conditioned media by using specific ELISA methods. The levels of Aβ40 (n = 4) were decreased by the extract at higher test concentrations **, P<0.01; ***, P<0.001.

In addition, this extract induced the decrease of sAPPβ-sw levels (Figure 3.7) and an increase of sAPPα level (Figure 3.8). At 50 µg/ml extract, the sAPPβ-sw level was reduced by $23.7 \pm 2.7\%$, while the sAPPα level was increased by $28 \pm 5\%$. At 100 µg/ml, the extract showed a higher reduction of sAPPβ-sw of $55.24 \pm 1.9\%$ while at the same concentration, sAPPα levels were increased by $45.08 \pm 11\%$.

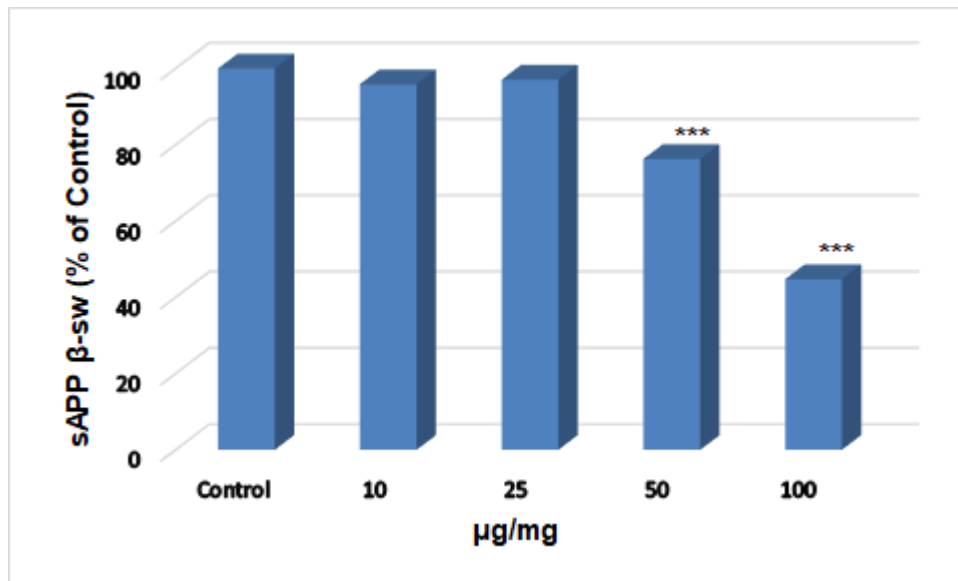


Figure 3.7: Change in the levels of sAPPβ-sw. Cells were incubated with indicated the concentrations of leaf extract of *S. brachypetala* for 8 h, the levels of sAPPβ-sw were measured from the conditioned media by using specific ELISA methods. The levels of sAPPβ-sw (n = 4) were decreased by the extract at higher test concentrations. **, P<0.01; ***, P<0.001.

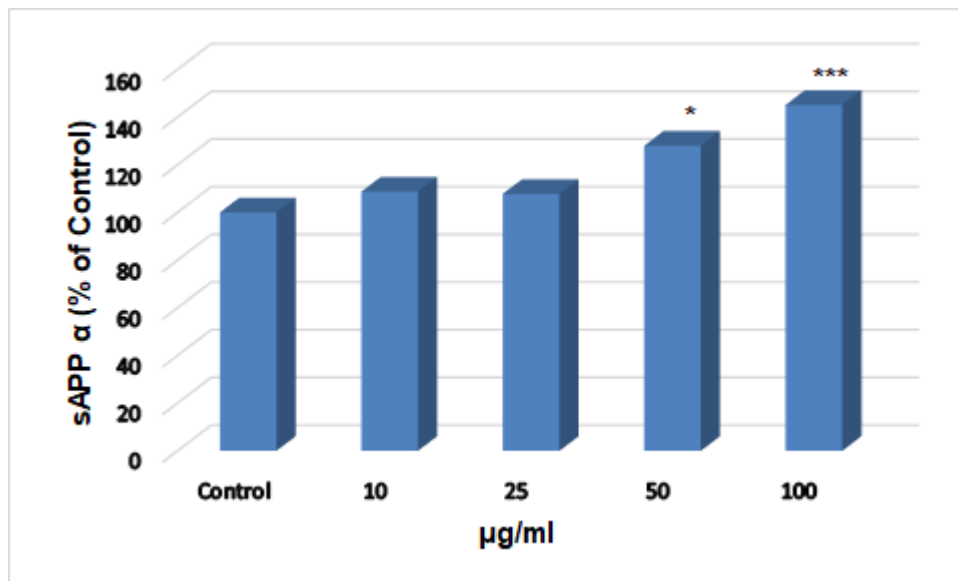


Figure 3.8: Change in the levels of sAPPα. Cells were incubated with the indicated test concentrations of leaf extract of *S. brachypetala* for 8 h, the levels of sAPPα were measured from the conditioned media by using specific ELISA methods. The levels of sAPPα (n = 4) were increased by the extract at higher test concentrations. **, P<0.01; ***, P<0.001.

From the dose dependent studies for the Aβ₄₂, Aβ₄₀, sAPPβ-sw and sAPPα bioassays, it was observed that only at the higher concentrations of 50 μg/ml and 100 μg/ml, *S.*

brachypetala leaf extract showed the significant results. These results suggest that the extract of *S. brachypetala* leaf extract decreased A β 42 production at non-toxic concentrations through decreasing amyloidogenic processing of APP and increasing the nonamyloidogenic processing of APP.

Two features, one involving the reduction of A β 42 through amyloidogenic process and other involving increase in the levels of sAPP α through nonamyloidogenic process are proven to exhibit the neuroprotective properties and also known to enhance the memory [38]. Thus these two features make the plant potent for the therapeutic target for Alzheimer's disease. In the present study, *S. brachypetala* leaf extract supported both the features to prove its efficacy to be potent towards the drug discovery. In a previous study, the plant was well researched for its activity towards the reduction of A β 42. The *S. brachypetala* (leaves and stalks) was found to significantly decrease the A β 42 peptide in the *in vivo* analysis on the mean mouse A β 42 concentration using ELISA assay [14]. The present study confirmed the previous findings for the potential of the plant to treat Alzheimer's disease.

3.5.3. Chemical characterization of DCM:MeOH leaf extract of *S. brachypetala*

A chemical analysis of DCM:MeOH leaf extract of *S. brachypetala* using the UPLC-QTOF-MS was undertaken to identify the major peaks and the compounds which could be responsible for the reduction of A β 42. In ESI negative mode (Figure 3.9), the peaks were mainly distributed within the 5.9 to 8 min range of the 18 min chromatogram. The ionization in ESI positive mode did not produce enough ionization of compounds for analysis and it was not further analysed.

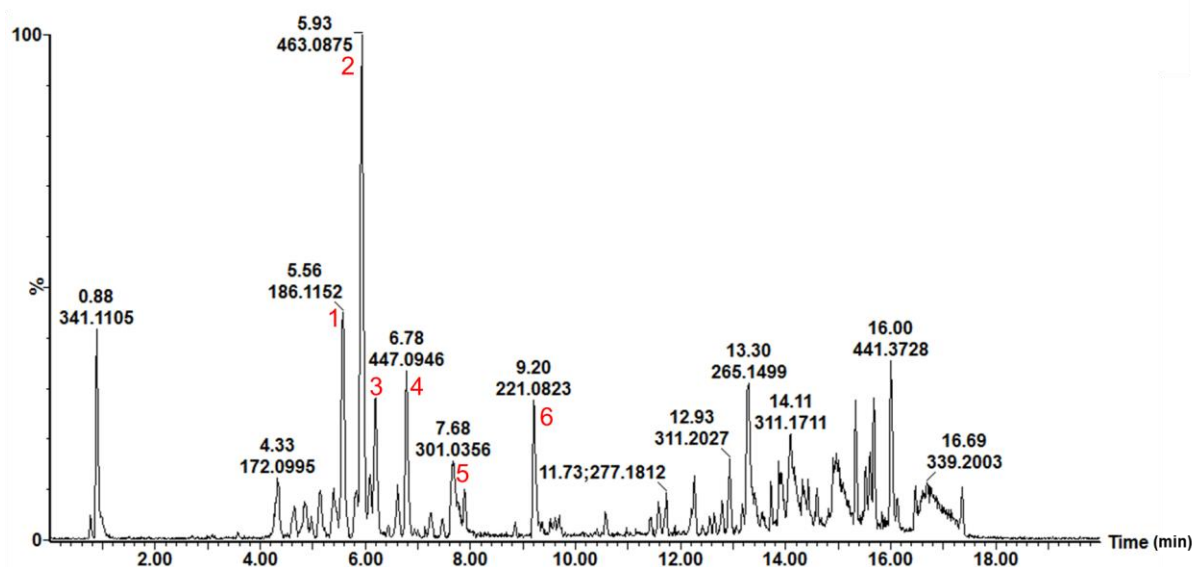


Figure 3.9: ESI negative mode BPI chromatogram of *S. brachypetala* leaf extract.

The chromatogram indicated the general elution region for polar to medium polar compounds. A total of five compounds were tentatively identified shown in Table 3.2, belonging to the class of flavonoids, peak 1 was unidentified; peak 2 identified as isoquercetin (**25**) with m/z 463.0875 [M-H]⁻; peak 3 identified as myricetin-3-O- α -L-rhamnopyranoside (**26**) with m/z 463.0872 [M-H]⁻; peak 4 identified as quercetin-3-O-rhamnoside (**27**) with m/z 447.0946 [M-H]⁻ and peak 5 identified as quercetin (**28**) with m/z 301.0356 [M-H]⁻ and peak 6 was unidentified.

Table 3.2: Tentative identification of compounds obtained from ESI-MS negative mode of DCM:MeOH extract of *Schotia brachypetala* (leaf)

Peak	RT (min)	Acquired [M-H] ⁻ m/z	Formula of possible structure	Theoretical [M-H] ⁻ m/z	Possible structure	Mass error (ppm)	MS/MS Data (Fragment s)	Ref.
1	5.56	186.1152	-	-	unidentified	-	-	-

2	5.93	463.0875	C ₂₁ H ₂₀ O ₁₂	463.0877	isoquercetin	-0.4	317.0262, 301.0377, 271.0243, 179.0014	[39]
3	6.23	463.0877	C ₂₁ H ₂₀ O ₁₂	463.0877	myricetin-3- O-alpha-L- rhamnopyra noside	0.0	301.0344, 300.0272, 271.0257	[40], [41]
4	6.78	447.0943	C ₂₁ H ₂₀ O ₁₁	447.0927	quercetin-3- O- rhamnoside	3.6	301.0352, 300.0436, 271.0420, 255.0608	[42], [40]
5	7.68	301.0351	C ₁₅ H ₁₀ O ₇	301.0348	Quercetin	1.0	151.0066, 121.0306, 107.0152	[40]
6	9.20	221.0823	C ₁₆ H ₁₃ O	221.0966	unidentified	64.6	-	-

The most intense peak (labelled as peak 2) in the DCM:MeOH leaf extract of *S. brachypetala* was observed at m/z 463.0875 [M-H]⁻ at retention time 5.93 min. The molecular formula C₂₁H₂₀O₁₂ was generated through the MassLynx with a normalized iFit value 0.212 (**supplementary data 1**). The compound was identified as Isoquercetin (**25**) shown in Figure 3.14.

The compound was further confirmed by using pure standard isoquercetin shown in Figure 3.10. The confirmation of compound was based on the comparison of close retention time and same accurate masses of peak 2 (retention time 5.93 min and m/z 463.0872) with that of pure standard isoquercetin (retention time 6.23 minutes and m/z 463.0882).

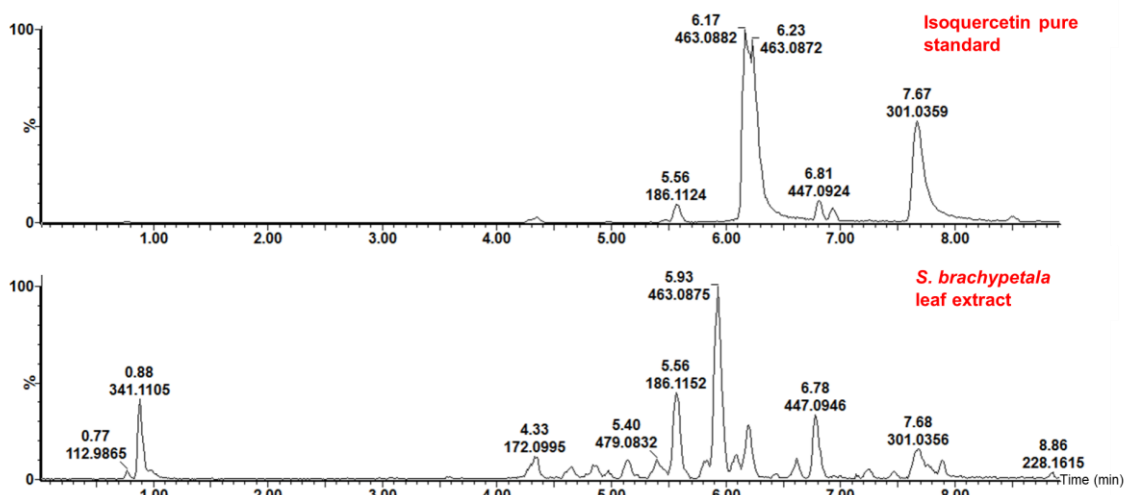


Figure 3.10: ESI negative mode BPI chromatogram of *S. brachypetala* leaf extract overlaid with isoquercetin pure standard. The pure standard peak was at retention time 6.23 min and m/z 463.0872 which corresponded to the peak 2 at retention time 5.93 min and m/z 463.0875 in the extract.

The MS/MS data analysis (**supplementary data 4**) gave a quercetin fragment at m/z 301.0344 $[M-162]^-$ with the loss of a glucoside moiety, m/z 271.0257 $[M-162-CO-2H]^-$. Compound was also confirmed by comparing fragmentation pattern with that of pure standard isoquercetin shown in Figure 3.11. The fragmentation pattern was also supported by the literature studies [40], [41].

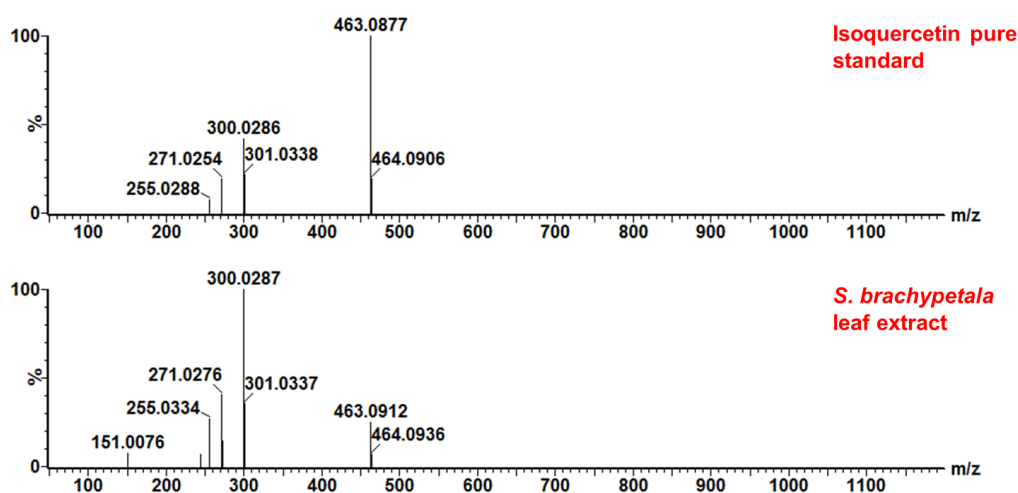


Figure 3.11: MS/MS peak 3 (isoquercetin) in *S. brachypetala* crude extract overlaid with the MS/MS of isoquercetin pure standard

Isoquercetin has many therapeutic properties including antioxidants, anti-allergic and anti-inflammatory. It was found to exhibit neuroprotective properties by increasing Nrf2 through ERK 1/2 pathways, preventing oxygen-glucose deprivation followed by reoxygenation (OGD/R) -induced increase in apoptosis, lactate dehydrogenase release and reduction in cell viability. These studies evaluated the potency of isoquercetin towards the treatment of ischemia and other associated diseases [43]. Isoquercetin significantly reduced the glutamate-induced oxidative neurotoxicity in HT22 hippocampal cells at a concentration of 100 μ M ($p < 0.001$) with an IC_{50} of $88.8 \pm 11.1^{**}$ ($**p < 0.001$) [44]. Isoquercetin also exhibited the protective effects against toxicity induced by 6-hydroxy dopamine (6-OHDA) in rat pheochromocytoma (PC12) cells. It showed a significant increase in cell viability at 10 μ M with a value of 73.2 ± 1.01 ($p < 0.001$) as compared to 6-OHDA (50.80 ± 1.66) at 100 μ M [45]. There is no previously reported studies which showed direct relation of isoquercetin with A β 42 reduction and Alzheimer's disease, however its neuroprotective properties may contribute to the fact that it could be an active ingredient in the present study.

Peak 3 was observed at m/z 463.0872 [M-H]⁻ with a retention time of 6.20 min. The molecular formula of compound was generated through MassLynx with a normalised iFit value of 0.176 (**supplementary data 3**). The structure of the compound corresponding to this peak was tentatively identified as myricetin-3-O- α -L-rhamnopyranoside (**26**) shown in Figure 3.14.

The second order mass spectrum of the precursor ion showed fragments at m/z 316.0229 [M-146-2H]⁻ with the loss of a rhamnosyl moiety, at m/z 271.0243 [M-146-CO-H₂O-H]⁻ and at m/z 179.0014 (**supplementary data 2**). The observed fragmentation pattern shown was in agreement with the fragmentation pattern obtained from literature and led to the identification of the compound [39].

In a previous study myricetin was found to exhibit an anti-tau effect when tested at 50 μ M in HeLa-C3 cells [46]. It has also been found to exhibit oxidative effects towards the Met35 residue in A β peptides by interfering in A β aggregate formation which was obtained in docking studies [47]. These studies supported the fact that myricetin and its derivatives play an

important role in the neuroprotective activities and could be responsible for the reduction of A β 42.

Peak 4 represent a precursor ion at m/z 447.0946 [M-H]⁻ with retention time 6.78 min. The molecular formula of compound, C₂₁H₂₀O₁₁ was generated through MassLynx with a Normalised iFit value 0.65 (**supplementary data 5**) identified as quercetin-3-O-rhamnoside (**27**) shown in Figure 3.14.

The MS/MS spectra (**supplementary data 6**) produced a quercetin fragment ion at m/z 301.0352 [M-146]⁻ which involved the loss of a rhamnosyl moiety, a typical flavon-3-O-monoglycoside fragment at m/z 271.0420 [M-146-CO-H]⁻. The same fragment pattern was described in literature [42], [40].

The flavonoid quercetin-3-O-rhamnoside was previously isolated from the aqueous extract of *S. brachypetala* leaves and stalks. *S. brachypetala* leaves (100 mg/kg/day) and *S. brachypetala* stalk (100 mg/kg/day) significantly reduced A β 42 concentration by 76.2% and 72.8% as compared to (Lipopolysaccharide) LPS group, on a treatment for 6 days. This study indicated that identified flavonoids including quercetin-3-O-rhamnoside could be responsible for the activity [14]. The same compound isolated from *Salsola grandis* showed no percentage inhibition when tested against AChE [48]. These previous studies suggest that the compound was inactive towards AChE, however based on its reported reduction of A β 42 concentration points to the compound being one of the active compounds in the plant extract.

The peak labelled as 5 had a precursor molecular ion at m/z 301.0356 with retention time 7.68 min and corresponded to the proposed molecular formula C₁₅H₁₀O₇ generated through MassLynx with a normalised iFit value of 0.026 (**supplementary data 7**). The compound was tentatively identified as quercetin (**28**) shown in Figure 3.14.

The retention time of the pure standard of quercetin of 7.65 min (m/z 301.0384) with that of peak 5 in the crude extract retention time 7.68 min (m/z 609.1465) confirmed the identity of peak 5 as quercetin (Figure 3.12).

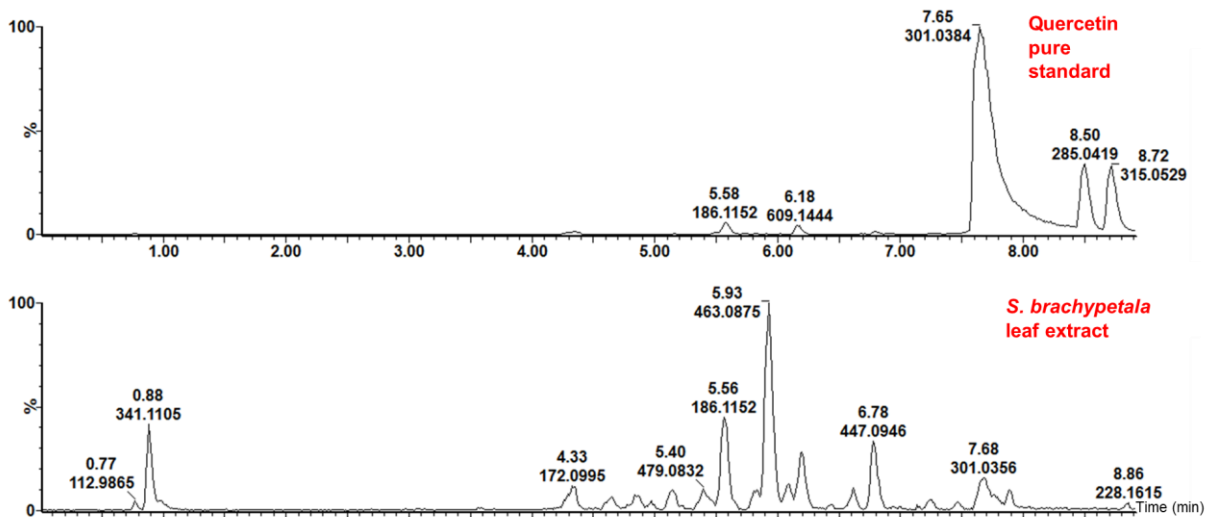


Figure 3.12: ESI negative mode BPI chromatogram of *S. brachypetala* leaf extract overlaid with isoquercetin pure standard. The pure standard peak was at retention time 7.65 minutes and m/z 301.0384 which corresponded to the peak at retention time 7.68 and m/z 301.0356 in the extract.

A comparison of the MS/MS fragmentation pattern of peak 5 (quercetin) in the crude extract with that of the pure standard of quercetin also confirmed the identity of peak 5 as quercetin (Figure 3.13).

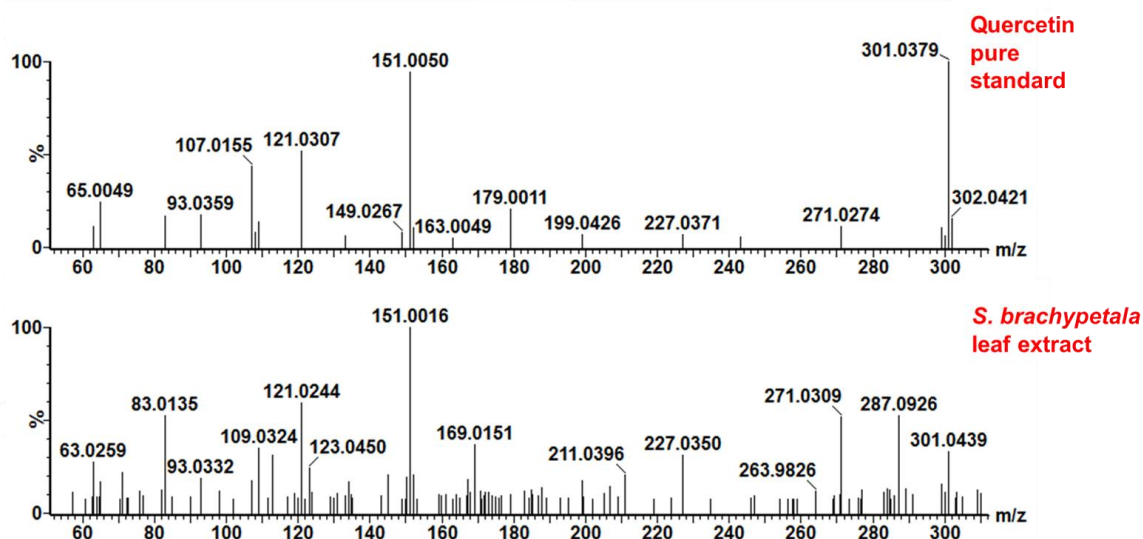
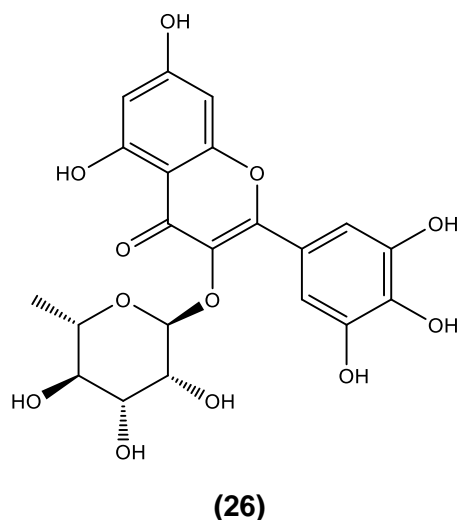
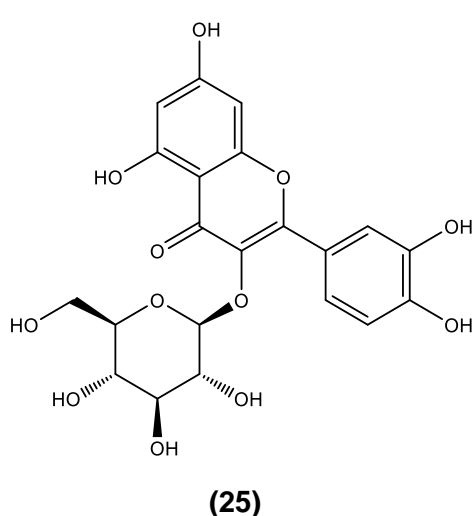


Figure 3.13: MS/MS peak 5 (quercetin) in *S. brachypetala* crude extract overlaid with the MS/MS of quercetin pure standard.

The compound was also confirmed by comparing the fragmentation pattern with literature. The MS/MS spectra (**supplementary data 8**) gave the fragment ions at m/z 271.0274 [M-CO-H]⁻, m/z 179.0011 and m/z 151.0050 which are similar to the fragmentation pattern obtained from literature [40].

The quercetin flavonoid showed strong antioxidant and free radical scavenging properties in *in vitro* studies [49]. In an *in vivo* study quercetin significantly decreased β -amyloidosis in aged 3xTg-AD mice when treated for three months as compared to the vehicle-treated 3xTg-AD mice. The reduction in the $A\beta_{(1-40)}$ and $A\beta_{(1-42)}$ levels were detected in the hippocampus of the quercetin-treated 3xTg-AD mice compared to the vehicle-treated 3xTg-AD mice [50]. At the lower doses (5 and 10 μ M), quercetin on pre-treatment with neurons for 1 h significantly reduced the $A\beta_{(1-42)}$ - induced cytotoxicity [51]. Quercetin showed the neuroprotective effects by inhibiting Lipoxidation, neuronal cell apoptosis, hippocampal neuronal cell damage after exposure to cadmium [52]. Quercetin was also reported to induce NF-E₂-related factor-2 (Nrf₂) which increased the neuroprotection in cerebellar granule neuronal culture [53]. Quercetin protected HT22 murine neuroblastoma cells from $A\beta_{(25-35)}$ oxidative attack by inhibiting $A\beta$ fibril formation [51]. All these previous studies supported the fact that quercetin could be the active ingredient in the present study because of its ability to reduce the $A\beta$ levels in different bioassays reported in previous studies.



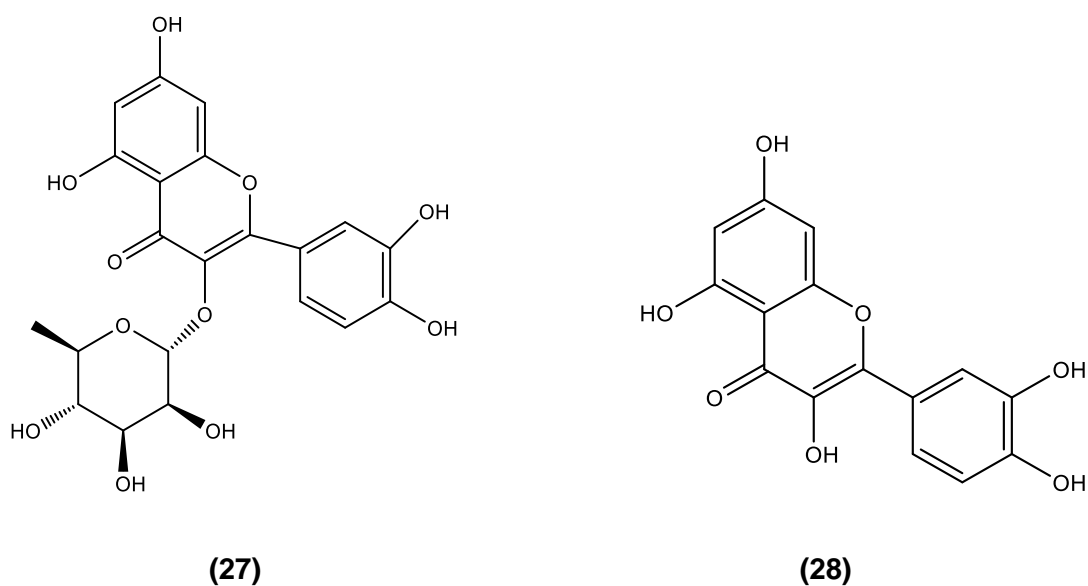


Figure 3.14: Chemical structures of compounds identified from *S. brachypetala* leaves, Peak 2: isoquercetin (**25**), Peak 3: myricetin-3-O- α -L-rhamnopyranoside (**26**), Peak 4: quercetin-3-O-rhamnoside (**27**), Peak 5: quercetin (**28**)

After LC-MS analysis, it was found that main class of compounds identified in the *S. brachypetala* leaves is flavonoid glycosides. The flavonoid glycosides were identified from *S. brachypetala*, with a typical characteristic fragmentation pattern of flavonol-3-O-glycosides produce a fragment ion at m/z 301 which revealed the presence of a quercetin moiety. The common secondary fragment ions at m/z 271, m/z 255, m/z 179 and m/z 151 were produced because of the fragmentation of quercetin. The fragments m/z 271 and m/z 255 were produced due to the loss of carbon monoxide (-28 amu) and carbon dioxide (-44 amu) respectively. The fragment ions at m/z 151 and m/z 179 were produced as a result of Retro Diel's alder fragmentation pattern. The fragmentation pathway followed by quercetin derivatives is shown in Figure 3.15.

Flavonoid glycosides are found in a wide variety of natural foods and have shown positive effects against dementia and Alzheimer's disease in various cell cultures and animal models [54]. In a transgenic mouse model, a flavonoid named nobiletin was found to improve the A β induced memory impairment. It also reduced the A β plaques formation in the hippocampus of APP-SL 7-5 Tg mice [55]. Luteolin another flavonoid, was found to inhibit the A β peptides with

an inhibition percentage >70% and >85% at the concentrations of 20 and 40 μM , respectively in both human “Swedish” mutant APP N2a and Tg2576 mouse-derived primary neuronal cells [56]. These previous studies indicated that the flavonoid glycosides identified from *S. brachypetala* could also play an important role in the anti-Alzheimer’s drug discovery.

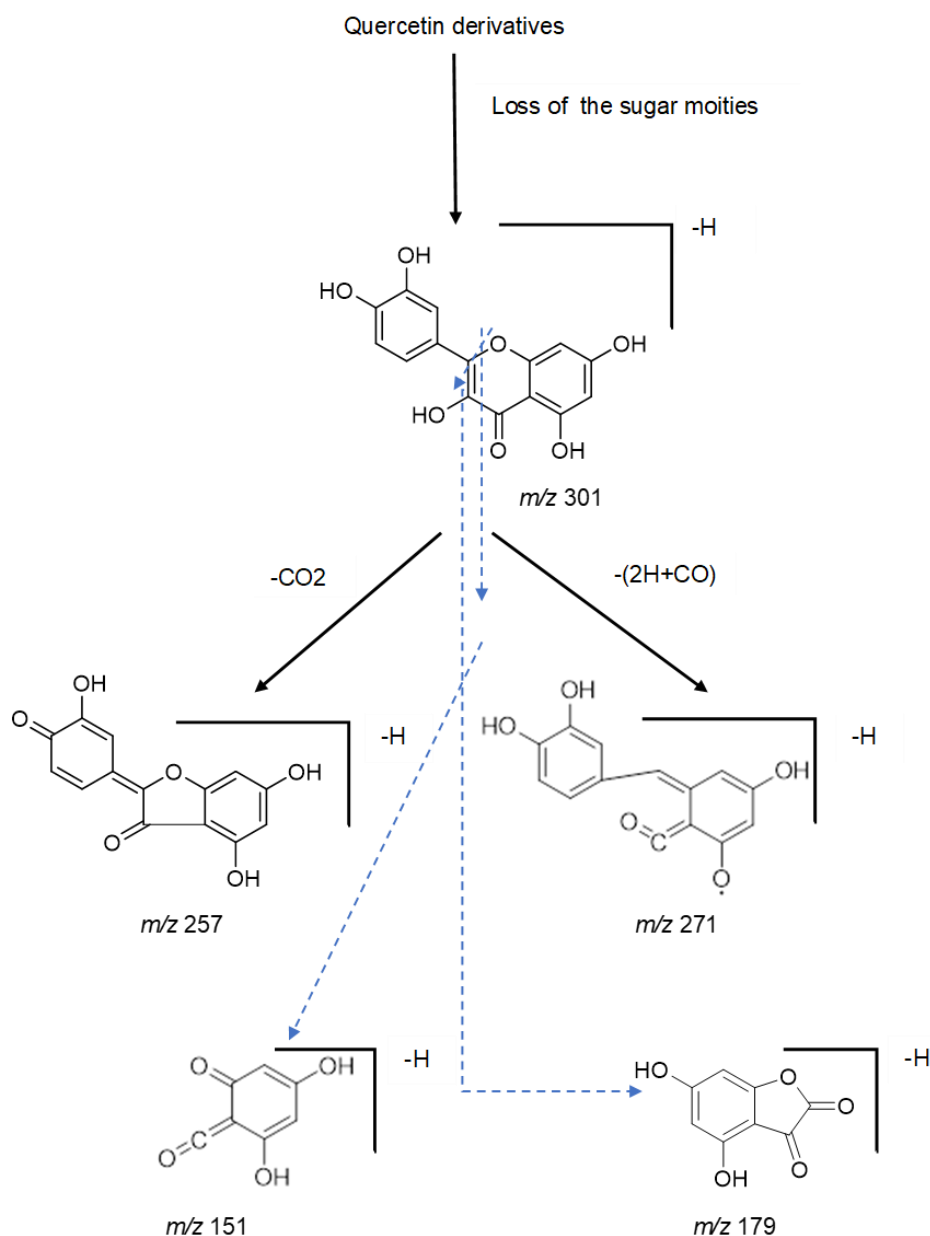


Figure 3.15: Fragmentation pattern of quercetin derivatives adapted from [43] and [39]

3.5.4. The effect of *C. paniculata* on APP processing and cytotoxicity

The extract of leaves of *C. paniculata* significantly reduced A β 42 production in a dose-dependent manner. At concentrations 10, 25, 50 and 100 μ g/ml, the leaf extract significantly reduced the A β 42 levels by $29.03 \pm 1.4\%$, $42.40 \pm 1.7\%$, $47.50 \pm 5.8\%$ and $61.14 \pm 9.3\%$, respectively (Figure 3.16).

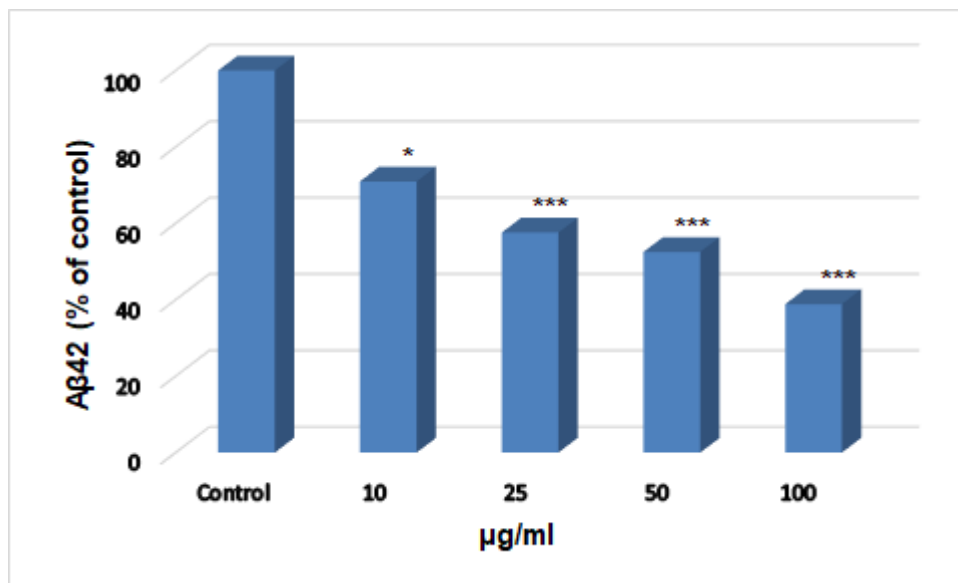


Figure 3.16: Change in the levels of A β 42. Cells were incubated with indicated concentrations of leaf extract of *C. paniculata* for 8 h, the levels of A β 42 were measured from the conditioned media by using specific ELISA methods. The levels of A β 42 (n = 4) were decreased by the extract in a dose dependant manner. **, P<0.01; ***, P<0.001.

C. paniculata leaf extract (Figure 3.17) did not influence cell viability at low test concentrations but at high concentrations it induced some cytotoxicity. At the lower test concentrations of 10, 25 and 50 μ g/ml *C. paniculata* leaf extract maintained cell viability at 105.4, 97.2 and 91.5%, respectively while at the high test concentration of 100 μ g/ml, the leaf extract induced cytotoxicity with cell viability of 74.52%.

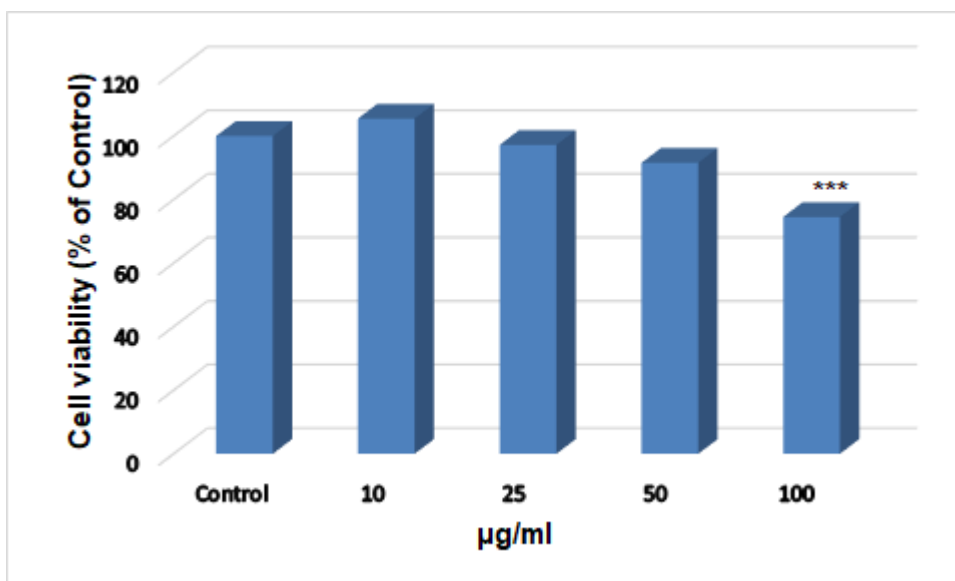


Figure 3.17: Cell viability measurement. Cells were incubated with indicated concentrations of extracts of leaves of *C. paniculata* for 8 h, and then incubated with EZ-Cytox solution for 1 h. The leaf extract of *C. paniculata* induced the decrease of cell viability at the highest test concentration of 100 µg/ml (n = 6).

The levels of APP proteolytic products from the conditioned media using specific ELISA kits for A β 40, sAPP β -sw and sAPP α were then measured. The level of A β 40 was significantly decreased in a dose dependent manner by $47.7 \pm 9.1\%$, $79.6 \pm 2.2\%$ and $90.19 \pm 2.5\%$ at 25, 50 and 100µg/ml of extract, respectively (Figure 3.18).

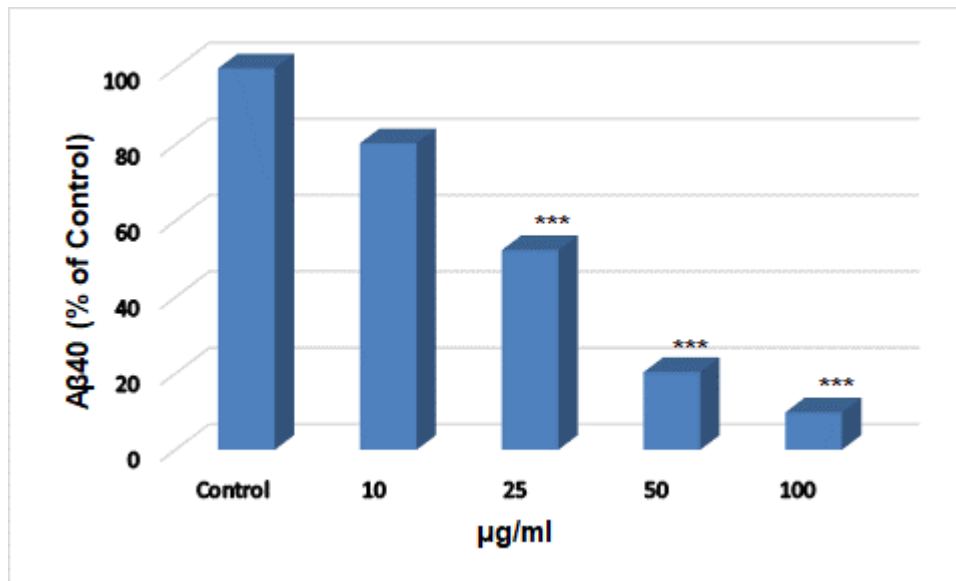


Figure 3.18: Change in the levels of Aβ40. Cells were incubated with indicated concentrations of leaf extract of *C. paniculata* for 8 h, the levels of Aβ40 were measured from the conditioned media by using specific ELISA methods. The levels of Aβ40 (n = 4) were decreased by the leaf extract in a dose dependant manner. **, P<0.01; ***, P<0.001.

The level of sAPPβ-sw was also significantly decreased in a dose dependent manner by $37.9 \pm 4.4\%$, $56.6 \pm 0.8\%$ and $74.57 \pm 1.5\%$ at 25, 50 and 100 μg/ml of extract, respectively (Figure 3.19.). In addition, this extract also significantly reduced sAPPα production by $33.99 \pm 3\%$ and $49.47 \pm 10\%$ at 25 and 50 μg/ml, respectively as shown in Figure 3.20.

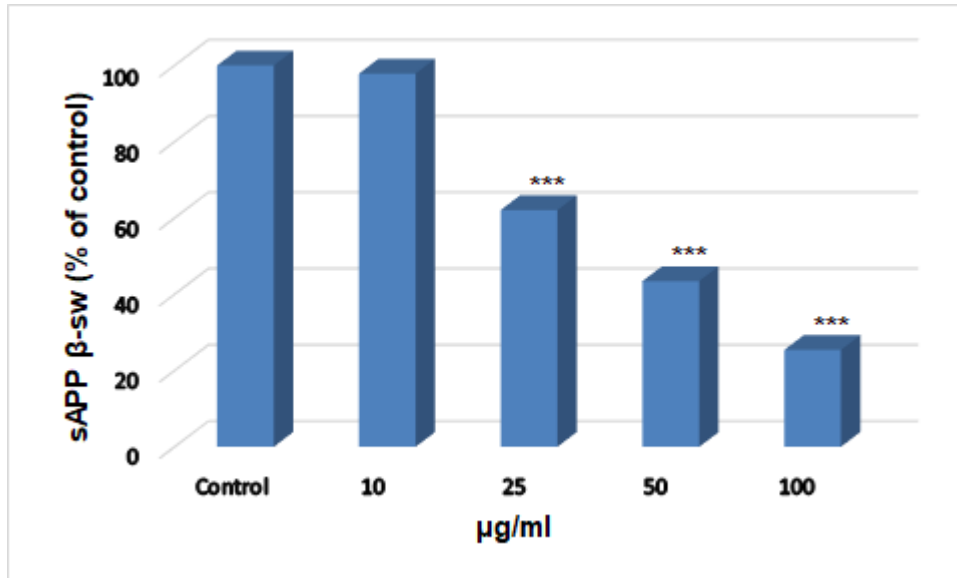


Figure 3.19: Change in the levels of sAPP β . Cells were incubated with indicated concentrations of leaf extract of *C. paniculata* for 8 h, the levels of sAPP β were measured from the conditioned media by using specific ELISA methods. The levels of sAPP β (n = 4) were increased by the extract in a dose dependant manner **, P<0.01; ***, P<0.001.

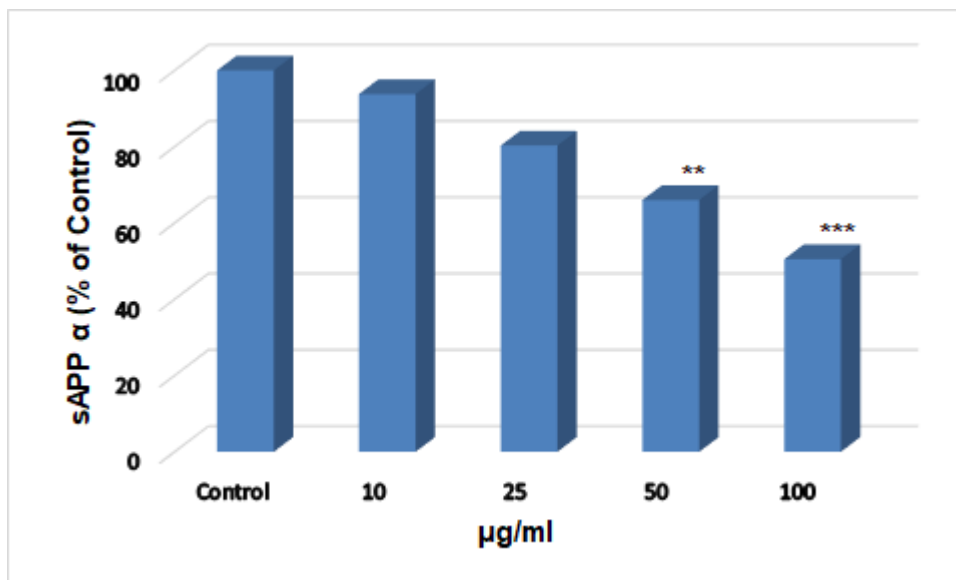


Figure 3.20: Change in the levels of sAPP α . Cells were incubated with indicated concentrations of leaf extract of *C. paniculata* for 8 h, the levels of sAPP α were measured from the conditioned media by using specific ELISA methods. The levels of sAPP α (n = 4) were increased by the extract in a dose dependant manner **, P<0.01; ***, P<0.001.

These results suggested that the DCM:MeOH extract of *C. paniculata* leaves decreased A β 42 production at non-toxic concentrations through decreasing amyloidogenic processing of APP but there was no increase in the nonamyloidogenic processing of APP. The ability of *C. paniculata* leaf extract to reduce A β 42 supported its efficacy in the development new drugs for the treatment of Alzheimer's disease. This study on the plant is the first report showing potent activity towards the reduction of A β 42.

3.5.5. Chemical characterization of DCM:MeOH leaf extract of *C. paniculata*

The DCM:MeOH leaf extract of *C. paniculata* was analysed using the UPLC-QTOF-MS to identify the major peaks and the compounds which could be responsible for the reduction of A β 42. The ionization was poor in the positive mode so it was not used for further analysis. In ESI negative mode, the peaks were mainly distributed within 6.16 to 11.38 minute range of the 18 minute chromatogram (Figure 3.21).

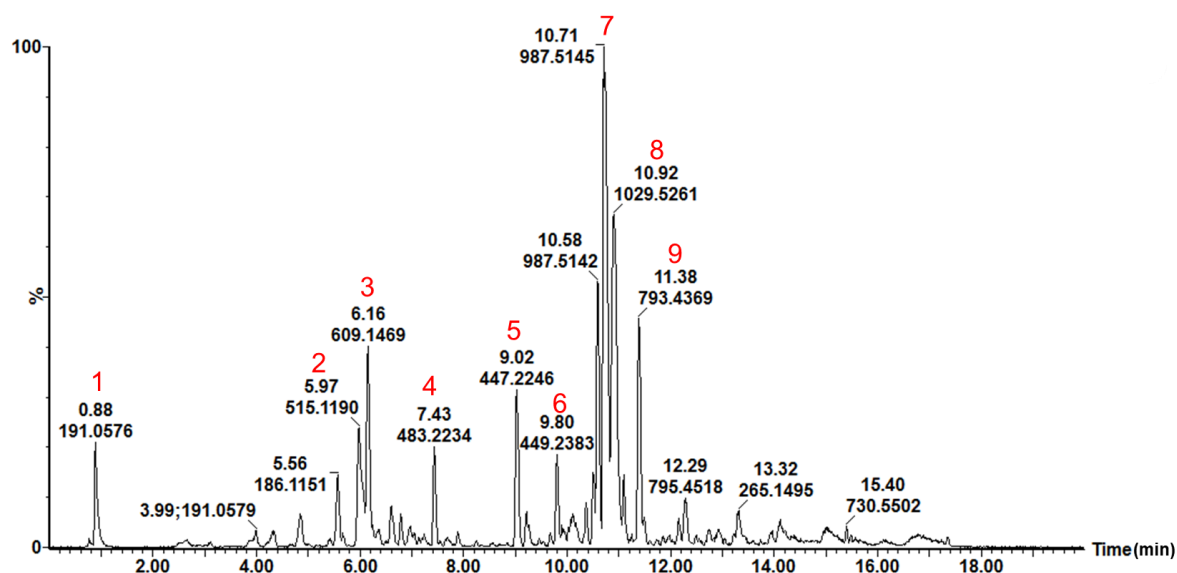


Figure 3.21: ESI negative mode BPI chromatogram of *C. paniculata* leaf extract.

This indicated the general elution region for polar to medium polar compounds. Using ESI negative mode, a total of nine compounds were tentatively identified shown in Table 3.3. Peak 1 was identified as quinic acid (**29**) with m/z 191.0576 $[M-H]^-$, peak 2 identified as 3,5-dicaffeoylquinic acid (**30**) with m/z 515.1190 $[M-H]^-$, peak 3 identified as rutin (**31**) with m/z 609.1469 $[M-H]^-$, peak 4 identified as valeriananoid E (**32**) with m/z 483.2234 $[M-H]^-$, peak 5 identified as acuminoside (**33**) with m/z 447.2246 $[M-H]^-$, peak 6 identified as dictamnocide D (**34**) with m/z 449.2383 $[M-H]^-$, peak 7 identified as 2''-O- β -D-glucopyranosylsaikosaponin B2 (**35**) with m/z 987.5145 $[M-H]^-$; peak 8 identified as clinoposaponin C (**36**) with m/z 1029.5261 $[M-H]^-$ and peak 9 identified as spinasaponin C (**37**) with m/z 793.4369 $[M-H]^-$.

Table 3.3: Tentative identification of compounds obtained from ESI-MS negative mode of DCM:MeOH extract of *C. paniculata* leaves

Peak	RT (min)	Acquired $[M-H]^-$ m/z	Formula of Possible structure	Theoretical $[M-H]^-$ m/z	Possible structure	Mass error (ppm)	MS/MS Data (Fragments)	Ref.
1	0.88	191.0576	C ₇ H ₁₂ O ₆	191.0556	quinic acid	10.5	161.0456, 131.0485	-
2	5.97	515.1190	C ₂₅ H ₂₄ O ₁₂	515.1190	3,5-dicaffeoylquinic acid	0.0	353.0883, 191.0572, 179.0363	[57]
3	6.16	609.1489	C ₂₇ H ₃₀ O ₁₆	609.1514	rutin	-4.1	301.0353, 300.0292, 271.0259	[58]
4	7.43	483.2234	C ₂₄ H ₃₆ O ₁₀	483.2230	valeriananoid E	0.8	277.1818, 173.0477, 135.0458	-
5	9.02	447.2246	C ₂₁ H ₃₆ O ₁₀	447.2230	Acuminoside	3.6	315.1824, 161.0469	-
6	9.80	449.2383	C ₂₁ H ₃₈ O ₁₀	449.2387	dictamnocide D	-0.9	161.0472, 101.0252	-
7	10.71	941.5082	C ₄₈ H ₇₈ O ₁₈	941.5086	2''-O- β -D-glucopyranosylsaikosaponin B2	-0.4	471.3481, 469.1565	[59]
8	10.92	983.5211	C ₅₀ H ₈₀ O ₁₉	983.5216	clinoposaponin C	-0.5	471.3488, 469.1578	[60]
9	11.38	793.4372	C ₄₂ H ₆₆ O ₁₄	793.4374	spinasaponin A	-0.3	631.3863	[61]

Peak 1 in the DCM:MeOH leaf extract of *C. paniculata* was observed at m/z 191.057 $[M-H]^-$ at retention time of 0.88 min. The molecular formula of the compound, C₇H₁₂O₆ was generated

through MassLynx with a normalized iFit value 0.025 (**supplementary data 9**). The structure of the compound corresponding to this peak was tentatively identified as quinic acid (**29**) shown in Figure 3.26. The MS/MS spectra (**supplementary data 10**) gave fragment ions at m/z 161.0456 with the loss of one hydroxyl ion and one oxygen atom. The other fragment ion was produced at m/z 131.0485 with the loss of formic acid and one hydroxyl ion.

In a previous study, it was reported that quinic acid did not prevent A β 42 aggregation when tested on SH-SY₅Y cell line using thioflavin T (Th-T) bioassays with an IC₅₀ value >100 μ M [62]. Although in the previous studies quinic acid itself did not show any significant activity, its derivatives (discussed for peak 2) were active towards the prevention of A β 42 aggregation [62].

Peak 2 had a m/z 515.1190 [M-H]⁻ with retention time 5.97 min. The molecular formula, C₂₅H₂₄O₁₂ was generated through MassLynx with a normalized iFit value 0.444 (**supplementary data 11**). The compound was tentatively identified as 3,5-dicaffeoylquinic acid (**30**) and the structure is shown in Figure 3.26. The MS/MS spectrum (**supplementary data 12**) produced the fragment ions at m/z 353.0883 [M-H-162]⁻ with the loss of a caffeoyl moiety, a quinic acid fragment m/z 191.0572 [M-H-162-162]⁻ with the loss of another caffeoyl moiety and a caffeic acid fragment ion at m/z 179,0363 [M-H-336]⁻. The MS³ base peak produced at m/z 191.0572 distinguished 3,5-dicaffeoylquinic acid from the isomers 3,4-dicaffeoylquinic acid and 4,5-dicaffeoylquinic acid. The isomers 3,4-dicaffeoylquinic acid and 4,5-dicaffeoylquinic acid produced MS³ base peak at m/z 173.0 [63], [57]. Therefore, the fragmentation pattern supported the identification of 3,5-dicaffeoylquinic acid.

The fragmentation pathway followed by 3,5-dicaffeoylquinic acid as an example is shown in Figure 3.23.

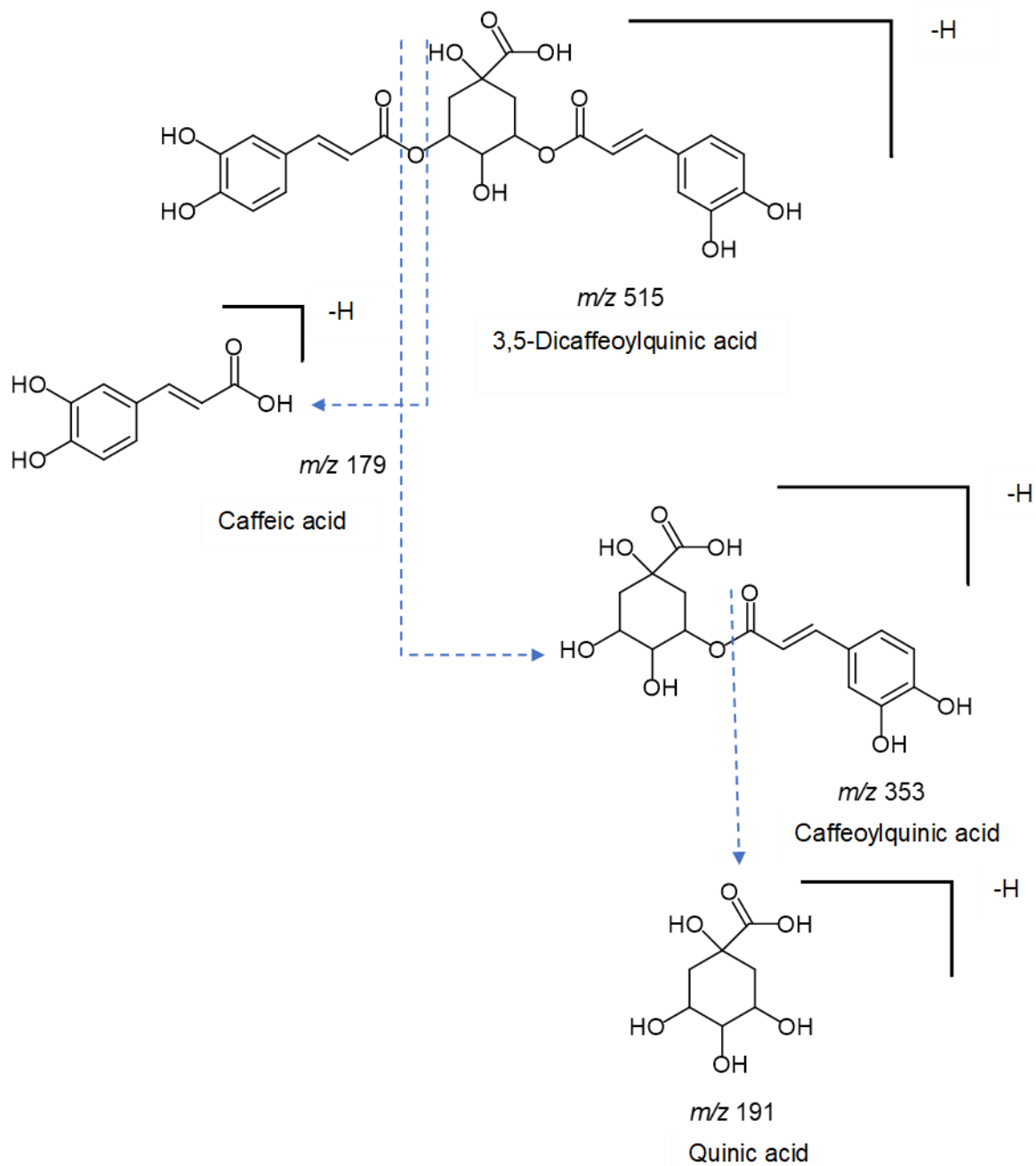


Figure 3.22: Fragmentation pattern of 3,5-Dicaffeoylquinic acid adapted from [57]

The 3,5-dicaffeoylquinic acid significantly improved the cognitive dysfunction caused because of the severe TMT induced neurotoxicity. It also inhibited AChE by recovering acetylcholine, a neurotransmitter in the cholinergic system [64]. The 3,5-dicaffeoylquinic acid isolated from

Aster scaber exhibited neuroprotective effects in A β -sensitive PC12 cells by significantly reducing the A β ₂₅₋₃₅ cytotoxicity with a cell viability 90.0% or 75% when measured by MTT [65]. The 3,5-dicaffeoylquinic acid was also tested in behavioural tests (Y-maze, passive avoidance and Morris water maze test) in which it significantly improved the learning and memory deficits. It was found to prevent neuronal apoptosis by protecting mitochondrial activities [64].

The peak labelled as 3 produced a m/z 609.1469 [M-H]⁻ with retention time 6.16 min. The molecular formula C₂₇H₃₀O₁₆ was generated through MassLynx with a normalised iFit value of 0.003 (**supplementary data 13**). The compound was tentatively identified as rutin (**31**) and the structure of the compound is shown in Figure 3.26. It was further confirmed by comparing peak 6 in the DCM:MeOH extract of leaves of *C. paniculata* with a pure standard of rutin (Figure 3.24). A comparison of the MS/MS (**supplementary data 14**) fragmentation pattern of peak 6 (rutin) in the crude extract with that of the pure standard of rutin also confirmed the identity of peak 6 as rutin. Further, the retention time of the pure standard of rutin of 6.16 minutes (m/z 609.1454) with that of peak 6 in the crude extract retention time 6.16 min at m/z 609.1469 confirmed the identity of peak 6 as rutin.

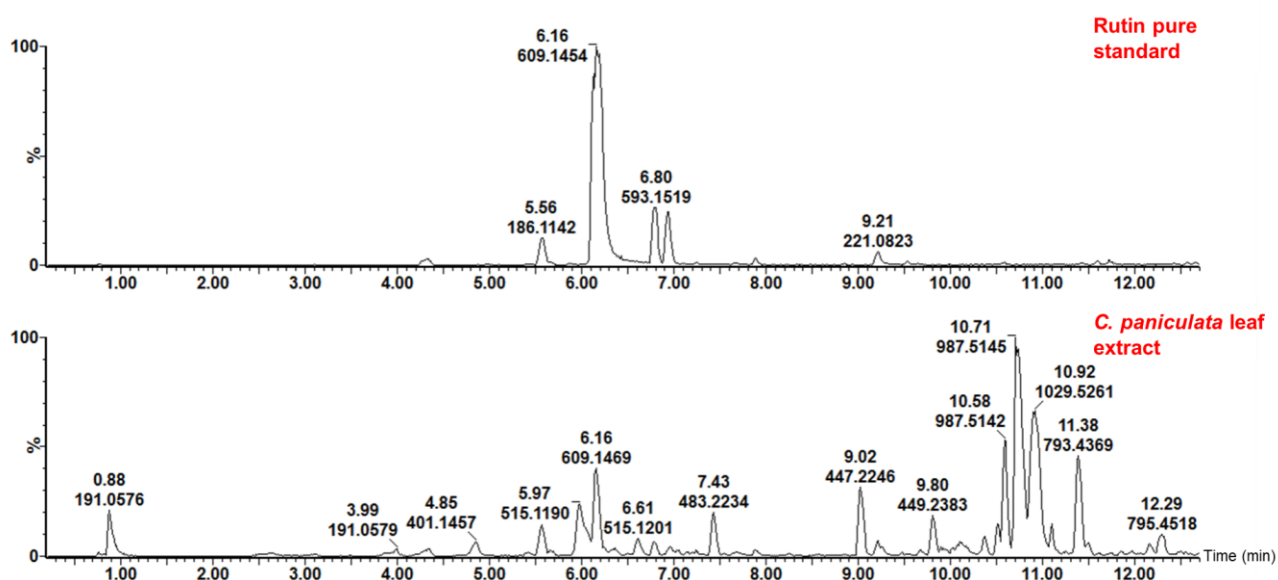


Figure 3.23: ESI negative mode BPI chromatogram of leaves of *C. paniculata* extracted with DCM:MeOH overlaid with standard rutin. The standard rutin had a retention time of 6.16 min and an m/z of 609.1454 which corresponded to peak 3 at retention time 6.16 min and m/z 609.1469 in the extract.

The MS² analysis gave a quercetin fragment at m/z 301.0344 [M-308]⁻ with the successive loss of two hexose moieties, m/z 271.0257 [M-308-CO-2H]⁻ and m/z 179.0221 due to the RDA fragmentation pattern. The fragmentation pattern was also supported with literature studies [40], [41]. Comparison of the accurate mass, retention time and the fragmentation pattern of peak 3 and the pure standard confirmed its structure (Figure 3.25).

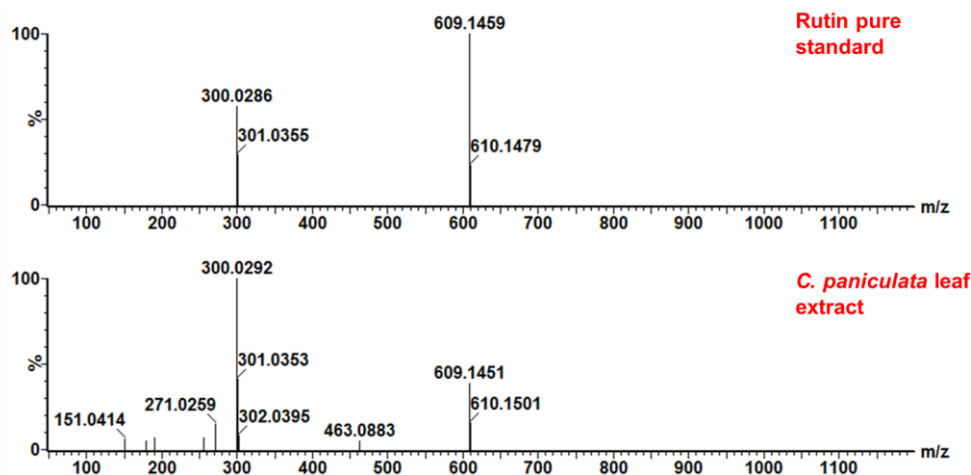


Figure 3.24: MS/MS peak 3 (rutin) in *C. paniculata* crude extract overlaid with the MS/MS of rutin pure standard

Rutin, is one of the major flavonoid glycosides isolated from *Ginkgo biloba*, which is one of the most investigated plant remedies for Alzheimer's disease, however its efficacy for the treatment of Alzheimer's is still controversial [66]. Rutin on oral administration significantly reduced the memory deficits in APP^{swe}/PS1^{dE9} transgenic mice by decreasing A β oligomer levels. It also increased super oxide dismutase (SOD) activity and glutathione (GSH)/glutathione disulfide (GSSG) ratio and also decreased interleukin (IL)-1 β and IL-6 levels in the brain. These results indicated that rutin as a promising agent for Alzheimer's disease [67]

The ESI negative mode for peak 4 gave a precursor ion peak at m/z 483.2234 and a molecular formula C₂₄H₃₆O₁₀ was generated through MassLynx with a normalised iFit value 0.075 (**supplementary data 15**). A sesquiterpenoid named valeriananoid E (**32**) was tentatively

identified and its structure is shown in Figure 3.26. MS/MS spectra gave the fragment ions at m/z 277.1818, m/z 173.0477 and m/z 135.0458 (**supplementary data 16**).

The compound was previously isolated from *Valeriana jatamansi* which is used traditionally for the treatment of hypochondriasis, nervous disorders, migraine, hysteria, stress, depression, insomnia, neuralgia and neuroasthenia. Valeriananoids D–E isolated from *V. jatamansi* showed less than 10% AChE inhibition activity at a concentration of 50 μ M as compared to the positive control, tacrine with an inhibition percentage, 47.6% at 0.33 μ M [68].

The ESI negative mode for peak 5 gave a precursor ion peak at m/z 447.2246 [M-H]⁻ and a molecular formula C₂₁H₃₆O₁₀ was generated through MassLynx with a normalised iFit value of 0.050 (**supplementary data 17**). The compound was identified as acuminoside (**33**) (6-O- β -D-apiofuranosyl- β -D-glucopyranoside) and the structure of the compound is shown in Figure 3.26. The MS/MS analysis (**supplementary data 18**) produced one fragment ion at m/z 315.1824 by the loss of apiofuranosyl and another fragment at m/z 161.0469 with the loss of glucopyranosyl. This diglycoside is found in grapes [62] and does not have any previous scientific study related to Alzheimer's disease.

The ESI negative mode for peak 6 gave a precursor molecular ion peak at m/z 449.2383[M-H]⁻ with retention time of 9.80 min. Molecular formula C₂₁H₃₈O₁₀ was generated through MassLynx with a normalised iFit value of 0.000 (**supplementary data 19**). The compound was tentatively identified the eudesmane-type-sesquiterpene glycoside, dictamnocide D (**34**) and the structure is shown in Figure 3.26. The MS/MS spectrum (**supplementary data 20**) yielded a fragment ion at m/z 161.0472 with the loss of glycoside molecule and another fragment ion at m/z 101.0252.

The compound was previously isolated from the root bark of a Chinese medicinal plant *Dictamnus dasycarpus* Turcz [69]. The compound does not have any previous reports with any direct relation to Alzheimer's disease, but the *Dictamnus* species were found to exhibit neuroprotective activities [70].

The ESI negative mode for peak 7 gave a pseudomolecular ion peak at 987.5164 [M+HCOO]⁻ and a molecular formula C₄₈H₇₈O₁₈ was generated through MassLynx with a normalised iFit value of 0.050 (**supplementary data 21**). The compound was identified as a saikosaponin derivative 2''-O-β-D-glucopyranosylsaikosaponin B2 (**35**) and its structure is shown in Figure 3.26. The fragment ions (**supplementary data 22**) were produced at *m/z* 471.3472 [M-(2×162)-146]⁻ with the successive loss of sugar moieties including two glucopyranosyl and one fucopyranosyl unit [59].

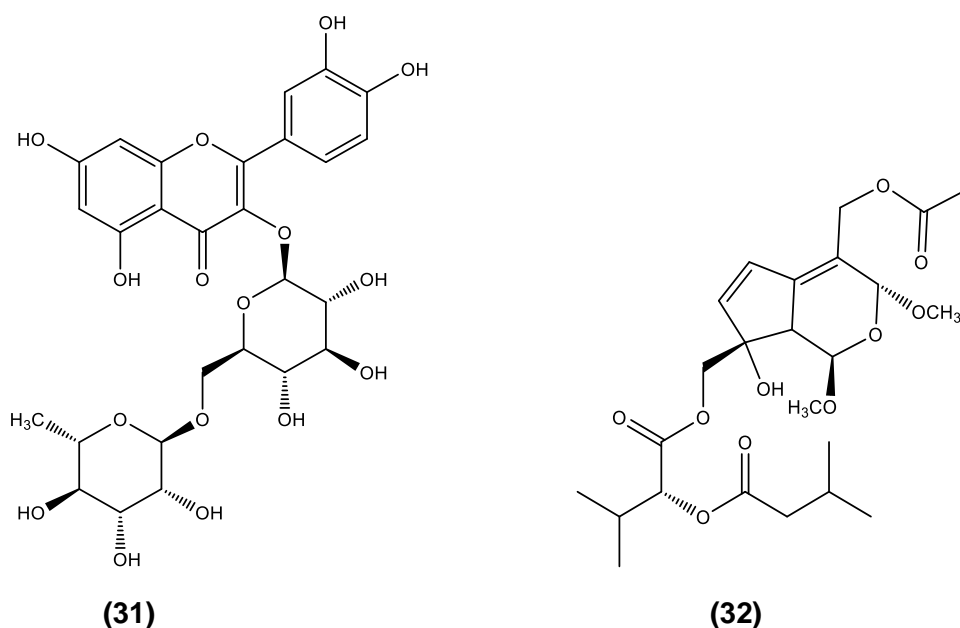
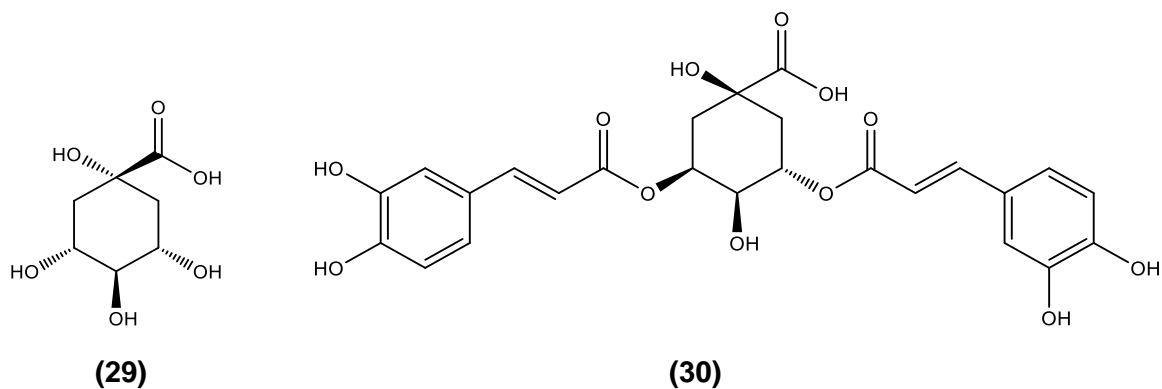
The compound was isolated from *Bupleurum longicaule* and from the roots of *Bupleurum wenchuanense* [59]. There is no previous study reported for this compound related to Alzheimer's disease. However there is some other saikosaponins of similar structure such as saikosaponin C which shown its effect on Alzheimer's disease by targeting amyloid β as well as tau protein [71]. Thus 2''-O-β-D-glucopyranosylsaikosaponin B2 could be the active ingredient in the present study.

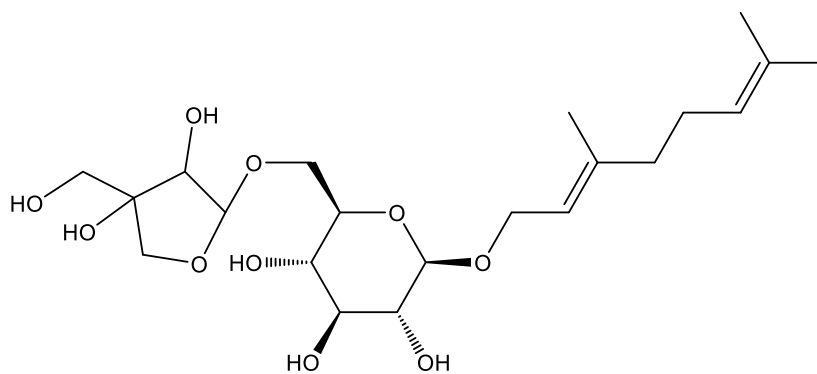
The ESI negative mode for peak 8 gave a pseudomolecular ion peak at 1029.5261[M+HCOO]⁻ with retention time 10.92. The molecular formula C₅₀H₇₉O₁₉ was generated through MassLynx with a normalised iFit value of 0.001 (**supplementary data 23**). The compound was identified as clinoposaponin C (**36**) and the structure of the compound is shown in Figure 3.26. The fragment ions (**supplementary data 24**) were produced at *m/z* 513.1736 [M-(2×162)-146]⁻ with the successive loss of sugar moieties including two glucopyranosyl and one fucopyranosyl. Another fragment ion was observed at *m/z* 471.3485 with the loss of the aglycone moiety.

The compound showed moderate cytotoxic activities with IC₅₀ values between 4.1 and 19.7 μM when it was tested against Hela, HCT-8, AGS, and MCF-7 human cancer cell lines [60]. The compound has not been reported for the treatment of Alzheimer's disease or other nervous disorders.

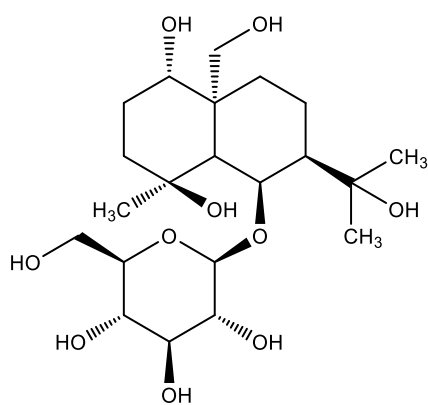
The ESI negative mode for peak 4 gave a pseudomolecular ion peak at 1029.5261 [M+HCOO]⁻ with retention time of 11.38 min. Molecular formula C₄₂H₆₆O₁₄ was generated through MassLynx with a normalised iFit value of 0.000 (**supplementary data 25**). The compound was tentatively identified as oleanane type triterpenoid saponin, spinasaponin A (**37**) and its structure is shown in Figure 3.26. The MS/MS spectrum (**supplementary data 26**) yielded a fragment ion at m/z 631.385 [M-H-glucose]⁻ corresponding to the loss of a glucose unit (162 amu).

Oleanane type triterpenoid saponins were found to possess antitumour, antiulcer, antiviral, antihyperlipidemic, antiinflammatory and other biological activities due to the presence of sachharide moities [72].

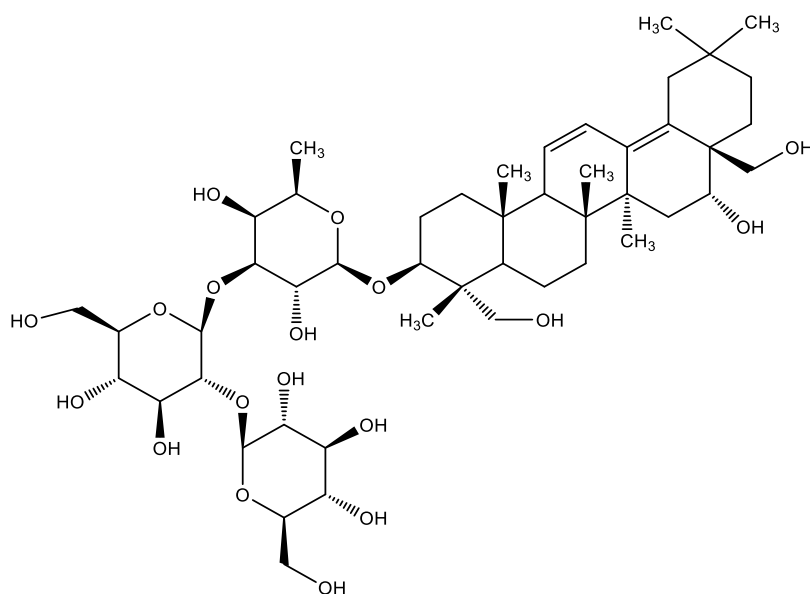




(33)



(34)



(35)

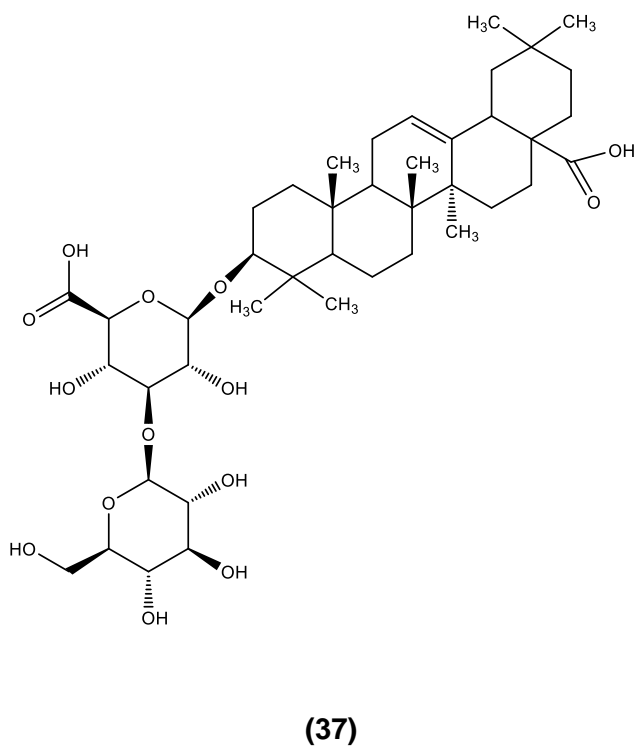
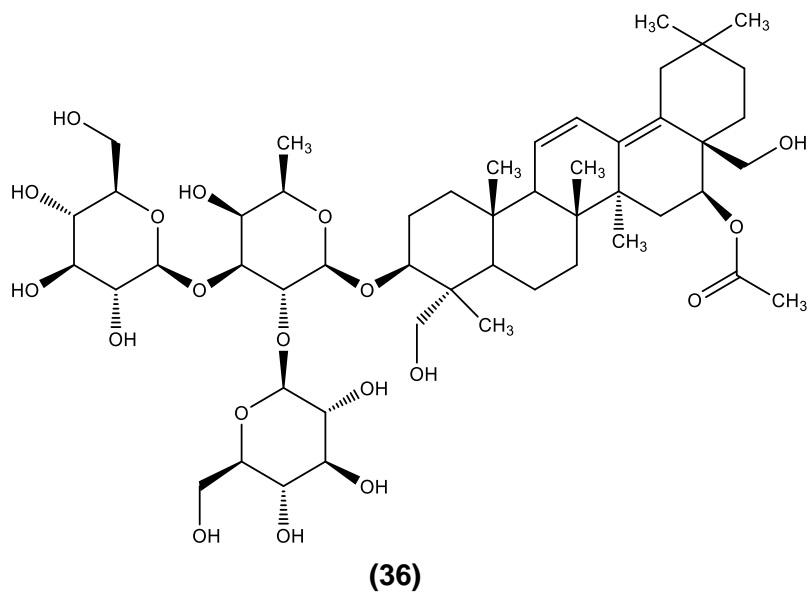


Figure 3.25: Chemical structures of the compounds identified from *C. paniculata* leaf extract. Peak 1: quinic acid (**29**), peak 2: 3,5- dicaffeoylquinic acid (**30**), peak 3: rutin (**31**), peak 4: valeriananoid E (**32**), peak 5: acuminoside (**33**), peak 6: dictamnocide D (**34**), peak 7: 2''-O- β -D glucopyranosylsaikosaponin B2 (**35**), peak 8: clinosaponin C (**36**), peak 9: spinasaponin A (**37**)

3.5.5. The effect of *H. arborescens* on A β 42 production

The stem extract of *H. arborescens* significantly reduced A β 42 production only at high test concentrations as shown in Figure 3.27. The A β 42 levels were reduced by $75.78 \pm 0.1\%$ at 50 $\mu\text{g/ml}$ and by $31.37 \pm 0.2\%$ at 25 $\mu\text{g/ml}$, while no reduction was seen at 5 and 10 $\mu\text{g/ml}$.

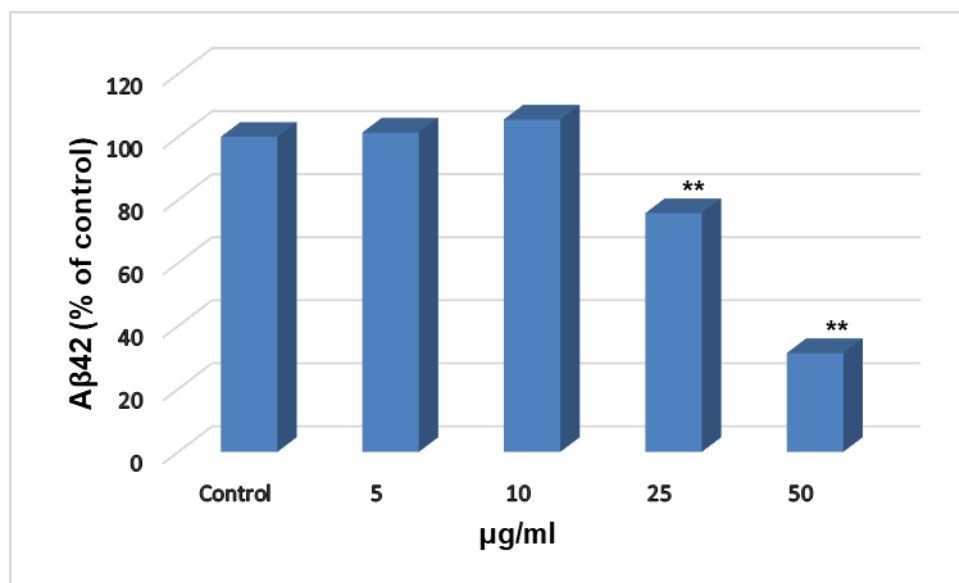


Figure 3.26: Change in the levels of A β 42. Cells were incubated with indicated concentrations of stem extract of *H. arborescens* for 8 h, the levels of A β 42 were measured from the conditioned media by using specific ELISA methods. The levels of A β 42 (n = 4) were decreased by the extract on at 25 and 50 $\mu\text{g/ml}$. **, P<0.01; ***, P<0.001.

H. arborescens has no previous reported scientific studies related to any central nervous system disorder. This plant species is for the first time reported in the present study for the reduction of A β 42 and supported the fact that plant is traditionally used for the nervous disorders. These results indicated that the compounds present in *H. arborescens* could be the active ingredients for the reduction of A β 42 and can be developed into a potent drug for the treatment of Alzheimer's disease.

3.5.6. Chemical characterization of DCM:MeOH stem extract of *H. arborescens*

The DCM:MeOH stem extract of *H. arborescens* was analysed on UPLC-QTOF-MS to identify the major peaks and establish whether if these could be the active ingredients responsible for the reduction of A β 42. The ES negative mode was not further analysed because of the poor ionisation. In ESI negative mode, the peaks were mainly distributed within 5.90 to 12.04 min range of the 18 min chromatogram (Figure 3.28).

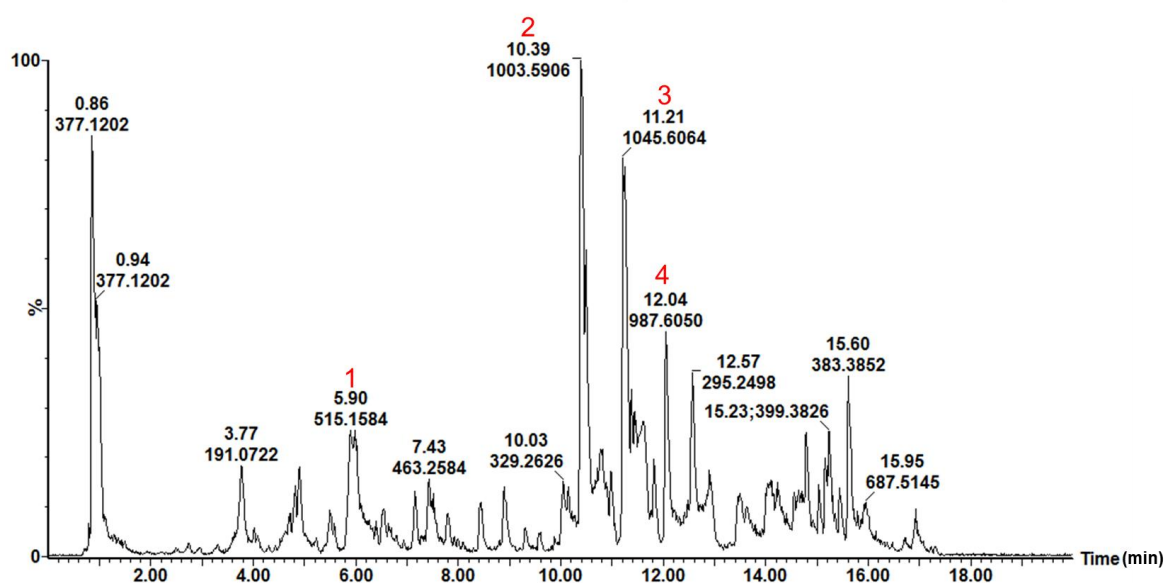


Figure 3.27: ESI negative mode BPI chromatogram of *H. arborescens* stem extract.

This indicated the general elution region for polar to medium polar compounds. Using ESI negative mode, mass fragmentation data and online search databases, a total of four compounds were tentatively identified which included one polyphenol and three triterpenoid saponins (Table 3.4), peak 1 identified as cynarin (1,5-dicaffeoylquinic acid) (**38**) with m/z 515.1584 [M-H]⁻; peak 2 identified as alternoside IX (**39**) with m/z 1003.5906 [M-HCOO]⁻; peak 3 identified as alternoside I (**40**) with m/z 1045.6064 [M-HCOO]⁻ and peak 4 identified as saikogenin B4 (**41**) with m/z 987.6050 [M-HCOO]⁻.

Table 3.4: Tentative identification of compounds obtained from ESI-MS negative mode of DCM:MeOH extract of *H. arborescens* (stem)

Peak	RT (min)	Acquired [M-H] ⁻ m/z	Formula of Possible structure	Theoretical [M-H] ⁻ m/z	Possible structure	Mass error (ppm)	MS/MS Data (Fragments)	Ref.
1	5.90	515.1192	C ₂₅ H ₂₄ O ₁₂	515.1190	Cynarin	0.4	353.1182, 191.0722, 135.0551, 179.0321	[73]
2	10.39	957.5052	C ₄₈ H ₇₈ O ₁₉	957.5059	alternoside IX	-0.2	795.5234, 633.4459	[74]
3	11.21	999.5168	C ₅₀ H ₈₀ O ₂₀	999.5165	alternoside I	0.3	837.5388, 675.4644	[74]
4	12.04	941.5130	C ₄₈ H ₇₈ O ₁₈	941.5110	saikogenin B ₄	2.1	151.0066, 121.0306, 107.0152	[75]

ESI negative mode of peak 1 gave a molecular peak ion at m/z 515.1584 [M-H]⁻ with a retention time 5.90 min. The molecular formula C₂₅H₂₄O₁₂ was generated through MassLynx with a normalised iFit value of 0.000 (**supplementary data 27**). The compound was identified as polyphenol named cynarin (1,5- dicaffeoylquinic acid) (**38**) and the structure of the compound is shown in Figure 3.26.

This compound was also identified through integrated analytical workflow using UPLC-QTOF-MS and UNIFI informatics platform. In UNIFI, the iFit confidence is 100% and exact mass error is -0.02 mDa (**supplementary data 28**). The identified compound met the match criteria with the incorporated traditional medicinal library. The compound is provided with detailed information including compound name with retention time, molecular structure and fragment ion in the component plot (**supplementary data 29**). The fragmentation patterns obtained

from Masslynx and UNIFI were compared and found to be similar and both gave same molecular formula.

The fragments were produced at m/z 353.1182, m/z 191.0722, m/z 179.0321 and m/z 135.0551 (**supplementary data 30**). The same fragmentation pattern was obtained from a previous study which supported its identification as cynarin (**38**) [73]. 1,5-dicaffeoylquinic acid was found to partially reduce the A β cytotoxicity in peptide treated cultures (50 μ M) in SH-SY5Y cell model system. It was also normalized A β peptide-induced tau gene expressions [76].

The ESI negative mode of peak 2 gave a pseudomolecular peak ion at 1003.5906 [M+HCOO]⁻ with a retention time 10.39 min. MassLynx was used to generate a molecular formula C₄₈H₇₈O₁₉ with a normalised iFit value of 0.050 (**supplementary data 31**). The compound was tentatively identified as oleanene-type of compound called alternoside IX (**39**) and the structure of the compound is shown in Figure 3.29.

The fragment ions (**supplementary data 32**) were produced at m/z 795.5234 [M-162]⁻ and m/z 633.4459 [M-162-162]⁻ with the successive loss of two sugar moieties, glucopyranoside and rhamnopyranoside (-162 amu). There is no scientific study reported for this compound related to Alzheimer's disease and other mental disorders.

The ESI negative mode for peak 3 gave a pseudomolecular peak ion at 1045.6064 [M+HCOO]⁻ with a retention time 11.21 minutes. The molecular formula C₅₀H₈₀O₂₀ was generated through MassLynx with a normalised iFit value of 0.000 (**supplementary data 33**). The compound was identified as oleanene-type compound named alternoside I (**40**) and its structure is shown in Figure 3.29.

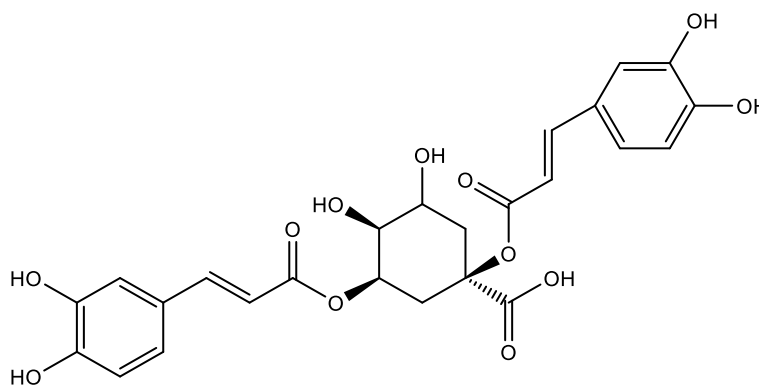
The fragment ions (**supplementary data 34**) were produced at m/z 837.5388 [M-162]⁻ and m/z 675.4644 [M-162-162]⁻ with the successive loss of two sugar moieties, glucopyranoside and rhamnopyranoside. The successive loss of the sugar moieties showed the characteristic

fragmentation patterns of saponins. The compound was not reported previously for the reduction of A β 42 or other bioassays related to Alzheimer's disease.

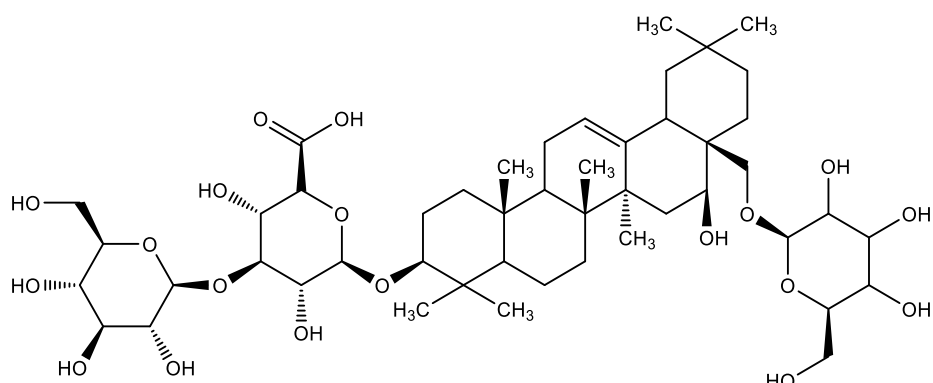
The molecular formula and fragmentation pattern found from the previous studies also supported the identification of both oleanene-type triterpenoids, alternoside IX **(39)** and alternoside I **(40)** [74].

The ESI negative mode for peak 4 gave a pseudomolecular ion peak at 987.6050 [M+HCOO]⁻ with a retention time 12.04 min. The molecular formula C₄₈H₇₈O₁₈ was generated through MassLynx with a normalised iFit value of 0.000 (**supplementary data 35**). The compound was identified as oleanene-type compound named saikogenin B₄ **(41)**, an oleanolic acid triterpenoid saponin with a structure shown in Figure 3.29.

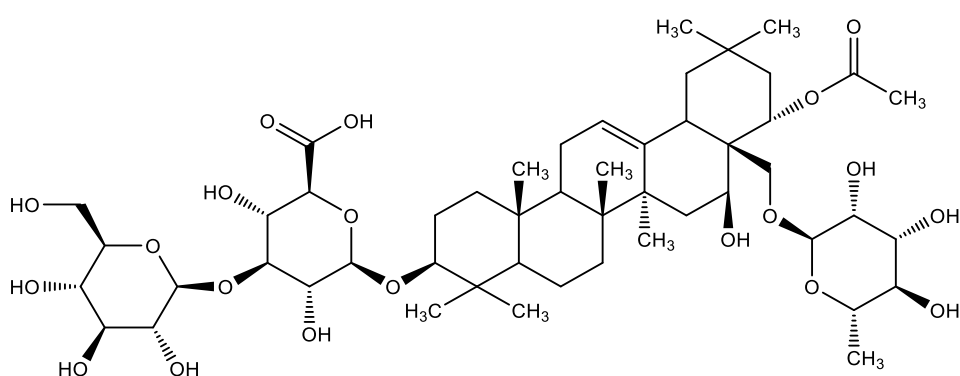
The fragment ions (**supplementary data 36**) were produced at *m/z* 779.4558 [M-162]⁻ and *m/z* 617.4025 [M-162-162]⁻ with the successive loss of two hexose moieties. The compound was previously isolated from the root extract of *Bupleurum chinese*, the plant which was reported for its potential as a mild sedative, antidepressant and anti-inflammatory. Furthermore, the oleanane triterpenoids including saikosapogenin B₄ isolated from *Bupleurum chinese* were found to exhibit neuroprotective properties and have the potential to be developed for the treatment of Parkinson's disease, Alzheimer's disease and depression [77]. Based on this, the compound is highly probable to be an active ingredient for the reduction of A β 42 in the stem of extract of the plant.



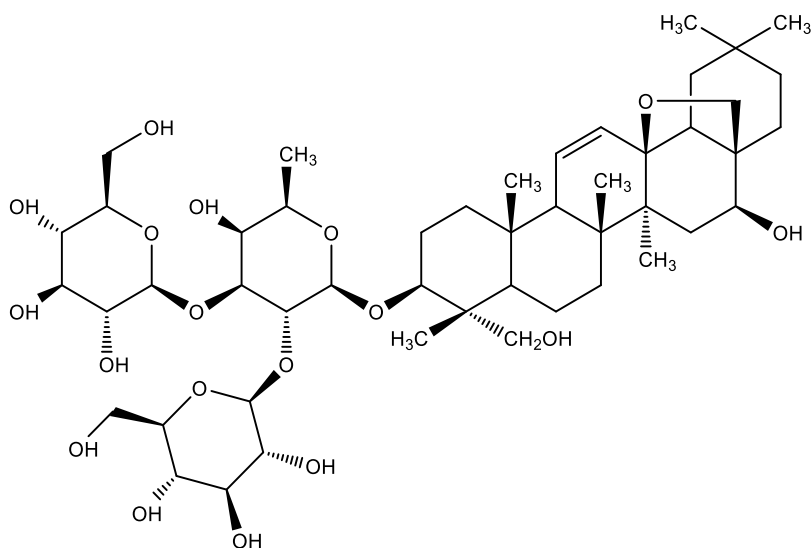
(38)



(39)



(40)



(41)

Figure 3.28: Chemical structures of the compounds identified from *H. arborescens* stem extract. Peak 1: cynarin (**38**) Peak 2: alternoside IX (**39**), peak 3: alternoside I (**40**), saikogenin B₄ (**41**)

The main class of compounds identified in *H. arborescens* stem extract using UPLC MS QTOF was saponins while a limited number of compounds have been identified in *C. paniculata* leaf extract. In previous studies it was found that most saponins contained an aglycone with molecular mass 448, 456, 458, 472 or 474 Da with the presence of two to five sugar moieties [78]. The ESI MS/MS negative mode of these saponins showed similar of fragment pattern with the successive loss of several sugar moieties such as hexose, pentose or deoxyhexose with m/z 162, 132 or 146 respectively [79]. This characteristic fragmentation pattern of triterpenoid saponins obtained from previous studies supported the tentative identification of saponins from leaf extract of *C. paniculata* and stem extract of *H. arborescens*.

Saponins were also found to exhibit neuroprotective activities in various bioassays. In an *in vitro* study using the Thioflavin T method, three acylated oleanane-type triterpene oligoglycosides, floraassamsaponins III, IV, and VII were found to significantly inhibit the A β 42 aggregation with an inhibition percentage $73.4 \pm 8.0 \%$, $p < 0.01$; $68.8 \pm 8.4 \%$, $p < 0.01$; and $56.9 \pm 13.2 \%$, $p < 0.01$ at 100 μ M, respectively as compared to a positive control, morin which had an inhibition percentage $54.0 \pm 11.2 \%$, $p < 0.01$ at 100 μ M [80].

Some other saponins including panax notoginseng saponins isolated from *Panax notoginseng*, ginsenosides from *Panax ginseng* C. A. Mey. and gypenosides isolated from *Gynostemma pentaphyllum* were found to exhibit potential anti-Alzheimer's activity [81].

These previously reported bioassays support the fact that the saponins identified from *C. paniculata* leaf extract and *H. arborescens* stem extract are in all likelihood the active compounds responsible for the reduction of A β 42 and could lead to the development of new anti-Alzheimer's therapeutics.

3.6. Conclusion

The three plants, *S. brachypetala* leaf extract, *C. paniculata* leaf extract and *H. arborescens* stem extract were found to significantly reduce the levels of A β 42 either at higher

concentrations or in a dose dependant manner. This provided some evidence which substantiates the traditional use of the plant species for treating Alzheimer's disease and therefore warranted their further research. The plant extracts were further analysed on UPLC-QTOF-MS to identify the major compounds which could be responsible for the decreased levels of A β 42. ESI negative mode was used to identify the active ingredients as the ionization was poor for the positive mode for further analysis.

S. bracypetala leaf extract showed significant reduction in A β 42 production only at higher concentrations i.e. $29.69 \pm 4\%$ at 50 $\mu\text{g/ml}$ and by $62.17 \pm 7\%$ at 100 $\mu\text{g/ml}$.

Using UPLC-QTOF-MS, a total of five compounds were tentatively identified including four flavonoids, myricetin-3-O-alpha-L-rhamnopyranoside (**25**), isoquercetin (**26**), quercetin-3-O-rhamnoside (**27**), quercetin (**28**). Isoquercetin was confirmed by using a pure standard on UPLC-QTOF-MS. As these compounds were previously studied for their potential to treating Alzheimer's disease through A β reduction in different bioassays, in all likelihood the compounds would be responsible for the A β 42 reduction in *S. bracypetala*.

The extract of leaves of *C. paniculata* significantly reduced the A β 42 levels in a dose dependent manner by $29.03 \pm 1.4\%$, $42.40 \pm 1.7\%$, $47.50 \pm 5.8\%$ and $61.14 \pm 9.3\%$ at concentrations 10, 25, 50 and 100 $\mu\text{g/ml}$, respectively. This is the first report for the reduction of A β 42 and thus provides scientific evidence which substantiates its traditional use related to treatment of nervous disorder treatments.

Using ESI negative mode of UPLC-QTOF-MS, a total of nine compounds were tentatively identified as quinic acid (**29**), 3,5-dicaffeoylquinic acid (**30**), rutin (**31**), valeriananoid E (**32**), acuminoside (**33**), dictamnoid D (**34**), 2''-O- β -D-glucopyranosylsaikosaponin B2 (**35**), clinoposaponin C (**36**), spinasaponin C (**37**). Rutin was confirmed by using pure standard on UPLC-QTOF-MS. From these identified compounds, rutin was the most researched compound for Alzheimer's disease in previous studies. 3,5-dicaffeoylquinic acid was previously reported for AChE inhibition. The remaining identified compounds from this extract were not reported to be studied for treating Alzheimer's disease either through A β 42 reduction

or by AChE inhibition. In the present study, rutin or 3,5-dicaffeoylquinic acid could be the active ingredients in *C. paniculata* leaf extract.

The stem extract of *H. arborescens* significantly reduced A β 42 levels only at high test concentrations by $75.78 \pm 0.1\%$ at 50 $\mu\text{g/ml}$ and by $31.37 \pm 0.2\%$ at 25 $\mu\text{g/ml}$.

Using ESI negative mode, a total of four compounds were tentatively identified which included one polyphenol, cynarin (1,5-dicaffeoylquinic acid) (**38**) and three triterpenoid saponin alternoside IX (**39**), alternoside I (**40**) and saikogenin B₄ (**41**). Cynarin was previously reported for the reduction of A β cytotoxicity while the other identified compounds, triterpenoid saponins were not examined for Alzheimer's disease. Thus in this study there is a possibility that cynarin could be the active ingredient from *H. arborescens* leaf extract for the reduction of A β 42.

From the present study it can be concluded that UPLC-QTOF-MS is an effective and quick technique to identify major compounds in complex plant extracts and based on published data of such compounds, the active ingredients can be established. The main class of compound identified from these plants were flavonoids, quinic acid derivatives and saponins. All these classes of compounds were previously scientifically tested for their neuroprotective effects in various bioassays and animal models and gave many positive results thereby no further development of the complex plant extracts was undertaken.

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Chapter 4

Isolation and characterization of compounds from *Xysmalobium undulatum* targeting A β 42 reduction for the treatment of Alzheimer's disease

4.1. Introduction

4.1.1. Botany and geographical distribution of *Xysmalobium undulatum*

Xysmalobium undulatum (Apocynaceae) is a robust annual herb, 0.5 - 2.0 m tall. It sprouts annually in spring and dies in winter. The leaves are large (80 - 270 × 10 - 75 cm), hairy, heart shaped and almost stalkless; the branches are thick, erect and hairy and the roots are fleshy, it has a white inside and latex present in all parts. Many flowered, stalked inflorescences develop in the axils of the leaves. The flowers are creamish green to yellow and produced mainly in December and the fruits are usually large, inflated ovoid follicle, pale green, short hairy and many are seeded [1], [2].

The common names of *X. undulatum* are milkbush, milkwort, Uzara, wild cotton and wave-leaved *Xysmalobium* while in Afrikaans it is known as bitterhout [2]. These plants grow mainly in grasslands and moist places. The genus *Xysmalobium* comprises of about 40 species in Africa and of the 40 species, about 24 species are found in southern Africa. The plant is extensively harvested for its medicinal use in South Africa.

The plant is also widely distributed in Kenya, Tanzania, Malawi, Zambia, Angola, Namibia, Botswana, Zimbabwe, Mozambique, Lesotho, Swaziland and found in all the provinces of South Africa [3].



Figure 4.1: Picture showing *X. undulatum* plant (image taken by Anuradha Thakur at the time of collection).

4.1.2. Traditional medicinal uses

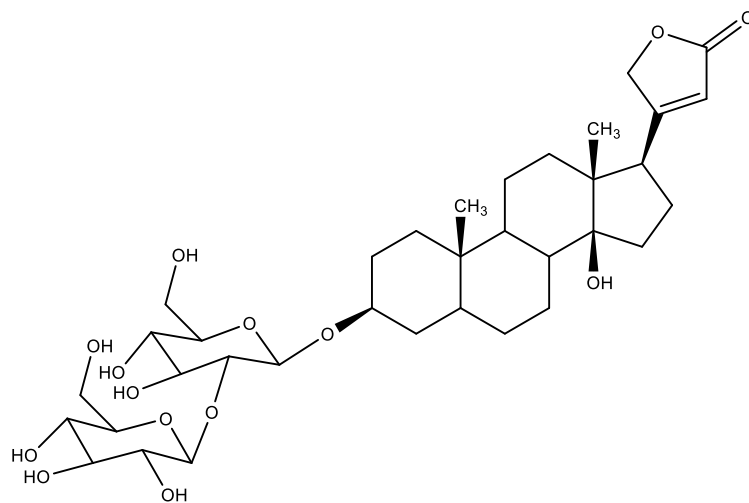
Traditionally the plant is used in Botswana for headache and abdominal pains [4] while it has also been recorded for the treatment of diarrhoea, dysentery, dysmenorrhoea, stomach cramps, wounds and indigestion [5]. The root extracts are used to treat acute diarrhoea and menstrual cramps since early 1900s in Europe and have been marketed as 'uzarae radix' [2]. The Xhosa people sometimes mixed *X. undulatum* with *Pachycarpus shinzianus* (Schltr.) N.E.Br. to treat intestinal problems as they have the same medicinal properties. The Tswana people chew a piece of the root as an antidote to food poisoning, the Mpondo people use powdered root or root decoction to treat dysentery and the Nama people ingest root as stomach carminative and for the treatment of diarrhoea and lumpy skin disease of cattle [2]. In Kenya a root decoction is taken to treat headache, in Zambia it is taken for the treatment of malaria,

typhoid and other fevers and in Zimbabwe the pulverized root is eaten with porridge as an aphrodisiac. The latex and plant decoction is used as a remedy for wounds, sores and snakebites. The Sotho people in South Africa eat the young leaves mixed in porridge [3]. The tuber infusions of this plant are also used for the treatment of diarrhoea and headache [6]. It is also used for the treatment of skin diseases, infected sores, wounds and urinary tract infections [7].

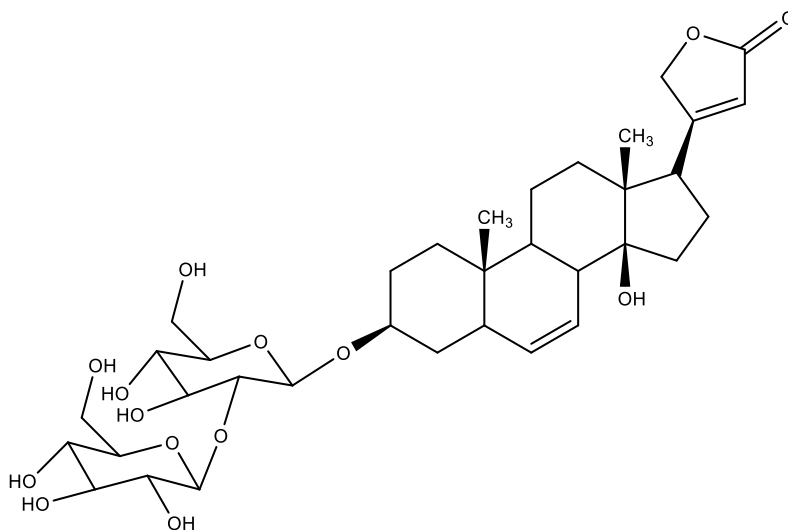
4.1.3. Phytochemistry

The phytochemical investigations showed that the main class of compounds in *X. undulatum* is cardenolide glycosides. The compounds isolated from *X. undulatum* root are uzarin, xysmalorin, uzarigenin, xysmalogenin, allouzarin, alloxysmalorin, allouzarigenin, alloxysmalogenin, ascleposide, corogluaucigenin, corogluaucigenin-3-O-glucoside, desglucouzarin, smalogenin, desglucoxysmalorin, uzaroside, pregnenolone and β -sitosterol [2].

Four major cardenolide glycosides were isolated named uzarin, allouzarin, xysmalorin and alloxysmalorin from the leaf extract [8] and the structures of these compounds are shown in Figure 4.2. A biglucosyl moiety was also identified through the presence of anomeric protons in the ^1H NMR while their β configuration was confirmed by the large coupling constants of the two anomeric protons (7.7Hz). The COSY spectrum in $\text{DMSO-}d_6$ established 1 \rightarrow 2, glycosidic linkage, i.e. glucopyranosyl-(1 \rightarrow 2)-glucopyranose.



(42)



(43)

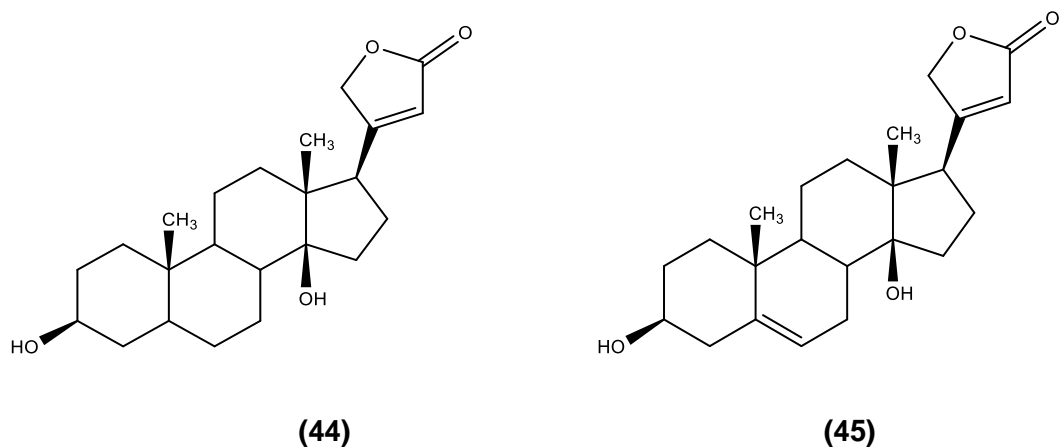


Figure 4.2: Structure of main compounds isolated from *X. undulatum*, uzarin (42), xysmalorin (43), uzarigenin (44) and xysmalogenin (45)

4.1.4. Previously reported biological assaying

Previous studies reported that *X. undulatum* possess a strong affinity towards serotonin uptake transport protein which plays an important role in the symptoms of depression and Alzheimer's diseases [9]. Another study has suggested that the ethyl acetate root extract of *X. undulatum* exhibited some acetylcholinesterase inhibitory activity by displaying an IC_{50} value of 0.0005 ± 0.000 mg/ml [10]. Previous findings showed that *X. undulatum* extracts exhibited antidepressant-like *in vitro* and *in vivo* effects [11]. The ethanol root extract of the plant exhibited antiplasmodial activity with an IC_{50} value $6\mu\text{g/ml}$. Uzara whole plant extract exhibited antimicrobial activity with a recorded MIC value of 3.125 mg/ml against four bacteria *Bacillus subtilis* (ATCC6051), *Escherichia coli* (ATCC11775), *Klebsiella pneumoniae* (ATCC13883) and *Staphylococcus aureus* (ATCC12600) using neomycin as a reference standard as well as *Candida albicans* (ATCC10231) with amphotericin B as a reference standard [2]. In an *in vitro* study for acetylcholinesterase (AChE) inhibition performed using Ellman's colourimetric method, the ethyl acetate root extract of *X. undulatum* exhibited the AChE inhibition with an IC_{50} value of 0.5 ng/ml as compared to control galanthamine with an IC_{50} value of 0.053 ng/ml [2]. These values are significantly lower for galanthamine as compared to the other $A\beta 42$ inhibition tests using galanthamine as control. As an example in one of the previous studies,

galanthamine was reported to inhibit amyloid aggregations at all the concentrations tested from 25-1000 μM [12]. In another study it was found that release of $\text{A}\beta_{42}$ was decreased by 37% with 10 μM galanthamine in SH-SY5Y cells [13].

4.1.5. History of commercialisation of *X. undulatum* (Uzara)

Carl Peter Thunberg, described Uzara under the synonymous name *Asclepias undulata* in European scientific literature [14]. The plant was a part of Thunberg herbarium and was said to be used as remedy for the colic [15]. Further Smith described the plant under a synonym name *Xysmalobium lapathifolium* and mentioned the uses of root as tonic and as a remedy for healing the wounds [16].

H.W.A Hopf from Melsungen, was the first person who brought Uzara roots to Europe. While travelling through South Africa, he suffered from severe dysentery and obtained Uzara roots from a traditional healer to treat dysentery. In this way he became familiar with Uzara roots and brought the roots with him when he returned to Melsungen, Germany, where he left the root specimen to the pharmacological institute for further investigations. At the institute, August Gurber (1864-1937) isolated three unidentified compounds from the roots during his investigation. Following further research Uzara was commercialized and introduced into German pharmaceutical market in 1909. It is marketed as a drug under the name "Uzara" for the treatment of acute diarrhoea [17].

4.1.6. Pharmacological effects of active ingredients from *X. undulatum*

The cardenolide glycosides, major compounds from *X. undulatum* are also known as cardiotonic steroids, a class of naturally occurring drugs which has both beneficial as well as toxic effects on the heart. Although the toxicity remains a serious problem, these compounds in lower doses are widely used in the treatment of congestive heart failure and cardiac arrhythmia. They have positive inotropic effect on myocardial contraction and atrioventricular conduction. The compounds increase the tone, excitability and contractility of the cardiac

muscle which helps the proper functioning of the weakened heart [18]. The cardenolide glycosides act by inhibiting the plasma membrane Na⁺, K⁺ ATPase thereby making them useful in the drugs for the treatment of heart problems [19]. The varying structures of cardenolides does not affect their binding at Na⁺, K⁺ ATPase [20]. They also exhibit diuretic activity due to increased renal circulation, which relieves edema often associated with heart failure. However in large doses they cause cardiac arrest in systole, and can also lead to death [18]. The major and best known cardenolide glycoside isolated from *X. undulatum* is uzarin. It inhibits intestinal motility through affecting the visceral smooth muscles and consequently proven to have antidiarrhoeal activity [18].

4.2. Materials and Methods

4.2.1. Collection and extraction of plant material

Fresh leaf material from *X. undulatum* were collected from the experimental farm of University of Pretoria, South Africa with the help of the curator, Jason Samuels. Plant identification was done at the H.G.W.J. Schweickerdt Plant Herbarium at University of Pretoria where a voucher specimen was kept for further reference. The collected leaves were oven dried at 60°C and ground to a fine coarse powder. The powdered leaves (300 g) were extracted repeatedly (3 times) for 8 hours with 2 l of DCM:MeOH (1:1) at room temperature using a magnetic stirrer. After filtration, the solvent was concentrated using a rotary vacuum evaporator at 50-60°C and then dried under high vacuum for 24 h and the resulting extract stored in the cold room (Figure 4.3).

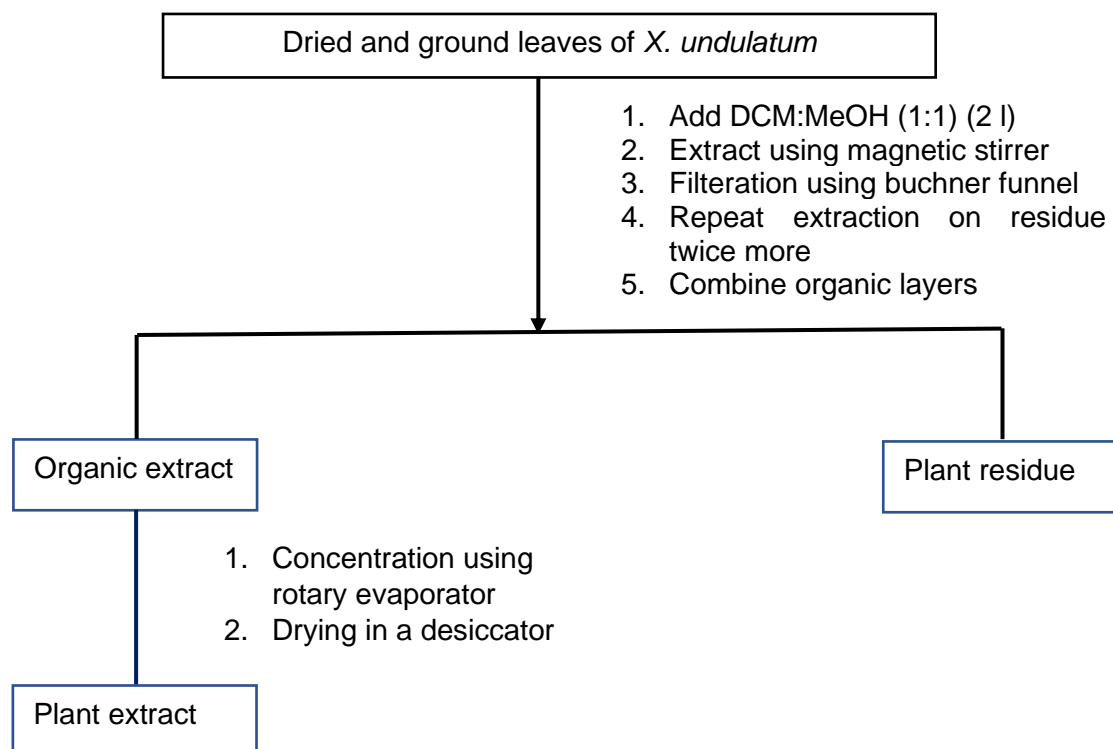


Figure 4.3: Flow diagram for the extraction of dried leaves of *X. undulatum*

4.2.2. Fractionation using silica gel column chromatography

The crude extract was subjected to silica gel (0.063-0.2 mm/70-230 mesh) gravity column chromatography using a glass silica column of 107 cm length and 7cm internal diameter. A stepwise gradient of DCM:Hexane (50:50 to 80:20), DCM (100%) and DCM:MeOH (95:5 to 70:30) was used for elution with the solvent gradient flow rate of 2 ml/min. The fractions were monitored using TLC silica gel plates and visualised under a UV lamp at 254 nm followed by staining using the vanilin stain. After visualisation, the similar fractions were combined based on R_f values. These fractions were then screened for A β 42 reduction activity. The active fractions were further purified using semi-preparative HPLC-MS.

4.2.3. Isolation and purification of compounds using preparative HPLC-MS

The active fractions were purified using preparative HPLC-MS on a Waters chromatographic system with Waters PDA (2998) and MS detector (Waters, Milford, MA, USA). Each of the active fractions (70mg) was diluted to 1 ml with methanol. A X-Bridge preparative C18 column (19×250 mm, i.d., 5µm particle size, Waters) was used for the separation while the mobile phase consisted of 0.1% ammonia in water (solvent A) and acetonitrile (solvent B) at a flow rate of 20 ml/min. Gradient elution was applied as follows: 95% A : 5% B for 1 min, 50% A : 50% B (1:00-8:00 min), 15% A : 85% B (8:00-13:00 min), 95% A : 5% B (13:00-13:20 min). Injection volume was 150 µl. Data were collected using MassLynx4.1™ (Waters, USA) software. The preparative HPLC system was interfaced with QDa mass spectrometer (Waters, USA) and the negative ion mode selected. The probe temperature and source temperature were set at 600°C and 120°C, respectively. The capillary and cone voltage were set to 800 and 10 V, respectively. Data was collected between 100 and 650 *m/z*. The pure compounds were collected using a fraction collector and subsequently combined and concentrated using the speed vacuum. The compounds were later analyzed by UPLC-MS and their structures are confirmed by NMR.

4.2.4. Chemical profiling using UPLC-QTOF-MS

The crude extracts and fractions were analyzed using a Waters Acquity UPLC system (Waters corp., MAUSA), equipped with a binary solvent delivery system and an autosampler. The instrument was centrally operated by MassLynx 4.1 software (Waters Inc., Milford, Massachusetts, USA) for data acquisition. The separation was achieved on Waters BEH C18 1.7 µm particle size (2.1 mm × 100 mm). The mobile phase consisted of solvent A: water with 0.1% formic acid and solvent B: methanol with 0.1% formic acid. The gradient elution was optimized as follows: 3% B (0-0.1 min), 100% B (0.3-14.00 min), 100% B (14.00-16.00 min), 3% B (16.00-16.50 min), 3% B (16.50-20.00 min). The flow rate was 0.3 ml/min for the entire run, giving a total run time of 20 min and the injection volume was 5µl. The instrument was calibrated by direct infusion of 5 nM sodium formate solution at a flow rate of 10 µl/min over a

mass range of 50-1200 Da. The following MS source parameters were set for both positive and negative mode: source temperature 100 °C, sampling cone 15 V, extraction cone 4.0 V, desolvation temperature 400 °C, cone gas flow 10.0 l/h, desolvation gas flow 700 l/h, capillary 2.0 kV. Both positive and negative modes were obtained, however the results were analyzed using the negative mode as the higher intensity peaks were obtained from this mode. Compounds were tentatively identified based on their accurate mass generated from MassLynx V 4.1, iFit value, MS/MS fragmentations (product ions) and by the use of Metlin, Metfusion, Pubchem, Reaxys, Scifinder, Scopus, Chemspider, Chemical Entities of Biological Interest (ChEBI) and Massbank libraries.

4.2.5. NMR analysis

Structure elucidation of isolated compounds was carried out using NMR spectroscopy. ¹H and ¹³C NMR spectra were acquired on a Bruker Fourier 400 spectrometer (¹H at 400.21 MHz and ¹³C at 100.63 MHz) and 500 spectrometer (¹H at 500.10 MHz and ¹³C at 125.75 MHz). Chemical shifts were reported in ppm, referenced to residual solvent resonances (methanol-d₄ δ_H 3.31, δ_C 49.0 ppm).

4.2.6. Cell culture

For cell culture methodology, see section 2.2.3 of chapter 2.

4.2.7. sAPP α , sAPP β , A β peptide assay

For sAPP α , sAPP β , A β peptide assay, see section 3.4.4 of chapter 3.

4.2.8. Cell viability measurement

For cell viability measurement, see section 3.4.5 of chapter 3.

4.2.9. Statistical analysis

For statistical analysis, see section 2.2.5 of chapter 2.

4.3. Results and discussion

4.3.1. Collection and extraction of plant material

Fresh leaves (900 g) of *X. undulatum* was collected from the experimental farm of University of Pretoria. A voucher specimen (PRU 124301) was deposited at H.G.W.J. Schweickerdt Herbarium of University of Pretoria. After oven drying of the leaves, 200 g of dried material was obtained. The three times repeated extraction using DCM:MeOH of the dried leaves gave 17 grams of the extract in a 8.5% yield. The weight and percentage yield of the plant material (fresh and dried leaves) and plant extract are shown in the Table 4.1.

Table 4.1: Weight and percentage yield of crude extracts from the collected of *X. undulatum*

Sample	Extract	Weight (g)	% yield (w/w)
<i>X. undulatum</i> leaf material	Fresh plant material	900	
	Dried and ground plant material	200	22.2
	Dry DCM:MeOH extract	17	8.5

4.3.2. The effect of the leaf extract of *X. undulatum* on A β 42 production

The bioassaying was made possible with support of Korean Institute of Science and Technology (KIST). Special thanks to the strong collaboration of Dr Yoon Sun Chun, Ansun Park in Dr Hyun OK Yang's laboratory at KIST. I also acknowledge the work done by KIST in the training on running the assays and interpreting of data during my research visit to their laboratories. The effect of *X. undulatum* leaf extract on A β 42 production at 0.5, 1, 5, and 10 μ g/ml extract for 8 h, showed significant decrease in the level of A β 42 production in a dose-

dependent manner (Figure 4.4). The leaf extract significantly reduced the A β 42 production by $49.8 \pm 4.0\%$, $71.4 \pm 1.6\%$ and $72.9 \pm 1.1\%$ at concentrations 1, 5 and 10 $\mu\text{g/ml}$, respectively.

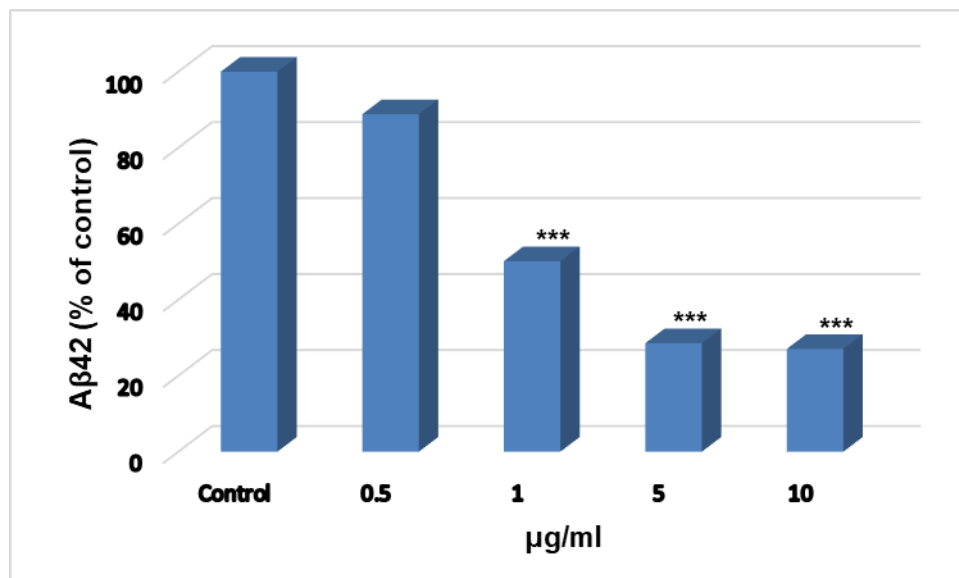


Figure 4.4: Change in the levels of A β 42. Cells were incubated with indicated concentrations of leaf extract of *X. undulatum* for 8 h, the levels of A β 42 were measured from the conditioned media by using specific ELISA methods. The levels of A β 42 ($n = 4$) were significantly decreased by the extract at concentrations 1, 5 and 10 $\mu\text{g/ml}$. **, $P < 0.01$; ***, $P < 0.001$.

The leaf extract of *X. undulatum* induced the decrease in cell viability to a small extent (Figure 4.5). The cell viability was decreased by $9.5 \pm 6\%$, $13.6 \pm 7.3\%$, $15.5 \pm 4.2\%$ and $22.0 \pm 2.8\%$ at the concentrations 0.5, 1, 5 and 10 $\mu\text{g/ml}$, respectively.

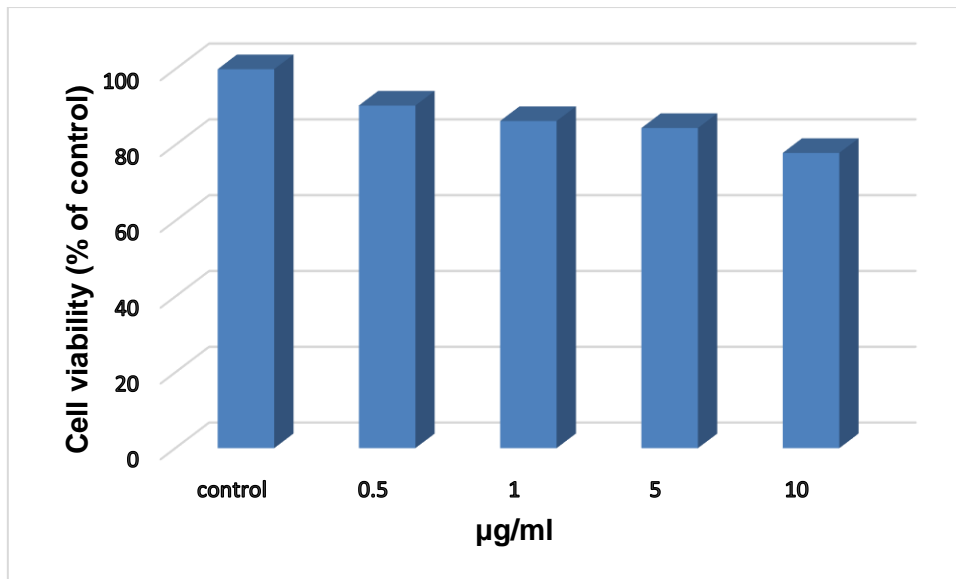


Figure 4.5: Cell viability measurements. Cells were incubated with indicated concentrations of extracts of leaves of *X. undulatum* for 8 h, and then incubated with EZ-Cytox solution for 1 h. The extract of leaves of *X. undulatum* induced the decrease of cell viability to small extent at all concentrations (0.5, 1, 5 and 10 µg/ml) (n = 6).

The secreted level of APP proteolytic products from the conditioned media using specific ELISA kits for A β 40, sAPP β -sw and sAPP α were then measured. Secreted level of A β 40 were decreased by $26.2 \pm 3.2\%$, $66.4 \pm 1.4\%$ and $73.3 \pm 1\%$ at 1, 5 and 10 µg/ml of extract, respectively (Figure 4.6).

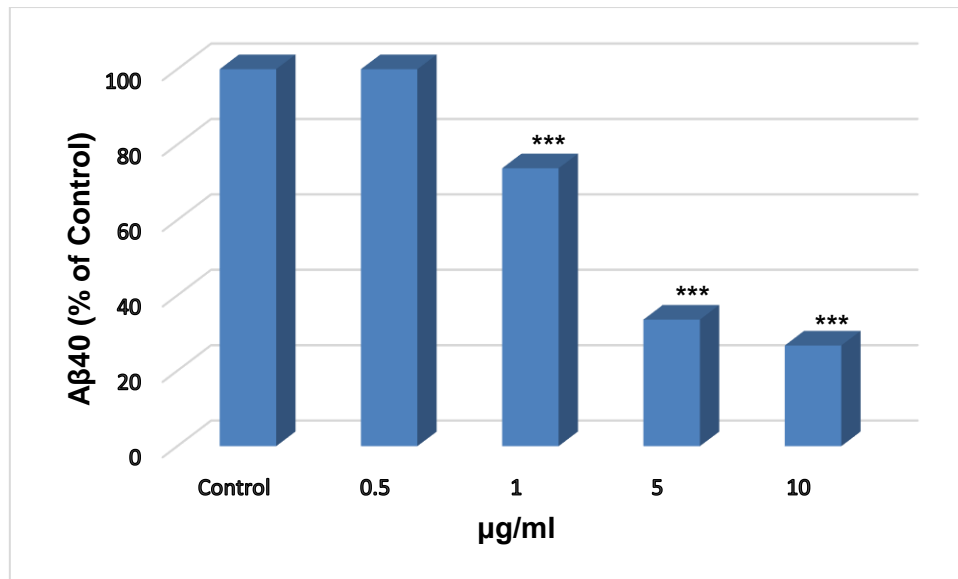


Figure 4.6: Change in the levels of Aβ40. Cells were incubated with indicated concentrations of leaf extract of *X. undulatum* for 8 h, the levels of Aβ40 were measured from the conditioned media by using specific ELISA methods. The levels of Aβ40 (n = 4) were decreased by the extract only at the higher concentrations. **, P<0.01; ***, P<0.001.

Secreted level of sAPPβ-sw were also reduced by 34.5 ± 1%, 49.1 ± 4.8% and 52.7 ± 2% at 1, 5 and 10 µg/ml of extract, respectively (Figure 4.7). In contrast, this extract significantly increased the secreted level of sAPPα in a dose-dependent manner. The sAPPα level was increased by two-fold at 10 µg/ml of the *X. undulatum* leaf extract. Secreted level of sAPPα level was increased by 23.2 ± 2.0 and 42.1 ± 2.0 at 5 and 10 µg/ml of extract, respectively (Figure 4.8). This data suggests that the leaf extract increased the non-amyloidogenic processing of APP, while it decreased amyloidogenic processing of APP, leading to the reduction of Aβ level.

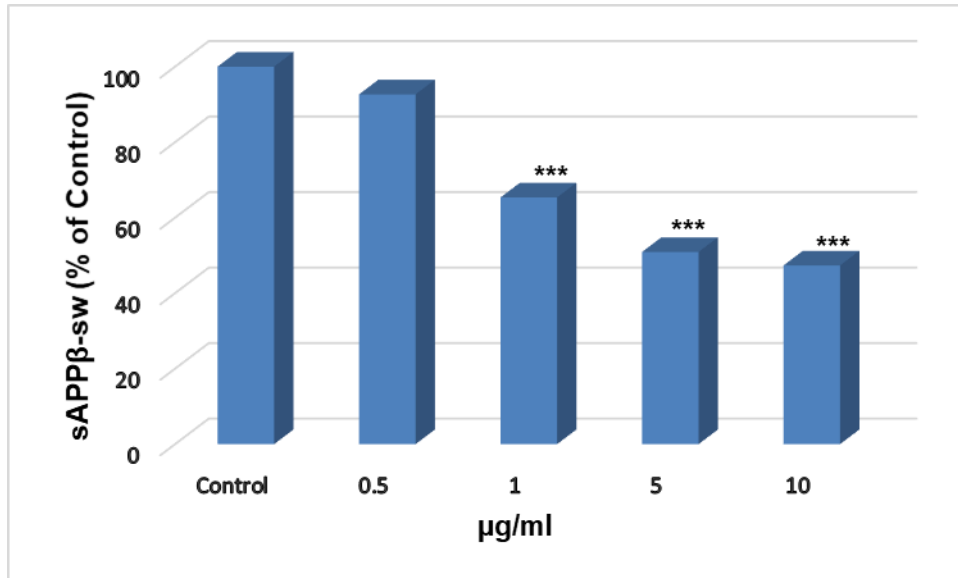


Figure 4.7: Change in the levels of sAPPβ-sw. Cells were incubated with indicated concentrations of leaf extract of *X. undulatum* for 8 h, the levels of sAPPβ-sw were measured from the conditioned media by using specific ELISA methods. The levels of sAPPβ-sw (n = 4) were decreased by the extract at higher test concentrations. **, P<0.01; ***, P<0.001.

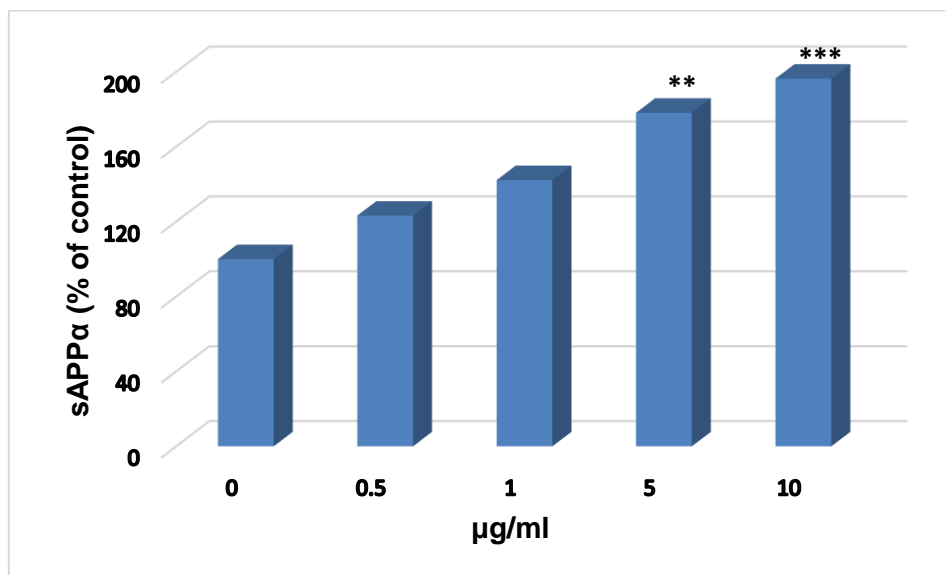


Figure 4.8: Change in the levels of sAPPα. Cells were incubated with indicated concentrations of leaf extract of *X. undulatum* for 8 h, the levels of sAPPα were measured from the conditioned media by using specific ELISA methods. The levels of sAPPα (n = 4) were increased by the extract. **, P<0.01; ***, P<0.001.

These results suggest that *X. undulatum* leaf extract potentially decreased A β 42 production at non-toxic concentrations through decreasing amyloidogenic processing of APP and increasing the nonamyloidogenic processing of APP.

Although the leaf extract *X. undulatum* has some cytotoxicity, it potentially induced the decrease in A β 42 production. The plant was chosen for further studies to determine the active ingredients responsible for its activity of A β 42 reduction. In addition, the leaf extract showed its efficacy to exhibit neuroprotective properties through inhibition of A β 42 and simultaneously increasing sAPP α secreted levels. Both these features are responsible for memory enhancement [21].

4.3.3. UPLC-MS-QTOF analysis of *X. undulatum* leaf extract

UPLC-MS-QTOF analysis was performed in ESI negative mode for the DCM: MeOH leaf extract of *X. undulatum* to identify the major peaks and the compounds which could be responsible for the reduction of A β 42. In ESI negative mode, the peaks were mainly distributed within the 3.20 to 6.40 min range of the 18 min chromatogram (Figure 4.9)

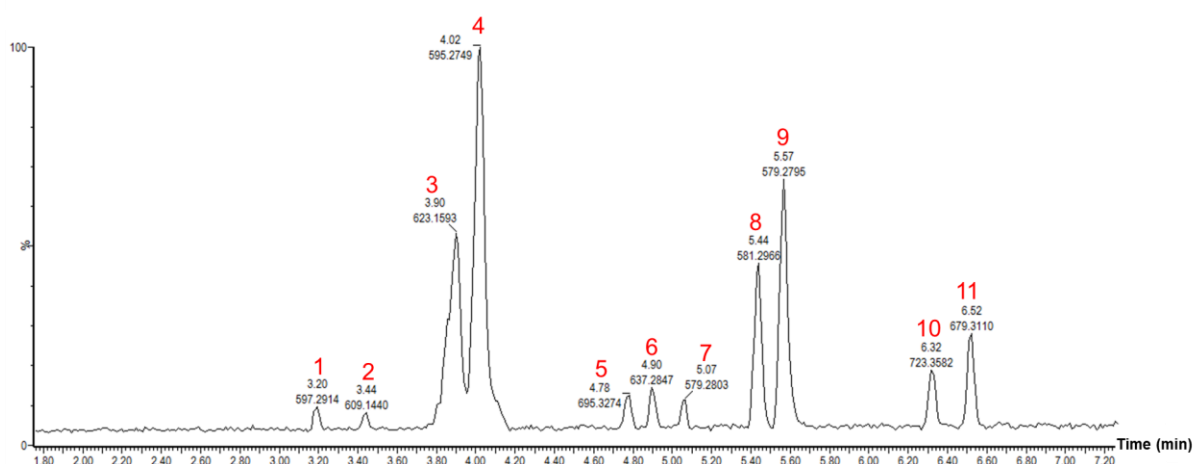


Figure 4.9: ESI negative mode BPI chromatogram of *X. undulatum* leaf extract.

In total, nine compounds were successfully detected and tentatively identified which included seven cardenolides, two flavone C-glycosides (rutin and isorhamnetin-3-O-rutinoside), based on the iFit values, mass error, mass fragmentation patterns and published data (Table 4.2.). The different electronic database, Chempider, Pubchem, Mass Bank, Scifinder, Scopus, Reaxys, Metalin Scripps and ChEBI (Chemical Entities of Biological Interest) were used for the tentative identification of compounds.

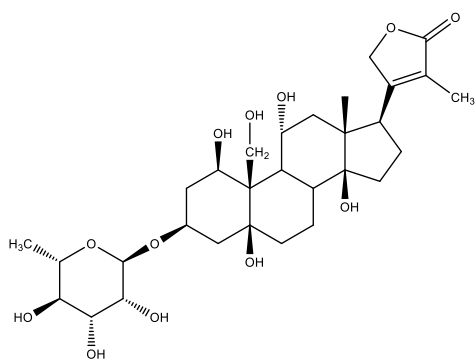
Table 4.2: Tentative identification of compounds obtained from ESI-MS negative mode of DCM: MeOH extract of *X. undulatum* (leaf)

Peak	RT (min)	Acquired [M-H] ⁻ m/z	Formula of Possible structure	Theoretical [M-H] ⁻ m/z	Tentative name	Mass Error (ppm)	MS/MS Data (Fragments)	Ref.
1	3.20	597.2914	C ₃₀ H ₄₆ O ₁₂	597.2911	22-methylouabain	0.5	587.26 50, 433.20 82, 304.91 43, 174.95 60	Reaxys
2	3.44	609.1440	C ₂₇ H ₃₀ O ₁₆	609.1455	rutin	-2.4	301.02 56, 300.03 09,	[22]
3	3.90	623.1593	C ₂₈ H ₃₂ O ₁₆	623.1612	isorhamnetin-3-O-rutinoside (ramnazin-3-O-rutinoside)	-3.0	315.04 90, 299.01 74, 271.02 22	[23]
4	4.02	549.2687	C ₂₉ H ₄₂ O ₁₀	549.2699	glucocorotoxinogenin	-2.1	387.21 44	http://www.chemspider.com
5	4.78	695.3274	C ₃₅ H ₅₂ O ₁₄	695.3278	xysmalorin	-0.5	533.27 62, 387.21 71	[24]
6	4.90	637.2847	C ₃₂ H ₄₆ O ₁₃	637.2860	unidentified	-2.0	533.27 56, 256.87 93, 174.95 65	-
7	5.07	579.2740	C ₃₀ H ₄₄ O ₁₁	579.2805	unidentified	-11.2	460.78 69 414.80 52 338.85 90	-

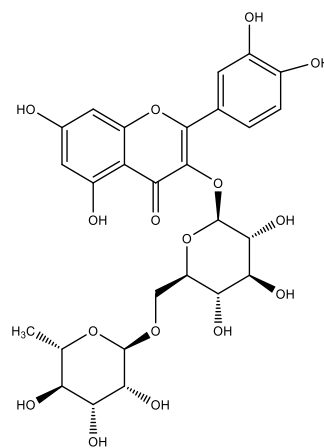
							274.87 97	
8	5.44	535.2877	C ₂₉ H ₄₄ O ₉	535.290 7	desglucouzarin	-5.6	373.23 82	[25]
9	5.57	533.2766	C ₂₉ H ₄₂ O ₉	533.275 0	gofruside	3.0	533.27 97, 387.21 74	Reaxys
10	6.32	677.3536	C ₃₆ H ₅₄ O ₁₂	677.353 7	corotoxigenin-3-O-β-digitalopyranosyl-(1→4)-O-β-digitoxopyranoside	-0.1	517.27 67	[26]
11	6.52	679.3110	C ₃₈ H ₄₈ O ₁₁	679.311 8	12β-(acetyloxy)-(3β,5β)-3-[4-carboxy-3-[4-(acetyloxy)-phenyl]-1-oxobutoxy]-14-hydroxycard-20(22)-enolide	-1.1	163.03 98, 145.02 96	https://pubchem.ncbi.nlm.nih.gov

Peak 1 exhibited a precursor ion [M-H]⁻ at *m/z* 597.2194 which corresponded with the molecular formula C₃₀H₄₆O₁₂. MS/MS spectra of deprotonated precursor ion showed different fragment ions at *m/z* 433.2082 [M-164.0112]⁻ with the loss of a glucoside moiety. The use of the electronic database, Reaxys, and the proposed molecular formula led to the tentative identification of a cardenolide, 22-methylouabain (**46**). The mass spectrum of peak 2 showed a molecular ion at *m/z* 609.1455 [M-H]⁻, producing a quercetin daughter ion at *m/z* 301.0256 [M-308.1199]⁻ with the loss of rhamnosyl-glucoside moiety. The elemental composition proposed the molecular formula C₂₇H₃₀O₁₆. From all the observed data, the compound was tentatively identified as rutin (**47**). The MS analysis of peak 3 showed a molecular ion at *m/z* 623.1593 with molecular formula C₂₈H₃₂O₁₆, releasing one MS² aglycone fragment ion of isorhamnetin at *m/z* 315.0490 [M-308.1103]⁻. This fragment ion showed the loss of rhamnosyl-glucoside moiety. The previous studies and the mass spectrum analysis led to the identification of isorhamnetin-3-O-rutinoside (ramnazin-3-O-rutinoside) (**48**) [27], [23]. Loss of rhamnosyl-glucoside fragment in rutin and isorhamnetin-3-O-rutinoside is a characteristic of flavone C-glycosides [27]. The MS analysis of peak 4 presented a pseudo molecular ion at *m/z* 595.2749 [M-HCOO]⁻ and was assigned the elemental composition C₃₀H₄₄O₁₂ (calculated

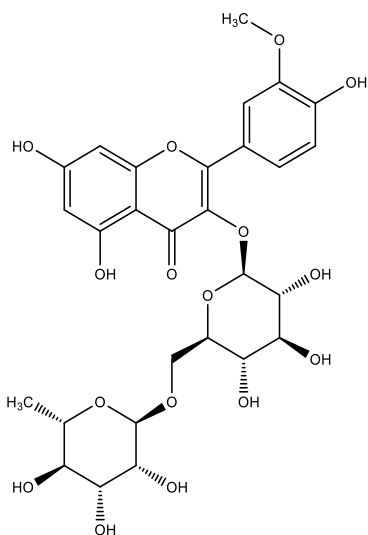
mass 550.2778). The MS/MS spectra gave the fragment ion at m/z 387.2144 [M-162.0543]⁻ with the loss of a glucoside moiety. This typical fragmentation pattern led to the tentative identification of the cardenolide, glucocorotoxigenin (**49**). The mass spectrum of peak 5 presented a molecular ion at m/z 695.3274 [M-H]⁻, corresponding to the formula C₃₅H₅₂O₁₄. Two MS² fragment ions, one at m/z 533.2756 [M-162.0518]⁻ and the other at m/z 387.2171 [M-162.0518-146.0585]⁻ were detected and corresponded to the loss of pentosylglucoside and rhamnosylglucoside, respectively. In accordance with literature and the mass fragmentation pattern, the compound was tentatively identified as xysmalorin (**43**) [24], for which the structure of the compound is shown in Figure 4.1. Peak 8 presented a pseudomolecular ion at m/z 581.2966 [M-HCOO]⁻ with a molecular formula C₃₀H₄₆O₁₁ (calculated mass 536.2985). The MS/MS spectrum produced the fragment ion at m/z 373.2382 [M-162.0495]⁻ with the loss of glucoside moiety. The typical fragmentation pattern of cardenolides and the literature study led to the tentative identification of compound as desglucouzarin (**50**) [25]. Peak 9 presented a pseudomolecular ion at m/z 579.2795 [M-HCOO]⁻ with a molecular formula C₃₀H₄₄O₁₁ (calculated mass 533.2750). The characteristic fragment signals at m/z 387.2174 [M-146.0582]⁻ indicated the loss of a deoxyhexose moiety. Based on the literature and fragmentation pattern, the compound was identified as the cardenolide, gofruside (**51**). The MS analysis of peak 10 showed a pseudo molecular ion peak at m/z 723.3582 with a predicted formula C₃₆H₅₄O₁₂ (calculated mass 678.3615). The fragment ions were detected at m/z 517.2767 [M-160.0926]⁻ with the loss of digitalose unit. Based on the previous study and the fragmentation pattern, this peak was tentatively identified as corotoxigenin-3-O-β-digitalopyranosyl-(1→4)-O-β-digitoxopyranoside (**52**) [26]. Finally, peak 11 presented molecular ion at m/z 679.3110 [M-H]⁻. Through the elemental composition, the molecular formula was predicted as C₃₈H₄₈O₁₁ with the release of fragment ion at m/z 163.0398 [M-516.2712]⁻. All this data including that in Pubchem led to the tentative identification of this peak as 12β-(acetyloxy)-(3β,5β)-3-[4-carboxy-3-[4-(acetyloxy)phenyl]-1-oxobutoxy]-14-hydroxycard-20(22)-enolide (**53**). The structures of the identified compounds are shown in Figure 4.10.



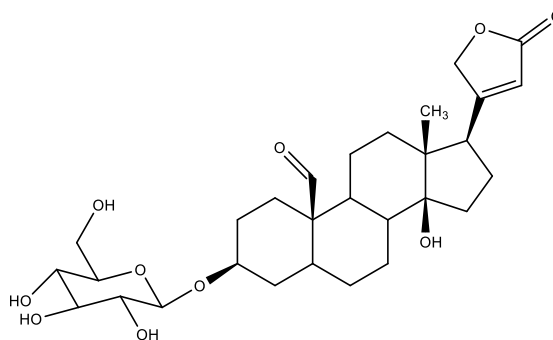
(46)



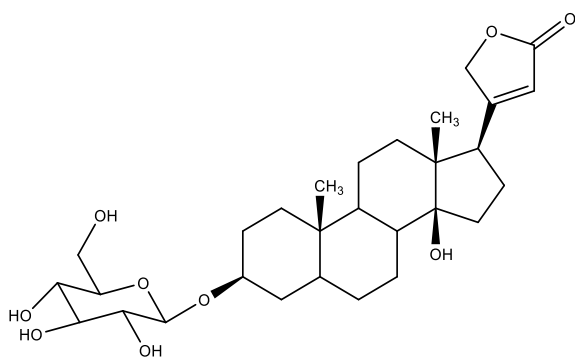
(47)



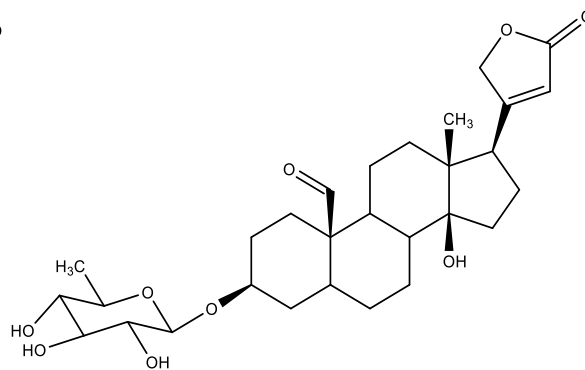
(48)



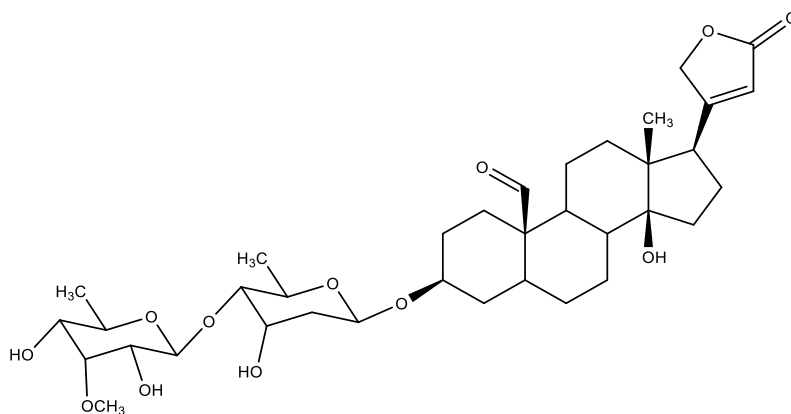
(49)



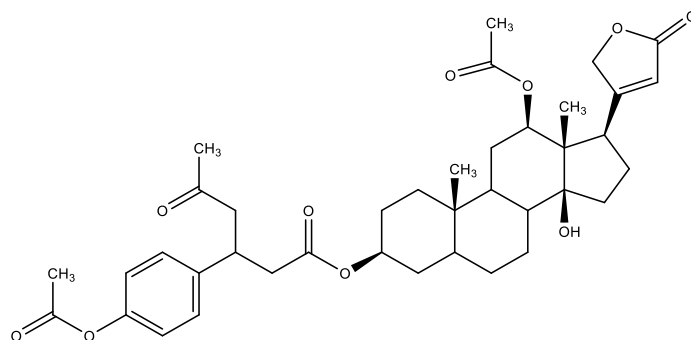
(50)



(51)



(52)

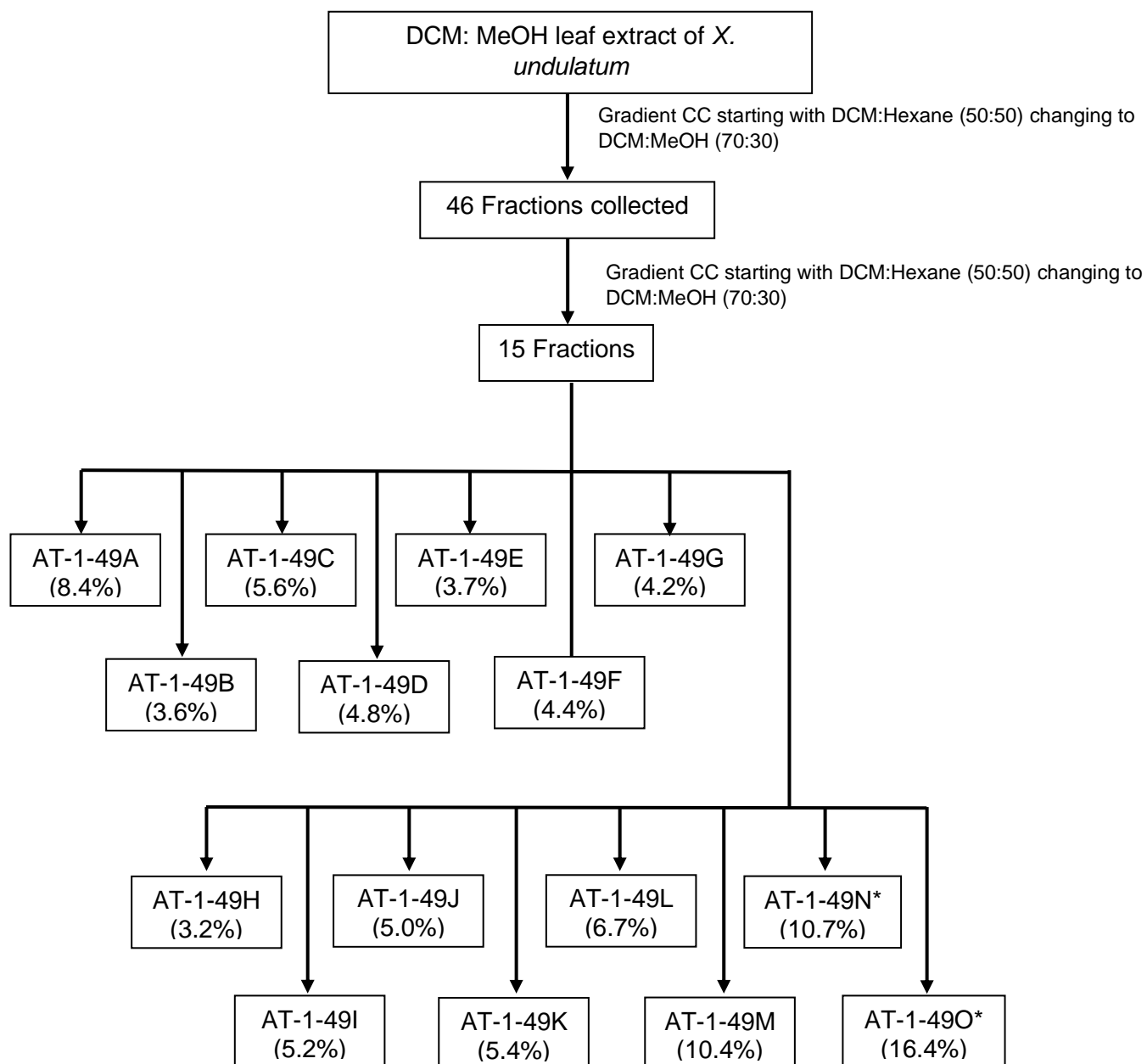


(53)

Figure 4.10: Structures of 22-methylouabain (**46**), rutin (**47**), isorhamnetin-3-O-rutinoside (Ramnazin-3-O-rutinoside) (**48**), glucocorotoxigenin (**49**), desglucouzarin (**50**), gofruside (**51**), corotoxigenin-3-O- β -digitalopyranosyl-(1 \rightarrow 4)-O- β -digitoxopyranoside (**52**), 12 β -(Acetyloxy)-(3 β ,5 β)-3-[4-carboxy-3-[4-(acetyloxy)-phenyl]-1-oxobutoxy]-14-hydroxycard-20(22)-enolide (**53**) identified in DCM:MeOH leaf extract of *X. undulatum*

4.3.4. Fractionation using silica gel column chromatography

Silica gel column chromatography of the leaf extract (17g) using the gradient solvent system resulted in the collection of 46 fractions. Fractions were analysed by TLC and those fractions which showed similar profiles based on compounds with similar R_f values were combined to give 15 fractions as shown in Figure 4.11 which were evaluated for their A β 42 production.



* active fractions

Figure 4.11:Flow diagram for the fractionation of the leaf extract of *X.undulatum* using silica gel column chromatography. Value in the brackets indicated the percentage yield (w/w) of the fractions obtained from silica gel column.

4.3.5. Bioassaying of fractions for A β 42 production

The 15 fractions (AT-1-49A to AT-1-49O) were evaluated for their activity on A β 42 production by incubating the HeLa cells with 10 μ g/ml of each fraction for 8 h and the levels of A β 42

measured from the conditioned media by specific ELISA. Two fractions (AT-1-49N and AT-1-49O) potently decreased the A β 42 secretion by $73.6 \pm 0.8\%$ and $65.3 \pm 0.01\%$, respectively, while all other fractions showed no or minimal reduction of A β 42 as shown in Figure 4.12. The results indicated that the silica gel column chromatography successfully separated the active from the non-active fractions. The two active fractions were further purified to isolate the active compounds.

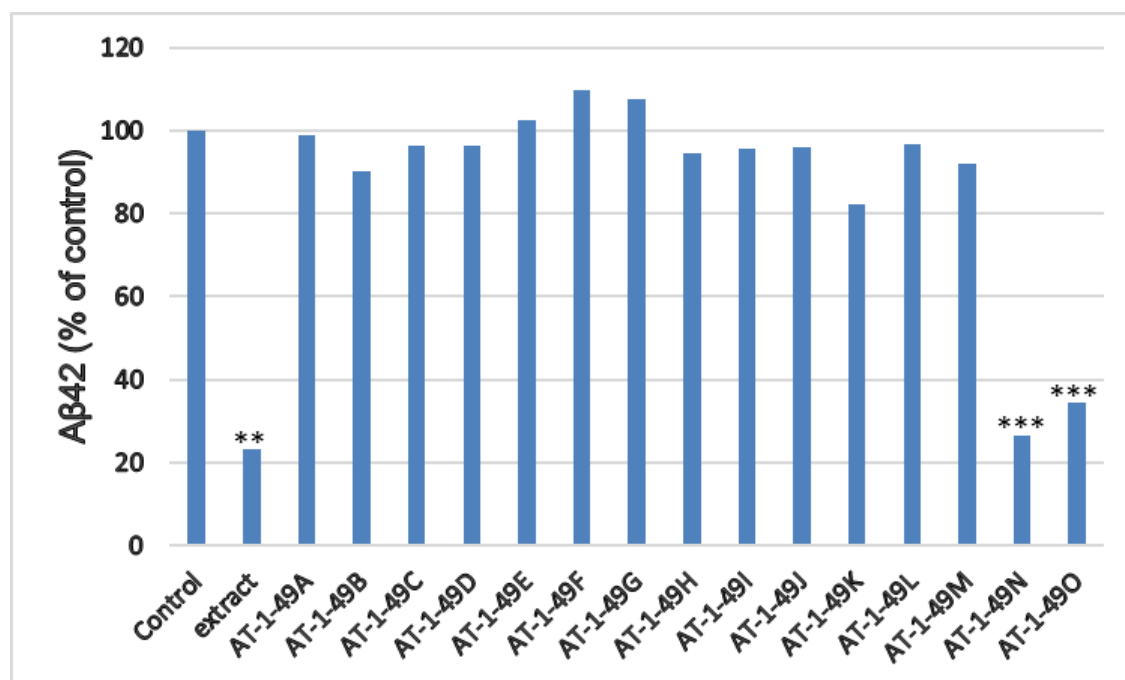


Figure 4.12: Effect of fractions on the production of A β 42 in APPsw-transfected HeLa cells. 15 Fractions were obtained from leaf extract of *X. undulatum* by silica gel column chromatography. Cells were incubated with 10 μ g/ml extract and 10 μ g/ml fractions for 8 h, and the level of A β 42 was measured from the conditioned media by specific ELISA methods. The two fractions, AT-1-49N and AT-1-49O, potently decreased the secreted level of A β 42 (n = 2). *, P<0.05; **, P<0.01; ***, P<0.001.

4.3.6. UPLC- MS analysis the active fractions

UPLC-MS analysis of the active fractions tentatively identified a mixture of cardenolides. Based on literature reports, it was found that the cardenolides in ESI MS negative mode formed abundant adduct ions [M+HCOO]⁻ while pseudo molecular ions [M-H]⁻ were obtained in ESI-MS/MS experiments [28]. The similar pattern was observed in the active fractions AT-

1-49N and AT-1-49O which supported the presence of cardenolides. In ESI-MS negative mode for fraction AT-1-49N (Figure 4.13), peak 1 with m/z 637.2856, peak 2 with m/z 433.2226, peak 3 with m/z 579.2813, peak 4 with m/z 723.3585 and peak 6 with m/z 709.3793 were obtained as formate adduct ions $[M+HCOO]^-$ while their pseudo molecular ions $[M-H]^-$ observed at m/z 591.2797, m/z 387.2164, m/z 533.2763, m/z 677.3526 and m/z 663.3740, respectively in the ESI MS/MS spectra (**supplementary data 1 to 5**). All the peaks except peak 2 and peak 6 were analysed earlier as described in section 4.1.2.

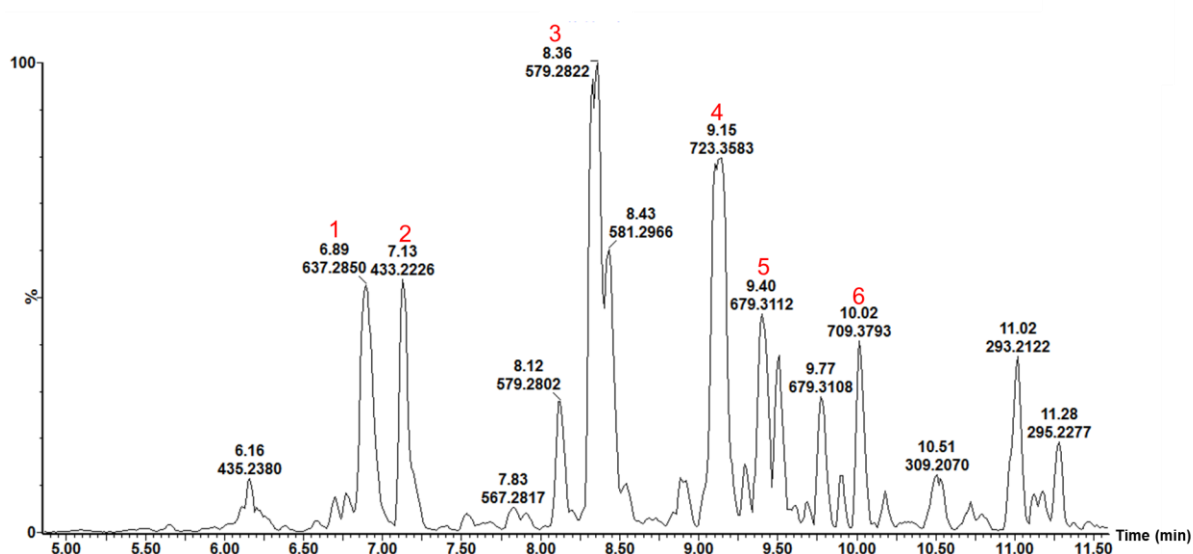


Figure 4.13: ESI negative BPI chromatogram of active fraction AT-1-49N

Peak 2 and peak 6 had not been detected in the crude parent DCM:MeOH leaf extract. This indicated either their poor ionization or low concentrations in the extract. Peak 1 (637.2850) showed a formic acid adduct ion in fraction AT-1-49N while this adduct did not appear in MS/MS fragmentation pattern of the DCM:MeOH extract due to its poor ionisation. The molecular formula for these compounds were generated through MassLynx and the compounds were identified using online database searches (Table 4.3). The structures of the compounds are shown in Figure 4.14.

Table 4.3: Tentative identification of compounds obtained from ESI-MS negative mode of fraction AT-1-49N obtained from DCM: MeOH extract of *X. undulatum* (leaf)

Peak	RT (min)	Acquired [M-H] ⁻ m/z	Formula of Possible structure	Theoretical [M-H] ⁻ m/z	Tentative name	Mass Error (ppm)	Ref.
1	6.89	591.2797	C ₃₁ H ₄₄ O ₁₁	591.2805	adonitoxigenin-(2-acetyl-rhamnosid)	-1.3	Reaxys
2	7.13	387.2164	C ₂₃ H ₃₂ O ₅	387.2171	crotoxinigenin	-1.8	Reaxys
3	10.02	699.3505	C ₄₉ H ₄₈ O ₄	699.3474	unidentified	-	-

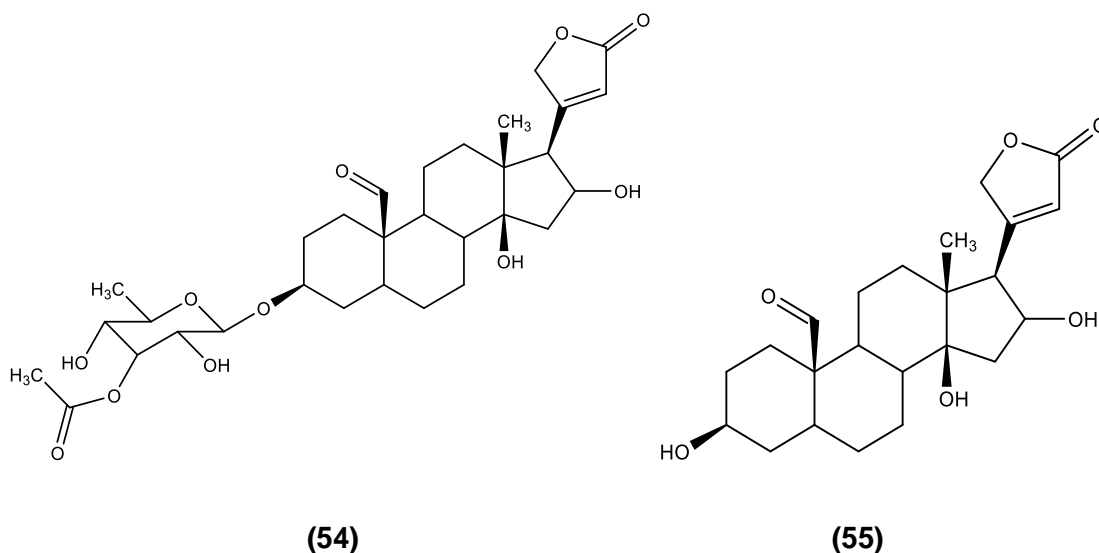


Figure 4.14: Structures of adonitoxigenin-(2-acetyl-rhamnosid) (54), crotoxinigenin (55)

Similarly formate adduct ions [M+HCOO]⁻ were observed in fraction AT-1-49O (Figure 4.15) with *m/z* 581.2970 and *m/z* 595.2756 while their pseudo molecular ions obtained in MS/MS spectra were *m/z* 535.2970 and *m/z* 549.2701, respectively (**supplementary data 6 and 7**). The analysis of these compounds were examined earlier as described in section 4.1.2.

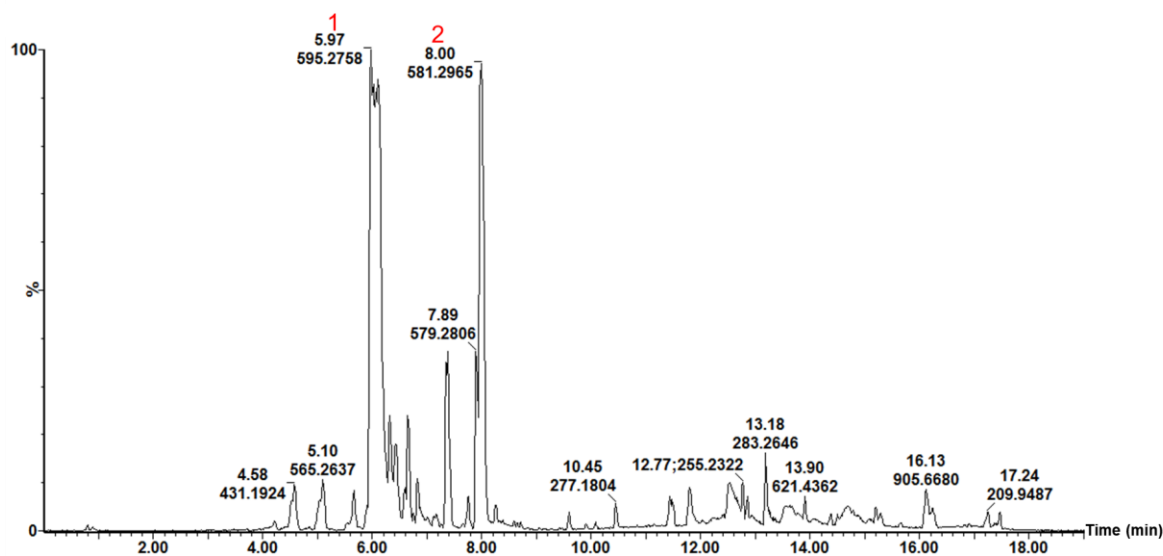


Figure 4.15: ESI negative BPI chromatogram of fraction AT-1-49O.

The common class of compounds, cardenolide glycosides was identified from both the active fractions AT-1-49N and AT-1-49O. This indicated that the cardenolide glycosides could be the active ingredients in the present study. These fractions were further purified using preparative HPLC.

4.3.7. Isolation and purification of compounds from active fractions using preparative HPLC

AT-1-49N was refined by using mass-directed purification using preparative HPLC. The fractions were collected in test tubes by using mass triggered fractionation. The fractions with the similar masses were combined and evaporated to dryness using rotary vacuum evaporator. The purity of compounds was further confirmed with UPLC-MS and NMR analysis. All m/z values in the ESI negative mode (Figure 4.16) corresponding to major peaks labelled as XU01 (637.2850), XU02 (433.2226), XU03 (579.2822), XU04 (723.3583), XU05 (679.3112), XU06 (709.3793) were targeted and collected. Of these major peaks, three XU01 (4 mg), XU03 (3.5 mg) and XU04 (4.9 mg) were collected in pure form with sufficient quantities to allow further

analysis. ESI negative-mode BPI chromatograms of pure compounds XU01, XU03 and XU04 is shown in figure 4.16.

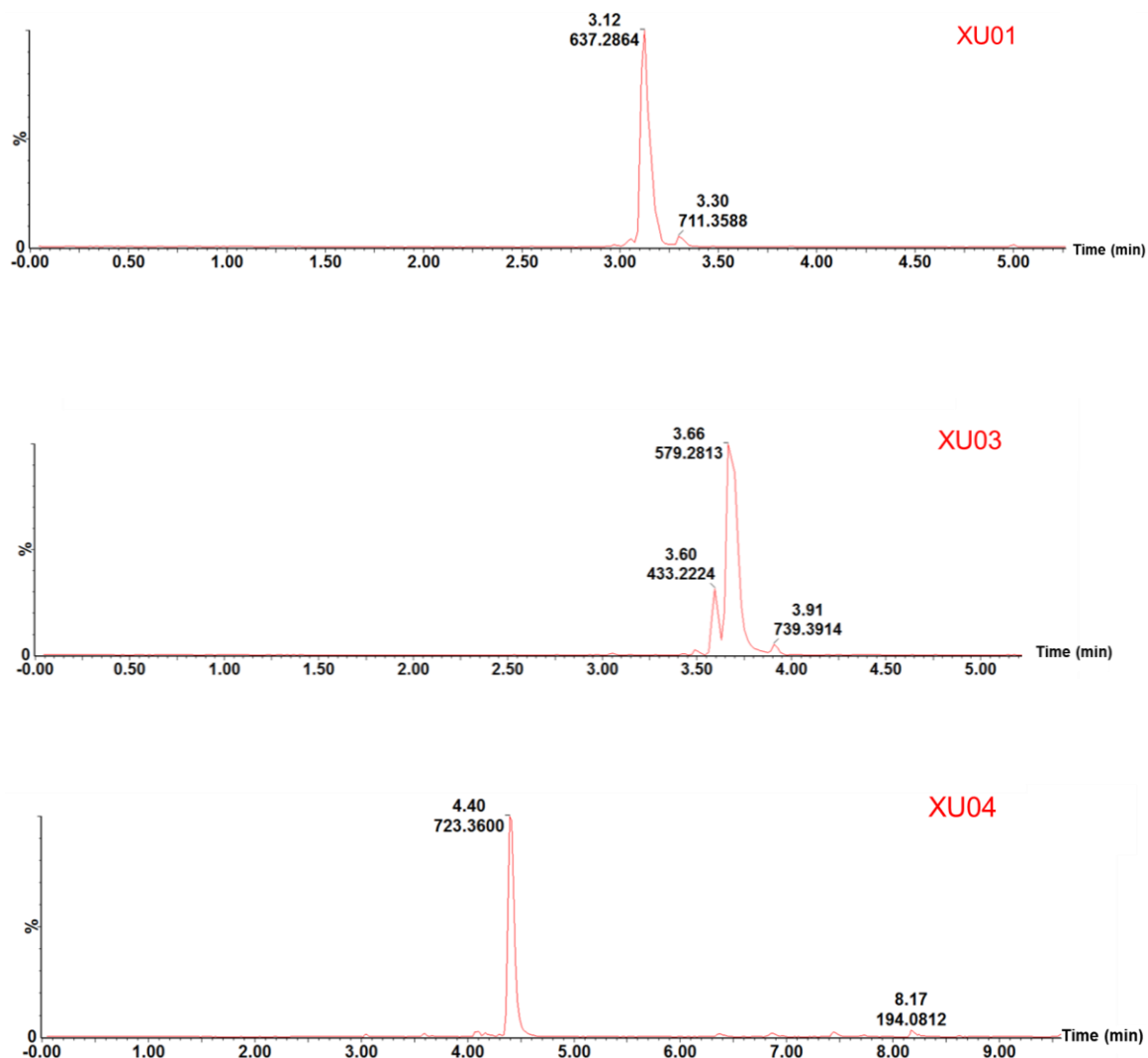


Figure 4.16: ESI negative-mode BPI chromatograms of pure compounds XU01, XU03 and XU04 from active fraction AT-1-49N.

Similarly, AT-1-49O was also resolved using mass-directed purification preparative HPLC. The m/z values in the ESI negative mode (Figure 4.13) corresponding to major peaks 595.2758 and 581.2965 were targeted and collected in pure form. These were labelled as XU07 (5 mg) and XU08 (4 mg). The ESI negative-mode BPI chromatograms of pure compounds XU07 and XU08 is shown in Figure 4.17.

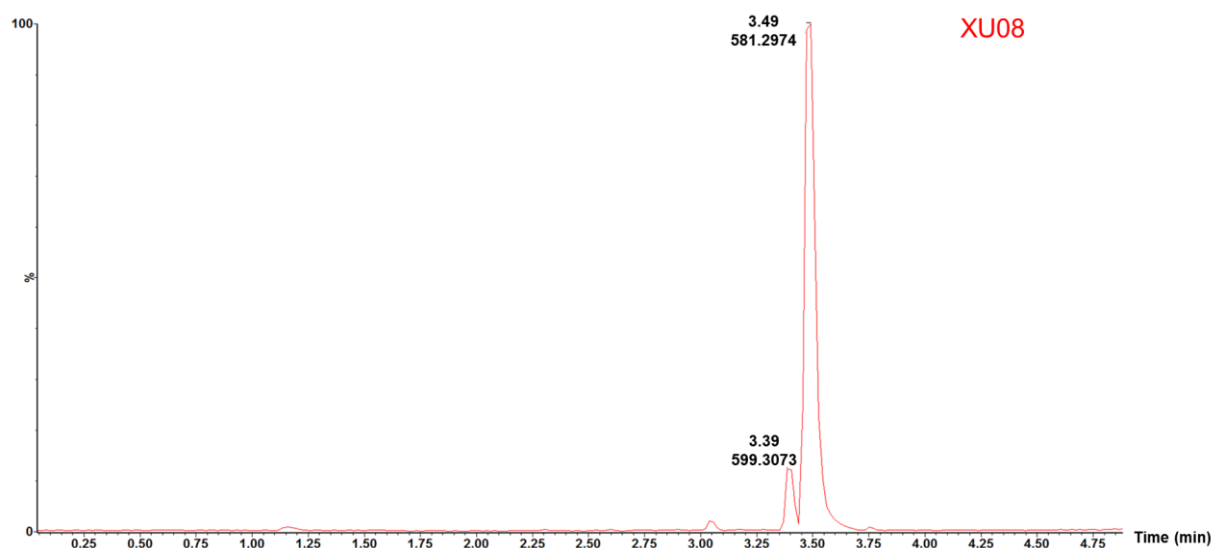
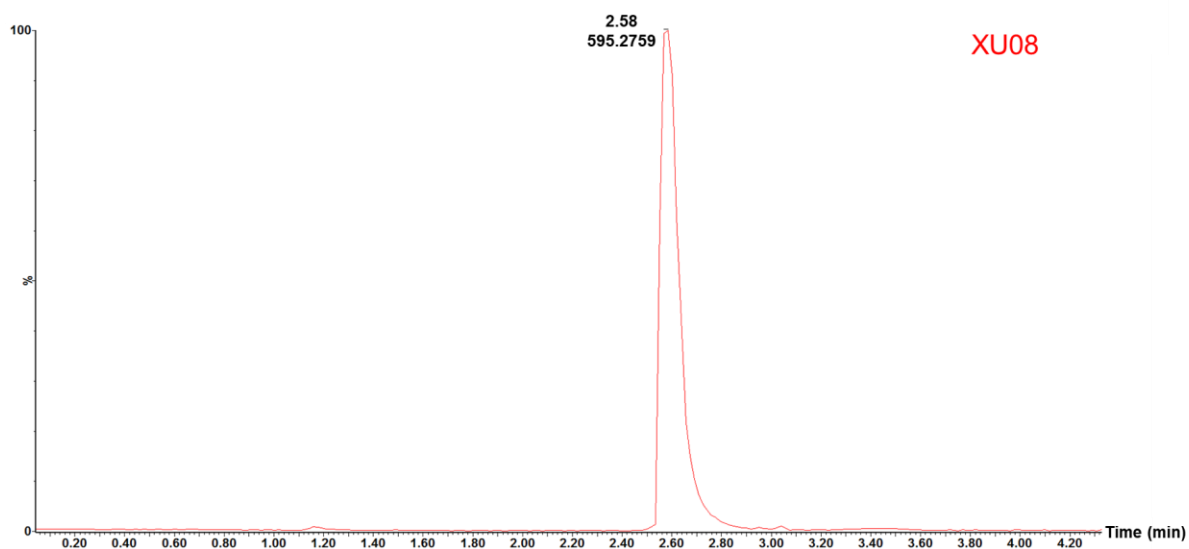


Figure 4.17: ESI negative-mode BPI chromatograms of pure compounds XU07 and XU08 from active fraction AT-1-49O.

4.3.8. Structural elucidation using NMR

4.3.8.1. Acetylated glycosydated crotoxigenin (56)

The pure compound (XU01) was identified as the novel acetylated glycosydated crotoxigenin (**56**) based on the NMR data and mass spectral analysis. It was isolated as a white amorphous powder with a mass of 4 mg. It had a molecular formula of $C_{31}H_{44}O_{11}$ (calculated mass

592.2883), as deduced from its pseudomolecular precursor ion at m/z 637.2851[M-HCOO]⁻ based on the QTOF mass spectrum. These data confirmed the presence of cardenolide glycoside. The formula was further confirmed by the number of protons in ¹H NMR and of carbon atoms in the ¹³C NMR spectrum (**Supplementary data 6, 7**). The structure of the compound (**56**) is shown in Figure 4.18.

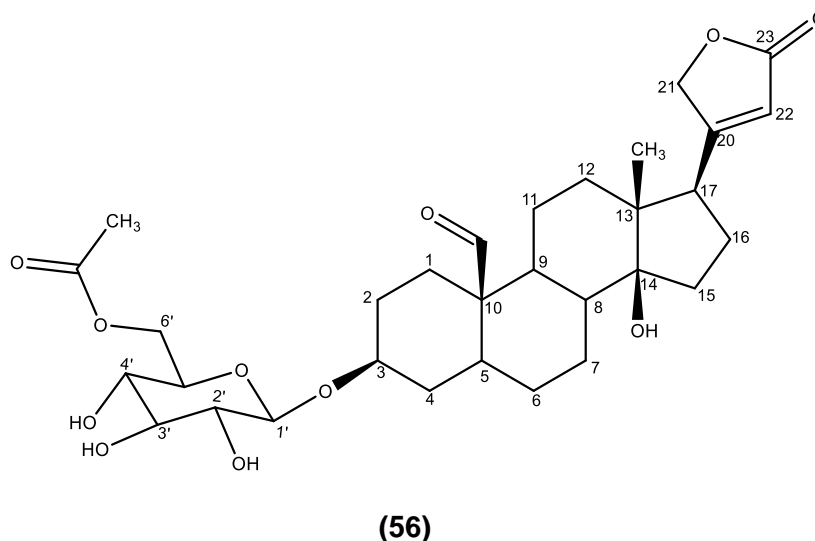


Figure 4.18: Structure of acetylated glycosydated crotoxogenin (**56**)

The ¹H NMR spectrum (MeOD-*d*₄, 400 MHz) of the compound showed the H-21 protons at δ 5.03 (1H, dd) and δ 4.96 (1H, dd) and an olefinic proton at δ 5.92 (1H, s, H-22) as part of the lactone ring which indicated the characteristic feature of the cardenolide system. The other prominent signal indicated the presence of one upfield methyl proton at δ 0.98 (3H, s, H-18). The spectrum also indicated one downfield aldehydic proton at δ 9.57 (1H, s, H-19) and overall pattern of the ¹H NMR indicated that the compound was a cardenolide with one sugar unit as an anomeric proton signal was observed at δ 4.40 (1H, d, H-1'). The chemical shift at δ 2.08 (3H, s, 5'-OCOCH₃) confirmed the presence of an acetate group, in all likelihood attached to the sugar moiety.

The ¹³C NMR spectrum had a total of 31 signals and the assignments were done together with the DEPT 135 spectra (**supplementary data 8**). The methylene signal at δ 73.9 was assigned

to C-21, δ 116.4 assigned to C-22 olefinic carbon, δ 176.9 assigned to the C-23 carbonyl group and δ 177.8 assigned to the quaternary C-20, all of which confirmed the presence of the butenolide ring. Two prominent signals at δ 14.8 assigned to C-18 and at δ 206.5 assigned to C-19 indicated the position of methyl carbon and carbonyl carbon, respectively. The β hydroxyl group at C-14 was confirmed by the downfield chemical shift at δ 84.5. One anomeric carbon was revealed at δ 101.3 which was assigned to C-1'. The carbon at δ 171.2 (5'-OCOCH₃) revealed the presence of acetate group and carbon at δ 19.3 represent the position of methyl group.

The HSQC and COSY correlations (**supplementary data 9, 10**) led to the assignment of the proton and carbon signals. The chemical shifts for ¹H and ¹³C were shown in Table 4.4.

Figure 4.19: ¹H NMR and ¹³C NMR data for glycosydated crotoxigenin in MeOD-*d*4

Position	¹³ C (ppm)	¹ H (ppm), J (Hz)	COSY	HMBC
1a	28.1	1.62, m,		C-9
1b		1.51, m		
2a	33.0	1.96, m,		C-3, C-10
2b		1.78, m		
3	77.8	3.74, m		
4a	25.9	1.87, m,		
4b		1.37, m		
5	41.6	1.89, m		C-9
6a	25.1	1.92, m,		C-5
6b		1.35, m		
7a	21.2	1.80, m,		C-6, C-8
7b		1.34, m		
8	35.0	1.98, m		C-5
9	35.1	1.83, m		
10	50.5	-		
11a	20.3	1.79, m,		C-9, C-10
11b		1.51 m		
12	39.3	1.88, m		C-9
12b		1.57, m		
13	49.6	-		
14	84.5	-		
15a	31.3	2.21 m,		
15b		1.72 m		
16a	26.5	2.17, m		C-14
16b		1.61, m		
17	50.4	2.80 (q, 14.6)		C-13, C-14, C-21, C-22
18	14.8	0.97, s		C-12, C-13, C-14
19	206.5	9.57, s		C-9
20	175.8	-		

21a	73.9	5.03 (dd, 18.5, 1.57),		C-22
21b		4.96 (dd, 18.1, 1.73)		
22	116.4	5.92 s		C-17, C-21, C-23
23	176.9	-		
1'	101.3	4.40 (d, 7.57)	H-2'	C-3
2'	73.6	3.17, m	H-1', H-3'	C-1'
3'	70.2	3.36, m	H-2'	
4'	73.7	3.30, m	H-5'	
5'	76.4	3.47, m	H-4'	
6'a	63.4	4.20 (dd, 11.6, 5.14)	H-5'	C-2', C-5' -OCOCH ₃
6'b		4.30 (dd, 11.6, 1.84)		
5'- OCO	171.2	-		
5'- CH ₃	19.3	2.08 s		C-5' -OCO

The strong HMBC correlation (**supplementary data 11**) between the proton at δ 2.80 (H-17) with C-21 (δ 73.9) and C-22 (δ 116.4) confirmed that the α,β -unsaturated γ -lactone was connected at C-17 (δ 50.4). The proton at δ 9.57 (H-19) is correlated to C-9 (δ 35.1) confirmed the position of aldehydic proton. HMBC correlations between C-5'- CH₃ with C-5'- OCO confirmed the presence of acetate group. HMBC correlations also established the linkages of the glycosidic bonds and the point of attachment of the saccharide chain to the aglycone based on the correlation between δ 4.40 (H-1') to δ 77.8 (C-3). The proton at δ 3.10 (H-2') correlating with δ 101.3 (C-1') and δ 73.7 (C-4'); proton δ 3.30 (H-4') correlating with δ 76.48 (C-5') and δ 63.40 (C-6') confirmed a single sugar moiety.

The large coupling constant of the anomeric proton at (δ 4.40, 7.57 Hz) indicated the β -orientation of the glycoside moiety. There was a COSY correlation between the anomeric proton at δ 4.4 (H-1') and the proton at δ 3.17 (H-2') while another correlation was observed between the proton at δ 3.17 (H-2') and the proton at δ 3.36 (H-3'). The proton at δ 3.30 (H-4') showed a COSY correlation with proton at δ 3.47 (H-5'). The proton at δ 3.47 (H-5') coupled with the two methylene protons at δ 4.20 (C6'a) and δ 4.30 (C6'b) to give a multiplet. These methylene protons also coupled with each other, hence the two sets of doublets of doublets at δ 4.20 (*dd*, 11.6 Hz, 5.14 Hz, H-6'a) and δ 4.30 (*dd*, 11.6 Hz, 1.84 Hz, H-6').

The HMBC and COSY relations are shown in the Figure 4.19.

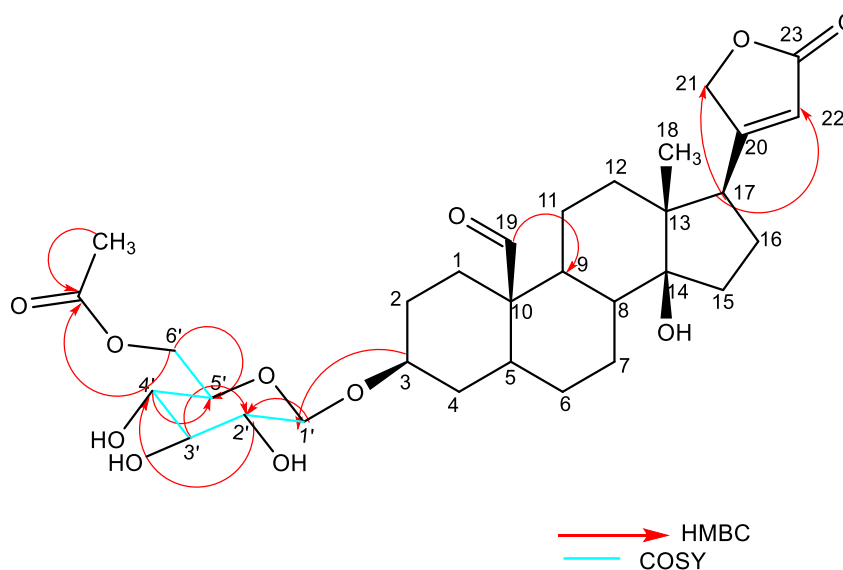


Figure 4.20: Key HMBC and COSY correlations of acetylated glycosydated crotoxegenin (**56**)

4.3.8.2. Xysmalogenin-3, β -D-glucopyranoside (**57**)

The pure compound (XU03) obtained from fraction AT-1-49N was identified as a known compound xysmalogenin-3, β -D-glucopyranoside (**57**) based on the NMR data, mass spectral analysis and previous published studies [2]. However, the exact and complete NMR data for this compound was not previously published. It was isolated as a white solid weighing 3.5 mg and had a molecular formula of $C_{29}H_{42}O_9$ (calculated mass 534.2828), as deduced from its pseudomolecular precursor ion at m/z 579.2822 $[M-HCOO]^-$ based on the QTOF mass spectrum. These data confirmed the presence of cardenolide glycoside. The formula was further confirmed by the number of protons in 1H NMR and of carbon atoms in the ^{13}C NMR spectrum (**supplementary data 12, 13**). The structure of the compound (**57**) is shown in Figure 4.20.

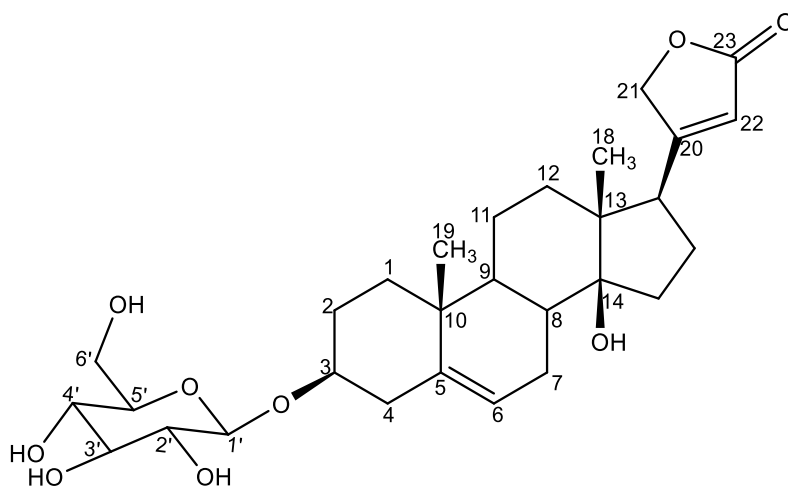


Figure 4.21: Structure of xysmalogenin-3, β -D-glucopyranoside (**57**)

The ^1H NMR spectrum (MeOD-*d*4, 400 MHz) of the compound showed the H-21 protons at δ 5.02 (1H, dd) and δ 4.92 (1H, dd) and an olefinic proton at δ 5.91 (1H, s, H-22) as a part of the lactone ring which indicated the characteristic feature of the cardenolide system. The spectrum showed the presence of one more double bonds at δ 5.44 (1H, s, H-6). The other prominent signals indicated one high field methyl proton at δ 0.92 (1H, s, H-18), and another high field proton at δ 1.03 (1H, s, H-19). The spectrum indicated the cardenolide with one sugar unit, with a signal for one anomeric proton at δ 4.38 (1H, dd, H-1'), where the large coupling constant (7.68 Hz) confirmed the β - coupling of the sugar unit to the aglycone.

The ^{13}C NMR spectrum had a total of 31 signals and the assignments were done together with the DEPT 135 spectra (**supplementary data 14**). The methylene signal at δ 75.2 was assigned to C-21, δ 117.8 assigned to C-22 olefinic carbon, δ 178.2 assigned to the C-23 carbonyl group and δ 177.1 assigned to the quaternary C-20, all of which confirmed the presence of butenolide ring. Two prominent signals at δ 16.1 assigned to C-18 and at δ 19.8 assigned to C-19 indicated the position of two methyl carbons. The β hydroxyl group at C-14 was confirmed by downfield chemical shift at δ 86.3. One anomeric carbon revealed at δ 102.3 was assigned to C-1'.

The proton and carbon signals are assigned based on the HSQC (**supplementary data 15**) and COSY correlations (**supplementary data 16**). The chemical shifts for ^1H and ^{13}C were shown in Table 4.5.

Figure 4.22: ^1H NMR and ^{13}C NMR data for xysmalogenin-3, β -D-glucopyranoside in MeOD-*d*₄

Position	^{13}C (ppm)	^1H (ppm), <i>J</i> (Hz)	COSY	HMBC
1	38.9	2.27 m		C-3, C-5, C-6
2a	30.5	1.92 m,		
2b		1.62 m		
3	79.6	3.60 m		
4	40.1	1.52 m		
5	140.8	-		
6	122.7	5.44 m	H-7	C-8
7a	27.4	2.26 m,	H-6	C-5, C-6
7b		2.22 m		
8	38.3	1.72 m		
9	47.7	1.24 m		
10	38.3			
11	22.0	1.53 m		
12	39.4	2.44 m		
13	50.7	-		
14	86.3	-		
15	33.8	1.74m		
16	28.1	1.73 m		C-14, C-15
17	52.1	2.86 (q, 14.62)		C-12, C-21, C-22, C-23
18	16.1	0.92 s		C-12, C-13, C-14
19	19.8	1.03 s		C-5, C-9, C-10
20	177.1	-		
21a	75.2	5.02, (dd, 18.44, 1.65)		C-22, C-23
21b		4.92, (dd, 18.44, 1.65)		
22	117.8	5.91 s		C-13, C-20, C-21
23	178.2	-		
1'	102.3	4.38 (dd, 7.68)	H-2'	C-3
2'	75.1	3.14 m		C-1', C-3'
3'	78.0	3.25 m		
4'	71.5	3.25 m		
5'	77.7	3.35 m		
6'a	63.6	3.84 m,	H-5'	C-4', C-5,
6'b		3.65 m		

The α,β -unsaturated γ -lactone was determined to be connected at C-17 through the strong HMBC (**supplementary data 17**) correlations between proton at δ 2.86 (H-17) to δ 177.1 (C-21), δ 117.8 (C-22) and δ 178.8 (C-23). The glycosidic linkage and the point of attachment of to the genin moiety was established by HMBC correlations from δ 4.38 (H-1') to δ 79.6 (C-3). HMBC correlations between δ 3.14 (H-2') with δ 102.3 (C-1') and δ 78.0 (C-3'); (δ 3.84) H-6'a with δ 77.7 (C-5') and δ 71.5 (C-4') confirmed the sugar unit.

A large coupling constant H1'/H2' (δ 4.38, 7.68 Hz) indicated a diaxial relationship for anomeric proton H1' and another proton H2'. This large coupling constant also indicated the β -pyranose form for glycoside by verifying the β -orientation of glycoside. COSY correlations were also observed between δ 5.44 (H-6) and δ 2.26 (H-7). The HMBC and COSY relations are shown in Figure 4.21.

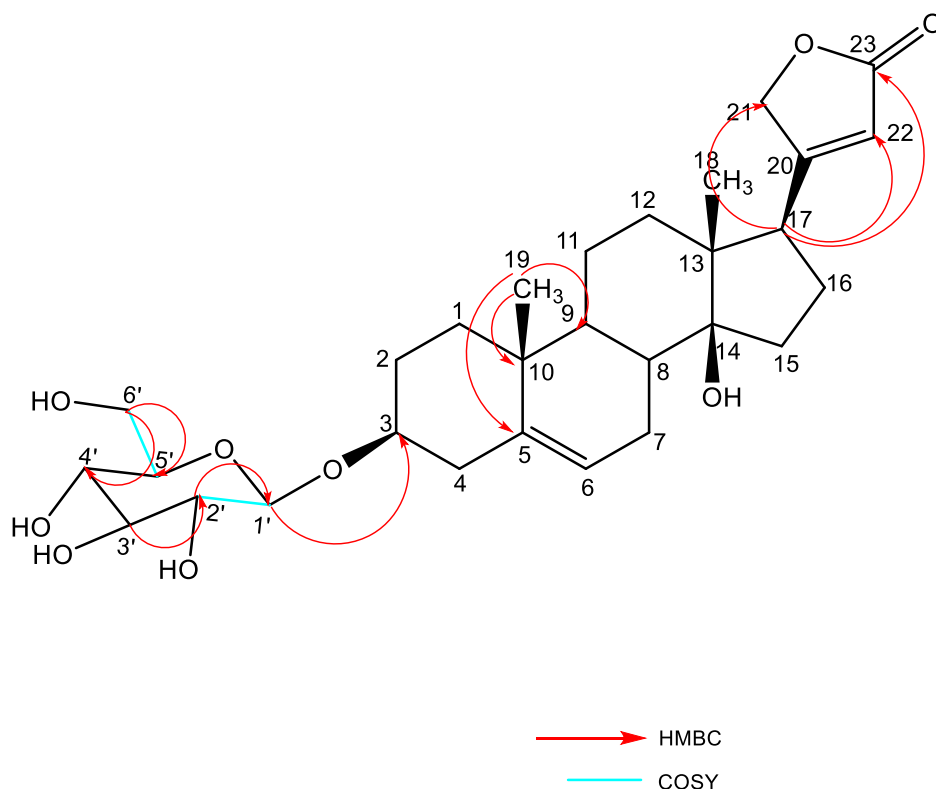


Figure 4.23: Key HMBC and COSY correlations of xysmalogenin-3, β -D-glucopyranoside (**57**)

4.3.8.3. Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**)

The pure compound (XU04) obtained from fraction AT-1-49N was identified as a known compound, crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**), based on the NMR data, mass spectral analysis and previous published studies [29], [26]. It was isolated as a white amorphous powder (4.9 mg). It had a molecular formula $C_{36}H_{54}O_{12}$ (calculated mass 677.3537), as deduced from its pseudo molecular precursor ion at m/z 723.3582 [M-HCOO]⁻ based on the QTOF mass spectrum. These data confirmed the presence of cardenolide glycoside. This is the first report of this cardenolide isolated from *X. undulatum*. The formula was further confirmed by the number of protons in ¹H NMR and of carbon atoms in the ¹³C NMR spectrum (**supplementary data 18, 19**). The structure of the compound (**58**) is shown in Figure 4.22.

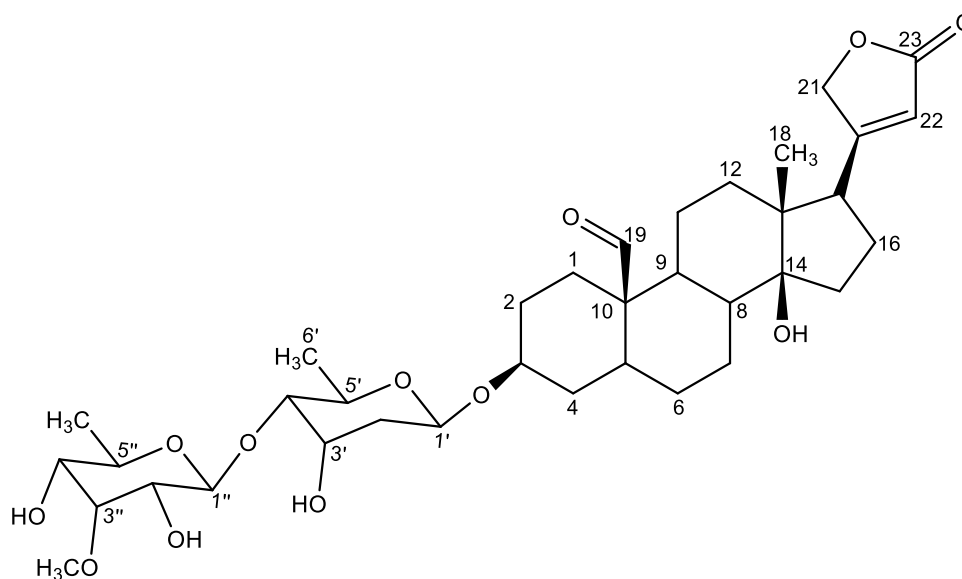


Figure 4.24: Structure of compound crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**)

The ^1H NMR spectrum (MeOD-*d*4, 400 MHz) of the compound showed the H-21 protons at δ 5.06 (1H, dd) and δ 4.90 (1H, dd) and an olefinic proton at δ 5.94 (1H, s, H-22) as part of the lactone ring which indicated the characteristic feature of the cardenolide system. The other prominent signal indicated the presence of one upfield methyl proton at δ 0.98 (3H, s, H-18). The spectrum also indicated one downfield aldehydic proton at δ 9.44 (1H, s, H-19). The ^1H NMR indicated the presence of two sugar units as one anomeric proton signal at δ 4.95 (1H, d, H-1') and other at δ 4.37 (1H, d, H-1''). The chemical shift at δ 3.48 (3H, s, 3''-OCH₃) confirmed the presence of methoxy group attached to one sugar moiety. The overall ^1H NMR confirmed the presence of cardenolide.

The ^{13}C NMR spectrum had a total of 36 signals and the assignments were done together with the DEPT 135 spectra (**supplementary data 20**). The methylene signal at δ 73.9 was assigned to C-21, δ 116.4 assigned to C-22 olefinic carbon, δ 176.1 assigned to the C-23 carbonyl group and δ 177.1 assigned to the quaternary C-20, all of which confirmed the presence of butenolide ring. Two prominent signals at δ 14.8 assigned to C-18 and at δ 206.7 assigned to C-19 indicated the position of methyl carbon and aldehydic carbon, respectively. The β hydroxyl group at C-14 was confirmed by downfield chemical shift δ 84.5. Two anomeric carbons were revealed at δ 95.7 assigned to C-1' and δ 104.7 assigned to C-1''. The carbon at δ 55.9 assigned to position 3''-OCH₃ revealed the presence of methoxy group.

The HSQC and COSY correlations (**supplementary data 21 and 22**) led to the assignment of the proton and carbon signals and the ^1H and ^{13}C NMR data of compound (**58**) corresponds to the NMR data published by [29] as shown in Table 4.6.

Table 4.4: ^1H NMR and ^{13}C NMR data for crotoxigenin 3-O- β -glucopyranoside in MeOD-*d*4 compared to literature reports for the compound in [29] in DMSO-*d*6.

Isolated crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (17)			^1H NMR and ^{13}C NMR literature data of compound (58) in DMSO- <i>d</i> 6	
Position	^{13}C (ppm)	^1H (ppm), <i>J</i> (Hz)	^{13}C (ppm)	^1H (ppm), <i>J</i> (Hz)
1	31.3	2.2, m	31.4	2.42 (dt, 13.5, 3.5, 3.5)
2	28.7	1.8, 1.6 m	32.5	2.1, 1.6 m
3	72.0	4.08 s	76.0	4.02 br, s

4	20.6	1.44 m	35.6	1.44
5	34.5	1.86 m	42.8	1.66
6	27.9	1.84, 1.53 m	30.3	1.88, 1.25 m
7	24.6	1.83, 1.59 m	28.2	1.71, 1.22 m
8	41.6	1.88 m	48.5	1.65 m
9	29.5	2.20 m	42.7	1.74 m
10	50.8	-	51.3	-
11	21.4	1.61 m	21.7	1.46, 1.19 m
12	39.4	1.54, 1.63 m	39.4	1.49, 1.35 m
13	49.6	-	49.4	-
14	84.7	-	85.0	-
15	21.7	1.77 m	26.8	2.41, 1.62 m
16	26.5	2.2, 1.9 m	27.7	2.11, 1.87 m
17	50.8	2.8 (dd, 8.7, 5.9)	50.9	2.76 (dd, 9.4, 5.4)
18	14.8	0.98 s	15.6	0.87 s
19	206.7	9.44 s	208.3	9.97 s
20	176.0	-	174.4	-
21	73.9	5.06 (dd, 18.4, 1.68), 4.9 (dd, 18.2, 1.79)	73.5	5.01 (dd, 18.2, 1.8) 4.80 (dd, 18.2, 1.8)
22	116.4	5.94 s	117.8	5.89 br.s
23	177.2	-	174.6	-
1'	95.7	4.95 (dd, 18.4, 1.4)	95.3	4.94 (dd, 1.9, 1.2)
2'	37.5	1.99, 1.77 m	37.3	1.97, 1.73 m
3'	67.4	4.30 (q, 6.17, 3.0)	66.8	4.24 (q, 6.4, 3.2)
4'	82.7	3.26 (dd, 9.4, 2.8)	82.8	3.22 (dd, 9.3, 2.9)
5'	67.1	3.87 m	67.9	3.85 m
6'	15.5	1.30 (d, 6.2)	16.5	1.36 (d, 6.5)
1' '	104.7	4.37 (d, 7.8)	103.8	4.32 (d, 7.8)
2' '	69.9	3.66 (q, 12.5, 6.7)	70.3	3.7 (dd, 9.5, 7.8)
3' '	82.9	3.15 (dd, 9.6, 3.1)	83.4	3.27 (dd, 9.5, 3.3)
4' '	68.3	3.89 m	67.8	3.22 m
5' '	70.0	3.58 (dd, 7.5, 1.9)	70.6	3.63 (dd, 6.1, 1.4)
6' '	17.1	1.32 (d, 6.2)	18.2	1.32 (d, 6.2)
3' ' -OCH ₃	55.9l	3.48 s	57.7	3.52 s

The strong HMBC correlation (**supplementary data 23**) from proton at δ 2.8 (H-17) with δ 73.9 (C-21), δ 116.4 (C-22) and δ 177.0 (C-23) indicated that the α,β -unsaturated γ -lactone to be connected at C-17 (δ 50.5); protons at δ 5.02 and δ 4.92 (C-21) correlated with δ 175.8 (C-20). The proton at δ 9.44 (H-19) correlated to δ 29.5 (C-9) and δ 34.5 (C-5) confirmed the position of aldehydic proton.

HMBC correlations also established the linkages of the glycosidic bonds and the point of attachment of the saccharide chain to the aglycone by a δ 4.95 (H-1') correlation to δ 4.02 (H-

3) and a proton at δ 4.37 (H-1'') correlation to δ 3.26 (H-4') which indicated a 1-4 linkage between the sugar units.

The two anomeric protons δ 4.95 (br dd) and δ 4.37 (d, 7.8Hz) indicated the presence of two sugar moieties (based on the large coupling you can indicate the alpha or beta linkage). The COSY correlations δ 4.95 (H-1') – δ 1.77 (H₂-2'); δ 4.30 (H-3') – δ 3.26 (H-4'); δ 3.89 (H-5') – δ 1.33 (H-6') confirmed the sugar to be digitoxose and the COSY correlations of the second sugar moiety, δ 4.37 (H-1'') – δ 3.58 (H-2''); δ 3.14 (H-3'') – δ 3.87 (H-4''); δ 3.66 (H-5'') – δ 1.30 (H-6'') confirmed the sugar to be digitalose. The chemical shifts for the digitoxose and digitalose were confirmed based the comparison to literatutre data [30],[31]. The HMBC and COSY correlations are shown in Figure 4.23.

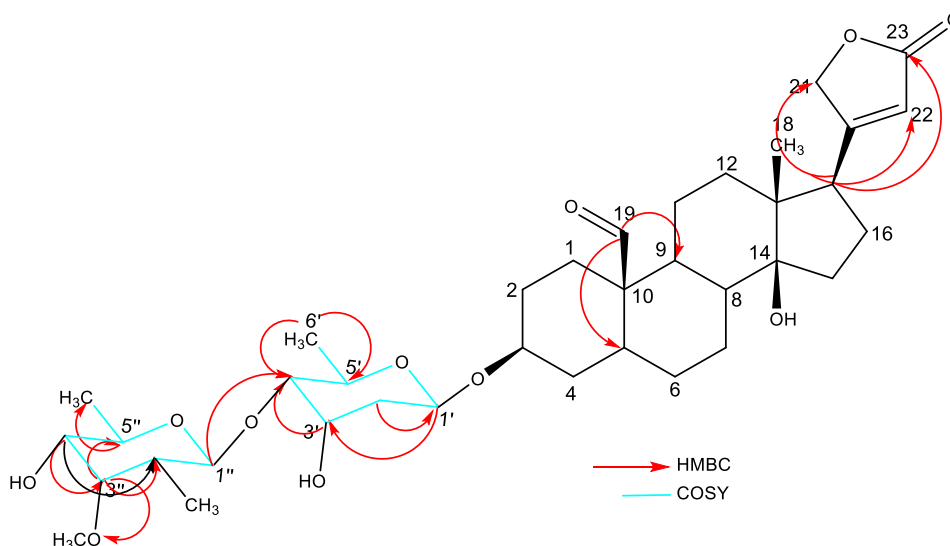


Figure 4.25: Key HMBC and COSY correlations of crotoxigenin-3-O-β-digitalopyranosyl-(1-4)-O-β-digitoxopyranoside (**58**)

Crotoxigenin-3-O-β-digitalopyranosyl-(1-4)-O-β-digitoxopyranoside, a cardenolide glycoside was originally isolated from *Alafia* sp. belongs to family Apocynaceae, a traditional medicinal plant from South of Madagascar for which the leaves of the plant are used traditionally for the treatment of malaria. This study reports the first isolation of the compound from the leaves of *X.undulatum*.

4.3.8.4. Crotoxigenin 3-O-glucopyranoside (59)

The pure compound (XU07) obtained from fraction AT-1-49N was identified as a known compound crotoxigenin 3-O-glucopyranoside (**59**) based on the NMR data, mass spectral analysis and previous published studies. It was isolated as a white crystalline solid (5 mg). It had a molecular formula of $C_{29}H_{42}O_{10}$ (calculated mass 552.2934), as deduced from its pseudomolecular precursor ion at m/z 595.2758 $[M-HCOO]^-$ based on the QTOF mass spectrum. These data confirmed the presence of cardenolide glycoside. The formula was further confirmed by the number of protons in 1H NMR and of carbon atoms in the ^{13}C NMR spectrum (**supplementary data 24, 25**). The structure of the compound (**59**) is shown in Figure 4.24.

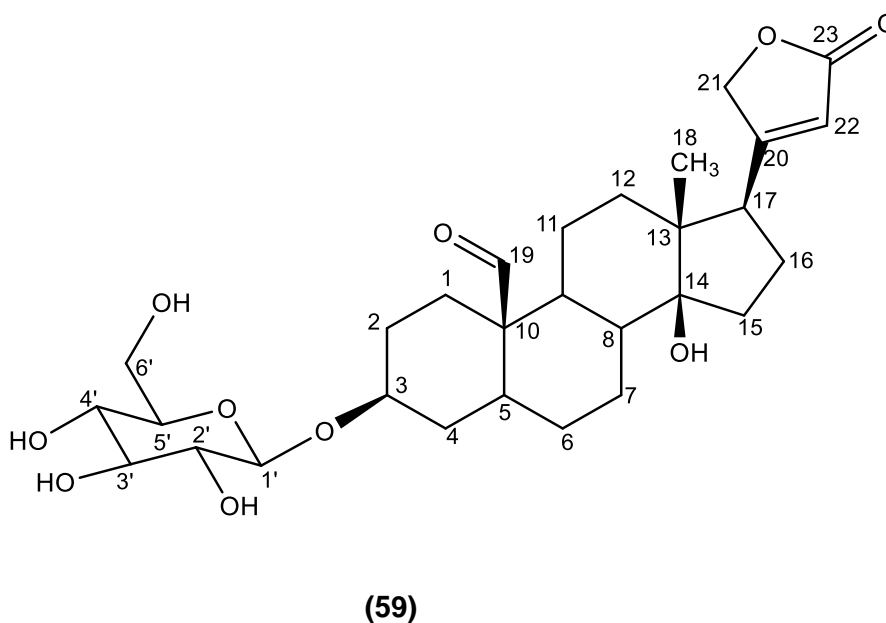


Figure 4.26: Structure of compound crotoxigenin 3-O-glucopyranoside (**59**)

The 1H NMR spectrum (MeOD- d_4 , 400 MHz) of the compound showed the H-21 protons at δ 5.04 (1H, dd) and δ 4.92 (1H, dd) and an olefinic proton at δ 5.92 (1H, s, H-22) as part of the lactone ring which indicated the characteristic feature of the cardenolide system. The other prominent signal indicated the presence of one high field methyl proton at δ 0.95 (3H, s, H-18). The spectrum also indicated one downfield aldehydic proton at δ 9.54 (1H, s, H-19) and

overall pattern of the ^1H NMR indicated that the compound was a cardenolide with one sugar unit as an anomeric proton signal was observed at δ 4.42 (1H, d, H-1').

The ^{13}C NMR spectrum had a total of 29 signals and the assignments were done together with the DEPT 135 spectra (**supplementary data 26**). The methylene signal at δ 73.6 was assigned to C-21, δ 116.4 assigned to C-22 olefinic carbon, δ 176.9 assigned to the C-23 carbonyl group and δ 175.8 assigned to the quaternary C-20, all of which confirmed the presence of butenolide ring. Two prominent signals at δ 16.1 assigned to C-18 indicated a methyl carbon and at δ 206.8 assigned to C-19 indicated the position of aldehydic carbon. The β hydroxyl group at C-14 was confirmed by downfield chemical shift δ 84.5. One anomeric carbon at δ 100.8 was assigned to C-1'.

The HSQC and COSY (**supplementary data 27, 28**) correlations led to the assignment of the proton and carbon signals. The ^1H and ^{13}C NMR data of compound (**59**) compared favorably to the NMR data based on the literature data for the compound [32] shown in Table 4.7.

Table 4.5: ^1H NMR and ^{13}C NMR data for desglucouzarin in MeOD-*d*4 compared to those reported by [32] in DMSO-*d*6.

Isolated Crotoxogenin 3-O-glucopyranoside (18)			^1H NMR and ^{13}C NMR literature data for compound (59) in DMSO- <i>d</i> 6	
Position	^{13}C (ppm)	^1H (ppm), <i>J</i> (Hz)	^{13}C (ppm)	^1H (ppm), <i>J</i> (Hz)
1	28.0	1.83 m, 1.54 m	28.03	1.85 m, 1.50 m
2	31.3	1.85 m	30.36	1.80 m
3	77.0	3.82 m	76.78	3.05 m
4	25.9	1.85 m, 1.33 m	30.20	1.82 m, 1.14 m
5	41.6	1.89 m	42.03	1.45 m
6	24.9	1.76 m	26.27	1.74 m
7	20.3	1.48 m	21.41	1.60 m
8	35.2	1.96 m	47.57	1.43 m
9	35.0	1.79 m	35.49	1.81 m
10	50.4	-	49.99	-
11	21.2	1.82 m, 1.32 m	21.41	1.60 m
12	39.4	1.51 m	42.18	1.44 m
13	49.6	-	49.19	-
14	84.5	-	83.37	-
15	32.9	1.71 m	31.59	1.94 m, 1.55 m
16	26.5	2.16 m	27.17	2.13 m
17	50.5	2.8 (q, 14.6)	51.23	2.72 m
18	14.8	0.95 s	15.54	0.69 s

19	206.8	9.54 s	209.66	9.94 s
20	175.8	-	173.96	-
21	73.7	5.04 (d, 1.65), 4.92 (dd, 18.5, 1.74)	73.48	4.90 dd
22	116.4	5.91 s	116.38	5.89 s
23	176.9	-	176.31	-
1'	100.8	4.42 d (7.80)	100.71	4.20 d
2'	73.9	3.14 (dd, 9.03, 1.07)	75.75	3.59 m
3'	76.5	3.26	73.23	2.85 m
4'	70.3	3.27 m	70.11	3.02 m
5'	76.7	3.35m	76.78	3.06 m
6'	61.4	3.87 m, 3.65 m	61.14	3.64m, 3.40m

The strong HMBC correlation (**supplementary data 29**) of proton at H-17 (δ 2.8) with δ 73.7 (C-21), δ 116.4 (C-22) and δ 176.9 (C-23) established that the α,β -unsaturated γ -lactone to be connected at C-17 (δ 50.53); the methyl protons at δ 0.95 (H-18) correlated to δ 39.4 (C-12), δ 49.6 (C-13) and δ 84.5 (C-14) which confirmed the position of the methyl group at C-18; the proton δ 9.54 (H-19) correlated with δ 35.0 (C-9) which indicated the position of the aldehydic group at C-19.

Correlations between proton δ 4.42 (H-1') with δ 77.0 (C-3) established the glycosidic linkage to be on position C-3 of the aglycone; proton δ 3.14 (H-2') with δ 76.5 (C-4') and δ 100.8 (C-1'); proton δ 3.35 (H-3') with δ 70.3 (C-4') and δ 3.27 (H-5'); proton δ 3.26 (H-4') with δ 70.3 (C-5'); proton δ 3.27 (H-5') with δ 76.5 (C-4') and δ 61.4 (C-6'); protons at δ 3.68 and δ 3.48 (H-6') with δ 70.3 (C-5') confirmed the presence of the sugar moiety.

A large coupling constant H1'/H2' (δ 4.42, 7.8 Hz) indicated a diaxial relationship for anomeric proton H1' and proton H2'. This large coupling constant also indicated the β - glycosidic linkage. The second large coupling constant H2'/H3' (δ 3.14, 9.07) confirmed that H2' was also axial. The small coupling constant H3'/H4' indicated the proton at H' 4 to be equatorial assignment. COSY correlations were also observed between H5' (δ 3.29) with two protons at H6'a (δ 3.87) and H6'b (δ 3.67). These two methylene protons are also coupled to each other and gave a set of doublet of doublet at δ 3.67 (12.0Hz, 5.2Hz) while at δ 3.87, the spin multiplicity was not clear due to overlapping of signals. The HMBC and COSY relations are shown in Figure 4.25.

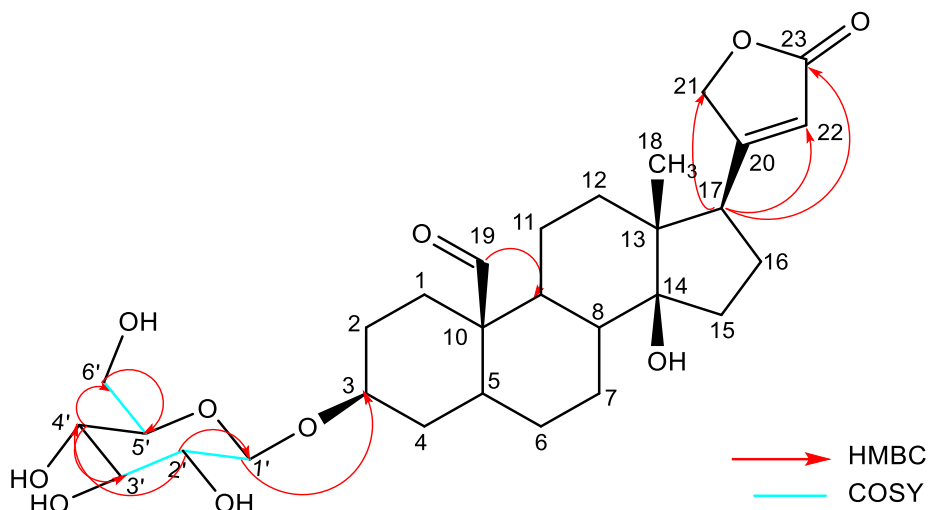


Figure 4.27: Key HMBC and COSY correlations of crotoxigenin 3-O-glucopyranoside (**59**)

4.3.8.5. Desglucouzarin (**60**)

The pure compound (XU08) obtained from fraction AT-1-49N was identified as a known compound desglucouzarin (**60**) based on the NMR data, mass spectral analysis and previous studies [25]. It was isolated as a white crystalline solid (4 mg). It had a molecular formula of $C_{29}H_{44}O_9$ (calculated mass 536.2985), as deduced from its pseudomolecular precursor ion at m/z 581.2965 $[M-HCOO]^-$ based on the QTOF mass spectrum. This data confirmed the presence of cardenolide glycoside. The formula was further confirmed by the number of protons in 1H NMR and of carbon atoms in the ^{13}C NMR spectrum (**supplementary data 30, 31**). The structure of the compound (**60**) is shown in Figure 4.26.

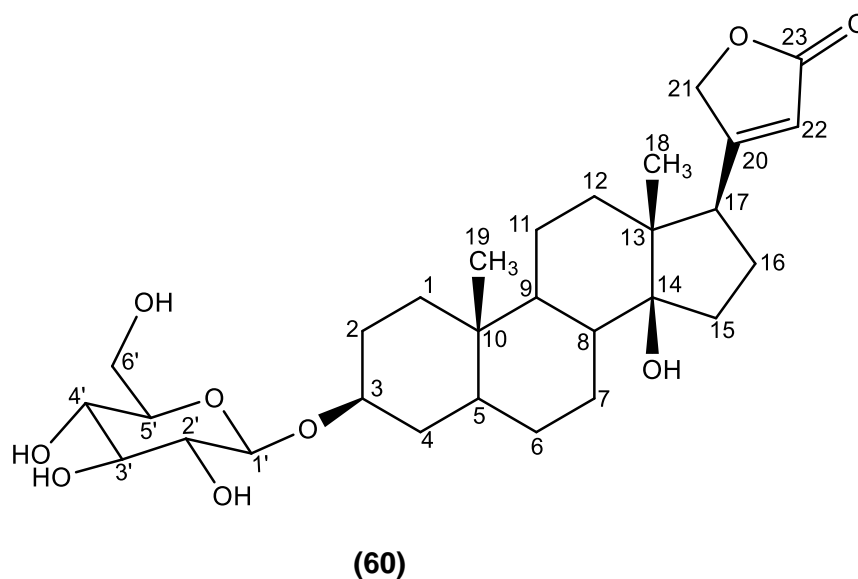


Figure 4.28: Structure of desglucouzarin (**60**)

The ^1H NMR spectrum (MeOD- d_4 , 400 MHz) of this compound showed the protons δ 5.01 and δ 4.95 (2H, dd, H-21) and an olefinic proton δ 5.92 (1H, s, H-22) attached to the lactone ring indicated the characteristic feature of cardenolide system. The other prominent signal of one high field methyl proton at δ 0.96 (3H, s, H-18), was suggestive of cardenolide nucleus together with the other high field proton at δ 0.92 (3H, s, H-19). The spectrum indicated a cardenolide with one sugar unit in which anomeric proton signal was observed at δ 4.40 (1H, dd, H-1'), where the large coupling constant (7.80 Hz) confirmed the β -coupling of the sugar unit to the aglycone.

The ^{13}C NMR spectrum had a total of 29 signals and the assignments were done together with the DEPT 135 spectra (**supplementary data 32**). The methylene signal at δ 73.7 was assigned to C-21, δ 116.8 assigned to C-22 olefinic carbon, δ 176.1 assigned to the C-23 carbonyl group and δ 177.1 assigned to the quaternary C-20, all of which confirmed the presence of butenolide ring. Two prominent signals at δ 22.3 assigned to C-18 and at δ 15.1 assigned to C-19 indicated the position of two methyl carbons. The β hydroxyl group at C-14

was confirmed by downfield chemical shift at δ 85.3. One anomeric carbon at δ 102.3 was assigned to C-1'.

The HSQC and COSY correlations (**supplementary data 33, 34**) led to the assignment of the proton and carbon signals. The ^1H and ^{13}C NMR data of compound. The ^1H and ^{13}C NMR data of compound (**60**) and the corresponding published data for the compound [25] is shown in Table 4.8.

Table 4.6: ^1H NMR and ^{13}C NMR data for desglucouzarin (**60**) in MeOD-*d*4 compared to that which is reported in DMSO-*d*6 [25]

Isolated desglucouzarin (19)			^1H NMR and ^{13}C NMR literature data of compound (60) in DMSO- <i>d</i> 6	
Position	^{13}C (ppm)	^1H (ppm), J (Hz)	^{13}C (ppm)	^1H (ppm), J (Hz)
1	34.6	1.07, 1.82 m	36.6	0.92, 1.75 m
2	33.9	1.65, 1.89 m	29.0	1.75 m
3	78.2	3.80 m	76.4	3.05 m
4	32.1	1.77 m	33.9	1.14, 1.63 m
5	41.6	1.45 m	43.7	0.95 m
6	26.9	1.41 m	28.5	1.13, 1.23 m
7	26.6	1.91 m	27.2	0.92, 1.94 m
8	41.8	1.67 m	40.8	1.42 (dt, 2.93)
9	36.2	1.83 m	49.4	0.87 (dt, 2.93)
10	34.7	-	35.4	-
11	20.8	1.49, 1.83 m	20.8	1.18, 1.4 m
12	39.6	1.54 m	38.8	1.3, 1.42 m
13	49.8	-	48.6	-
14	85.3	-	83.6	-
15	31.9	1.79, 2.2 m	32.1	1.55t, 1.92t
16	26.4	2.22, 1.86 m	26.3	1.75, 1.96 m
17	50.8	2.88 (dd, 9.1, 5.9)	50.1	2.72 (dd, 9.6, 5.4)
18	22.3	0.96 s	15.7	0.72 s
19	15.1	0.92 s	12.0	0.75 s
20	177.1	-	176.4	-
21	73.7	4.95, 5.01 (dd, 1.49, 18.4)	73.2	4.88 m, 4.95 (dd, 1.47, 18.32)
22	116.8	5.92 s	116.2	5.89 s
23	176.1	-	173.9	-
1'	100.8	4.4 (d, 7.80)	100.6	4.2 (d, 8.06)
2'	73.9	3.18 (dd, 9.07, 1.12)	73.5	2.85 m
3'	76.7	3.36 m	76.8	3.10 m
4'	70.2	3.28 m	70.1	3.01 m
5'	76.5	3.28 m	76.4	3.50 m
6'	61.3	3.87 m, 3.67 (ddd, 1.9, 5.2, 12.0)	61.1	3.35 m, 3.65 (ddd, 2, 5.4, 12.0)

The strong HMBC correlation (**supplementary data 35**) of proton at δ 2.88 (H-17) with δ 73.7 (C-21), δ 116.8 (C-22) and δ 176.1 (C-23) indicated that the α,β -unsaturated γ -lactone was connected at δ 50.8 (C-17). The methyl proton at δ 0.96 (H-19) correlated to δ 41.6 (C-5) and δ 34.7 (C-10) which confirmed the position of the methyl protons at C-19. Correlations between the anomeric proton H1' (δ 4.4) with δ 78.2 (C-3) established the glycosidic linkage and confirmed that the sugar unit is attached to C-3.

A large coupling constant H1'/H2' (δ 4.4, 7.8 Hz) indicated a diaxial relationship for anomeric proton H1' and other proton H2'. This large coupling constant also indicated the β -pyranose form for glycoside by verifying the β -orientation of glycoside. The second large coupling constant H2'/H3' (9.07 Hz) confirmed that H2' was also axial. The small coupling constant H3'/H4' indicated the proton at H4' to be equatorial assignment. COSY correlations were also observed between δ 3.29 (H5') with two protons at δ 3.87 and δ 3.67 (H6'). These two methylene protons also coupled to each other and gave a set of doublet of doublet at δ 3.67 (12.0 Hz, 5.2 Hz) while at δ 3.87, the spin multiplicity was not clear due to overlap of signals. The HMBC and COSY relations are shown in Figure 4.27.

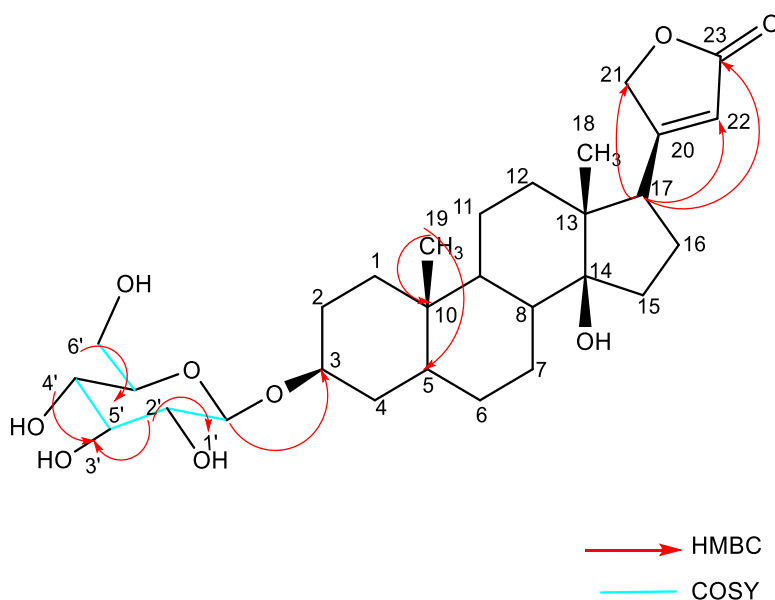


Figure 4.29: Key HMBC and COSY correlations of desglucouzarin (**60**)

The compound was previously obtained from *Asclepias subulata*, *Asclepias syriaca* L. and *Asclepias linaria* Cav., and showed significant antiproliferative activity on MCF-7, T47D, SK-

BR-3 and Hs578T cells. It showed a strong antiproliferative activity with an IC_{50} value 0.90 ± 0.02 [32]. This compound was also isolated from aerial parts of *Pergularia tomentosa* L. [25]. The compound was also previously reported in *X. undulatum* [2].

4.9. The effect of the isolated compounds on A β 42 production

The four compounds acetylated glycosydated crotoxigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), desglucouzarin (**60**) isolated from *X. undulatum* leaf extract significantly decreased the A β 42 levels as compared to the negative control by $75.04 \pm 0.85\%$, $74.66 \pm 0.1\%$, $65.55 \pm 0.19\%$ and $69.15 \pm 0.25\%$, respectively as shown in Figure 4.28. The compound crotoxigenin 3-O-glucopyranoside (**59**) did not show any significant reduction in A β 42 levels. The results clearly indicated that the isolated from compounds from the active fractions were responsible for the effect of A β 42 production in *X.undulatum* extract.

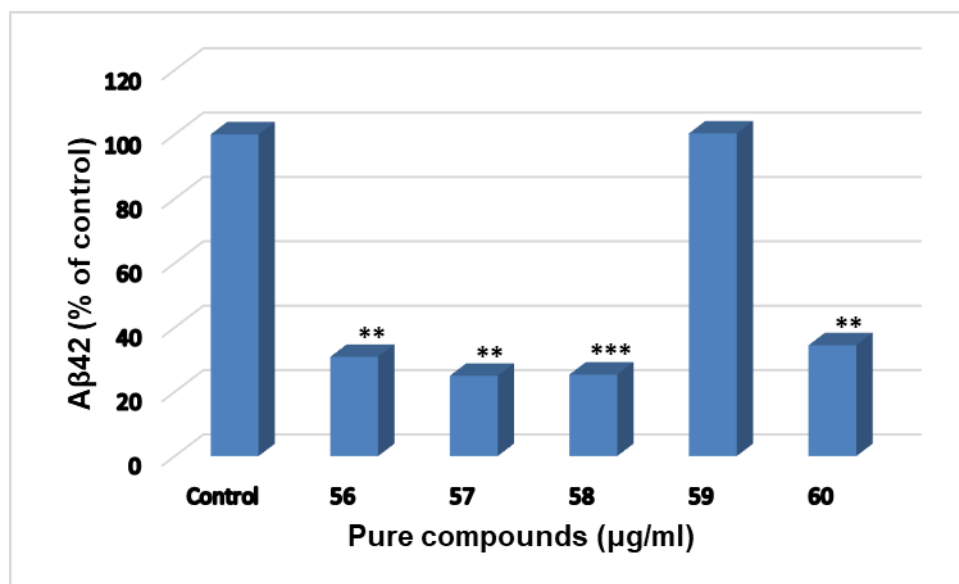


Figure 4.30: Change in the level A β 42 by compounds of *X. undulatum* extract. Cells were incubated with $10\mu\text{M}$ concentrations of compounds acetylated glycosydated crotoxigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), crotoxigenin 3-O-glucopyranoside (**59**) and desglucouzarin (**60**) for 8 h. The level of A β 42 was measured from the conditioned media by using specific ELISA. The level of A β 42 was decreased by compounds **56**, **57**, **58** and **60** ($n=4$), **, $P < 0.01$; ***, $P < 0.001$.

The active compounds were then treated at different concentrations in APPsw-transfected HeLa cells for 8 h. Compound **(56)** significantly decreased the levels of A β 42 by 35.5 ± 2.1 , 62.83 ± 1.6 %, 71.8 ± 2.4 % and 76.9 ± 4.4 % at concentrations 1, 2.5, 5 and 10 μ M, respectively (Figure 4.29). Compound **(57)** significantly decrease the A β 42 levels by 24.7 ± 0.0 , 37.6 ± 0.1 , 49.5 ± 0.0 and 52.3 ± 1.9 at 1, 2.5, 5 and 10 μ M respectively (Figure 4.30). Compound **(58)** significantly decreased the levels of A β 42 by 19.1 ± 1.2 %, 48.6 ± 7.2 %, 68.1 ± 3.1 % and 74.5 ± 0.9 % at concentrations 0.025, 0.05, 0.1 and 0.25 μ M, respectively (Figure 4.31) while the levels of A β 42 was significantly decreased by 31.2 ± 6.2 %, 63.8 ± 1.1 %, 73.2 ± 3.2 % and 74.9 ± 5.4 % at 1, 2.5, 5 and 10 μ M for compound **(60)**, respectively (Figure 4.32).

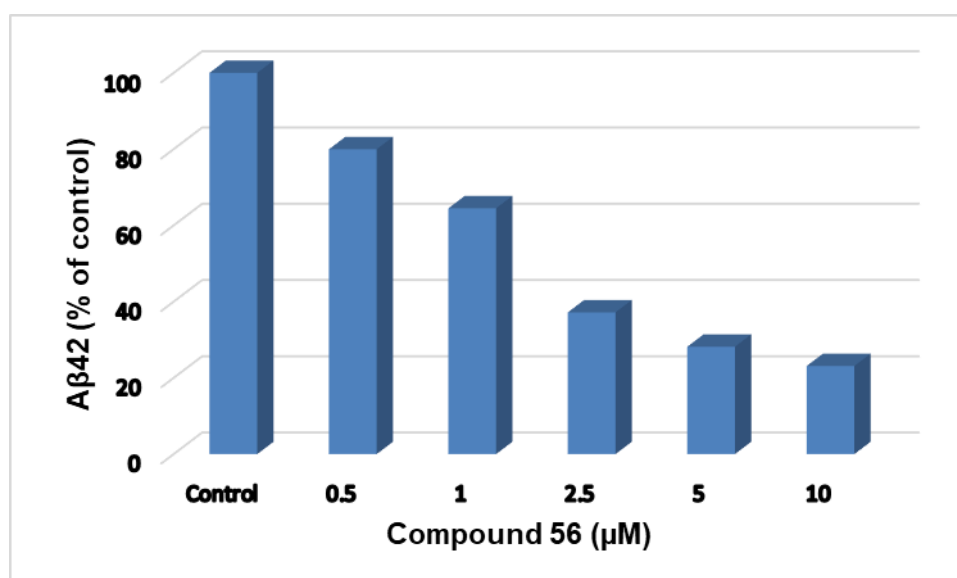


Figure 4.31: Change in the level A β 42 by acetylated glycosydated crotoxogenin **(56)**. Cells were incubated with indicated concentrations of compound **56** for 8 h. The level of A β 42 was measured from the conditioned media by using specific ELISA. The level of A β 42 was decreased by compound **56** in a dose dependant manner (n=4), **, P < 0.01; ***, P < 0.001.

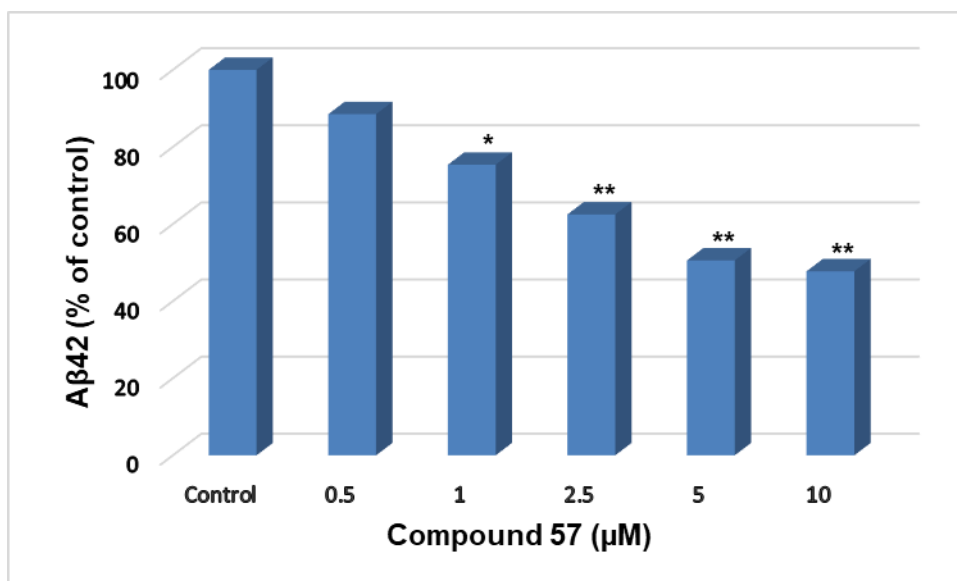


Figure 4.32: Change in the level Aβ42 by acetylated glycosydated crotoxogenin (**57**). Cells were incubated with indicated concentrations of compound **57** for 8 h. The level of Aβ42 was measured from the conditioned media by using specific ELISA. The level of Aβ42 was decreased by compound **57** in a dose dependant manner (n=4), **, P < 0.01; ***, P < 0.001.

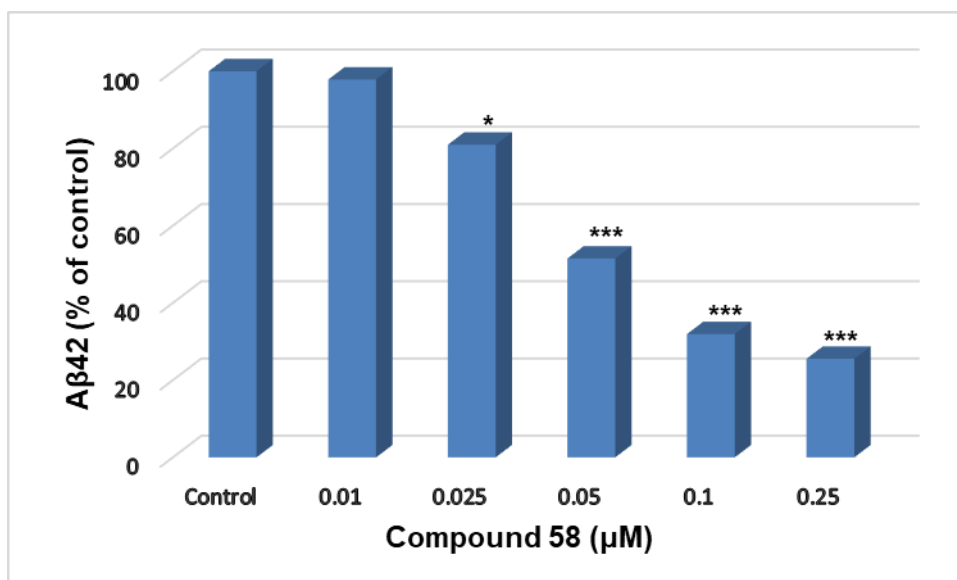


Figure 4.33: Change in the level Aβ42 by crotoxigenin-3-O-β-digitalopyranosyl-(1-4)-O-β-digitoxopyranoside (**58**). Cells were incubated with indicated concentrations of compound **58** for 8 h. The level of Aβ42 was measured from the conditioned media by using specific ELISA. The level of Aβ42 was decreased by compounds **58** in a dose dependant manner (n=4), **, P < 0.01; ***, P < 0.001.

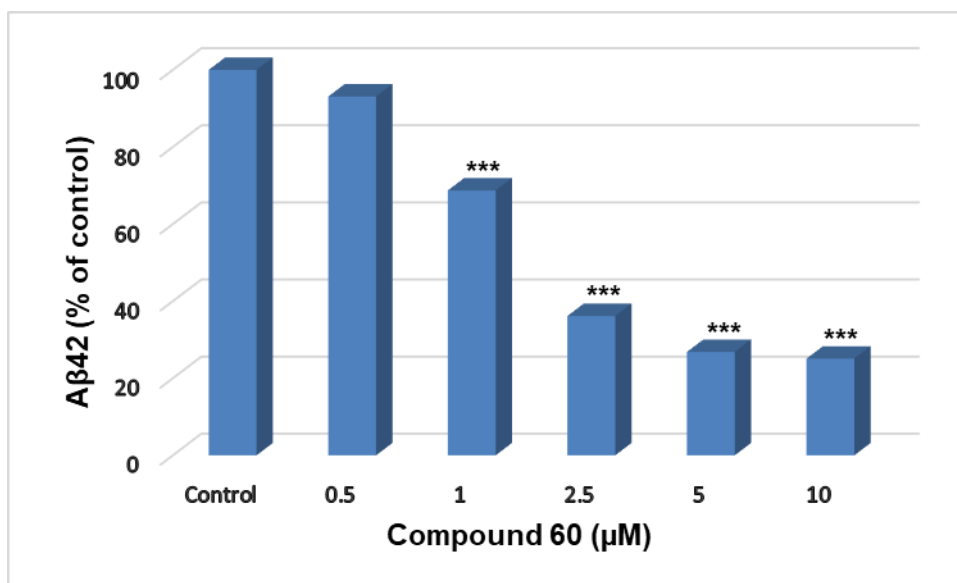
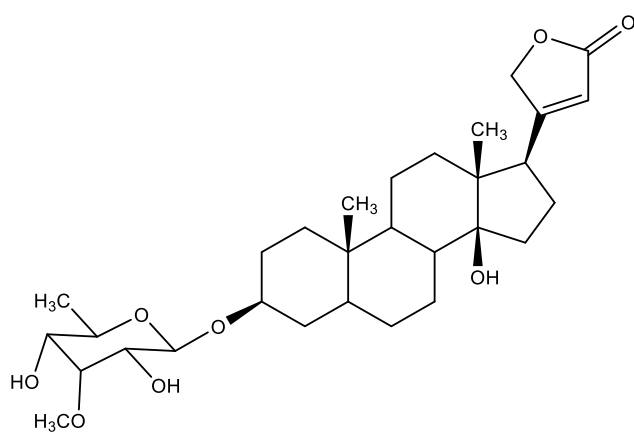


Figure 4.34: Change in the level Aβ42 by compound desglucouzarin (**60**). Cells were incubated with indicated concentrations of compound **60** for 8 h. The level of Aβ42 was measured from the conditioned media by using specific ELISA. The level of Aβ42 was decreased by compounds **60** (n=4), **, P < 0.01; ***, P < 0.001.

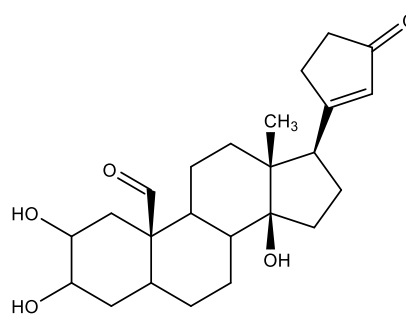
All these pure compounds acetylated glycosydated crotoxigenin (**56**), xysmalogenin-3, β-D-glucopyranoside (**57**), crotoxigenin-3-O-β-digitalopyranosyl-(1-4)-O-β-digitoxopyranoside (**58**), crotoxigenin 3-O-glucopyranoside (**59**) and desglucouzarin (**60**) isolated from the leaves of *X. undulatum* have examined for the first time for Aβ42 reduction in the present study. Four of the compounds, except crotoxigenin 3-O-glucopyranoside (**59**), showed significant activity. Since all the compounds have the butenolide ring attached to C-17 of the aglycone as part of their structures, it appears that this moiety may not be the pharmacophore responsible for the activity especially the fact that compound (**59**) had no activity at the highest test concentration of 10 μM (Figure 4.28). This is suggestive that the remaining units of the molecules i.e. the aglycone unit and the carbohydrate units a play a more significant the role in the reduction of Aβ42. This is promising as any future work should involve the removal of the butenolide ring and testing the glycosidated aglycones for their potential to reduce Aβ42. The added advantage of this is that the cardiotoxicity could be removed as it is known that the butenolide ring is mainly responsible for toxicity [33], making the compounds more pharmaceutically attractive. The compound (**58**), crotoxigenin-3-O-β-digitalopyranosyl-(1-4)-O-β-

digitoxopyanoside was for the first time isolated from this plant while compound **(56)**, acetylated glycosydated crotoxogenin has been reported as the novel cardenolide. The compound **(58)** was the most promising of the four active compounds since as it was tested at a very low concentrations (test concentration of 0.025 μM) compared to the other compounds (lowest test concentration 1 μM) and showed significant reduction of A β 42 at 0.025 μM , at least 40 time more potent than any of the other cardenolides.

In a brain slice assay for ischemic stroke, neriifolin **(60)** showed a significant neuroprotective effect included good delayed therapeutic potential exceeding 6 h as compared to no delay [34]. In a docking model generated for identifying the AChE inhibition, a cardenolide, calotropagenin **(61)** isolated from *Calotropis procera*, was found to exhibit significant binding affinity towards the AChE with a significant Grid based Ligand Docking with Energetics (GLIDE) score of -13.07 as compared to the GLIDE score of -11.23 for Aricept AChE complex [35]. In another study, a structurally similar cardenolide, odoroside H **(62)**, isolated from the leaves of *Nerium oleander* showed CNS depressant activity in *in vivo* studies. These compounds showed decrease in motor activity in mice at a dosage of 25 mg/kg [36].



(60)



(61)

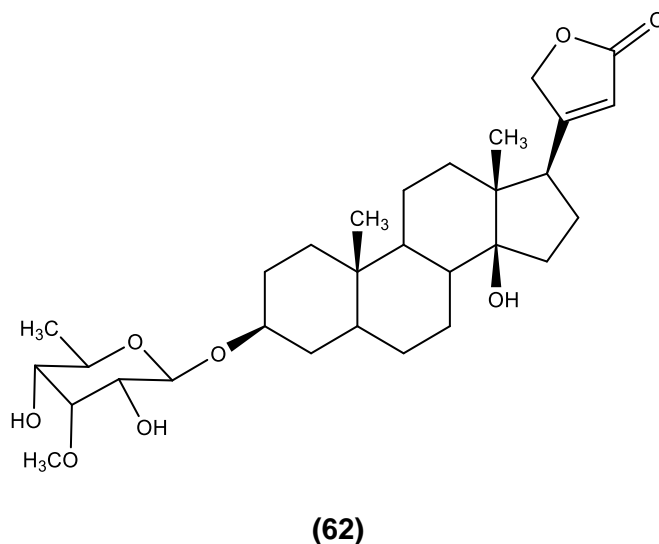


Figure 4.35: Structure of neriifolin (**60**), calotropagenin (**61**) and odoroside H (**62**)

Neriifolin (**60**) and odoroside (**62**) are structurally similar to the cardiac glycosides such as digitoxin, digoxin and ouabain which are medically recognised because of their Na^+/K^+ -ATPase inhibition action [34]. The isolated compounds in the present study also have similar skeletal structures. Therefore, it indicates that the isolated compounds in the present study can also be able to inhibit the AChE and can be examined for the *in vivo* animal models to check its ability to be develop into an Alzheimer's disease.

There is limited research showing the efficacy of cardenolides for the treatment of Alzheimer's disease with no reported research showing activity of cardenolides towards the reduction of $\text{A}\beta_{42}$ which is an important factor for the treatment of Alzheimer's disease. The current study has shown for the first time that cardenolides from *X. undulatum* can significantly reduce the levels of $\text{A}\beta_{42}$.

4.4. Conclusion

X. undulaum has many traditional uses such as treatment of diarrhoea, dysentery, dysmenorrhoea, stomach cramps, wounds and indigestion. *X.undulatum* is a commercialised plant in Europe for the treatment of diahrroea and traded as Uzara. Although the plant has

been extensively researched and well-documented for its antidepressant effect, acetylcholinesterase inhibitory effects and selective serotonin reuptake inhibitor (SSRI) activity, there is only limited literature available showing its inhibitory effects for the reduction of A β 42, an important factor in the treatment of Alzheimer's disease. In the present study, the DCM:MeOH leaf extract significantly reduced the A β 42 production by $49.8 \pm 4.0\%$, $71.4 \pm 1.6\%$ and $72.9 \pm 1.1\%$ at concentrations 1, 5 and 10 $\mu\text{g/ml}$, respectively. To the best of our knowledge, this study is the first report for the reduction of A β 42 as a treatment for Alzheimer's disease. The active leaf extract was then analysed on UPLC-QTOF-MS to identify the main classes of compounds. MS analysis revealed the presence of cardenolides and flavonoids. The DCM:MeOH leaf extract was subjected to bioassay guided fractionation using silica gel column chromatography. A total of 15 fractions were collected after the TLC analysis and combining fractions with common compounds based on their R_f values. Of these, two fractions AT-1-49N and AT-1-49O were found to be active towards the reduction of A β 42. These fractions were analysed by UPLC-QTOF-MS to identify the active ingredients. Cardenolide glycosides identified in both the active fractions through the MS analysis. These fractions were further purified using mass directed purification using preparative HPLC. Three peaks labelled as XU01 (637.2850), XU03 (579.2822) and XU04 (723.3583) and two peaks XU07 (595.2758) and XU09 (581.2965) were collected from the two active fractions. The structures of these compounds were elucidated using mass spectral analysis and ^1H , ^{13}C NMR and 2D NMR. The compounds were identified as acetylated glycosylated crotoxigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), crotoxigenin-3-O-glucopyranoside (**59**) and desglucouzarin (**60**) and were tested for A β 42 reduction. The four cardenolide glycosides, acetylated glycosylated crotoxigenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), desglucouzarin (**60**) significantly decreased the A β 42 levels at 10 μM as compared to the negative control by $75.04 \pm 0.85\%$, $74.66 \pm 0.1\%$, $65.55 \pm 0.19\%$ and $69.15 \pm 0.25\%$, respectively. The compound crotoxigenin-3-O-glucopyranoside (**59**) did not show any reduction in A β 42 levels. Dose response studies

showed that the A β 42 reduction for compound **(58)** potently reduced A β 42 levels at a low concentration of 0.025 μ M clearly indicating that it was the most potent compound of all the isolated cardenolide glycosides. All compounds were structurally similar and contained the characteristic lactone ring. As compound **(59)** was inactive, this indicated that the lactone ring may not be responsible for A β 42 reduction. Further research could focus on the reduction of the lactone ring in the compound **(58)** and further bioassaying of the analogues to measure their A β 42 reduction potential. The overall bioassay-guided fractionation was successful for the leaf extract of *X.undulatum* as the pure active ingredients were isolated and identified, often a major challenge to the natural product chemist.

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Chapter 5

Experimental

5.1. Materials

The plants under investigation were identified by the H.G.W.J. Schweickerdt Herbarium (PRU), University of Pretoria. The 18 selected plant species *Centella asiatica*, *Heteromorpha arborescens*, *Mondia whitei*, *Stapelia gigantea*, *Tabernaemontana elegans*, *Cussonia spicata*, *Cussonia paniculata*, *Bulbine natalensis*, *Catha edulis*, *Cotyledon orbiculata*, *Schotia brachypetala*, *Tetradenia riparia*, *Trichilia dregeana*, *Plumbago auriculata*, *Ruta graveolens*, *Zanthoxylum capense*, *Buddleja salvifolia*, *Cissus quadrangularis* were obtained from the Pretoria University gardens (University of Pretoria, cnr Lynnwood Road and Roper Street, Hatfield, Pretoria, South Africa); one plant, *Xysmalobium undulatum* was collected from the experimental farm of University of Pretoria; two plants, *Commelina africana* and *Ziziphus mucronata* were collected from Dinokeng Nature Reserve Pretoria (N1 North, Hammanskraal, 0400). Voucher specimens were preserved at H.G.W.J. Schweickerdt Herbarium (PRU), University of Pretoria.

For extraction and silica gel column chromatography, analytical grade dichloromethane ($\geq 99\%$) (Merck), methanol ($\geq 99.5\%$) (Merck), hexane ($\geq 99\%$) (Sigma-Aldrich), and ethyl acetate (99.5%) (Sigma-Aldrich) were used. For LC-MS and preparative HPLC-MS, the solvents used were ultra-pure HPLC grade methanol/ water/ acetonitrile (Romil-UpSTM, Microsep, South Africa), formic acid (99% purity) (Thermo Scientific, South Africa), ammonium hydroxide $\geq 25\%$ in H₂O which was the eluent additive (Fluka® Analytical, Sigma-Aldrich, South Africa). Vanillin (Analytical grade, Radchem) was used for staining TLC plates. Standards isoquercetin, quercetin and rutin were purchased from Sigma Aldrich, South Africa. Silica gel 60 (0.063-0.20 mm) was used for column chromatography and Silica gel plates 60 F254 (Merck, South Africa) was used for analysis and visualized using a UV Detector at 254

and 360 nm (Spectroline®, Model ENF-240/FE, Spectronics Corporation Westbury, New York, USA).

5.2. Extraction process

Plant parts were cut into small pieces and then dried in oven (PROLAB) at 60°C. The dried material was ground to fine coarse powder using a laboratory mill (POLYMIX® PX-MFC 90 D/ Kinematika). The dried ground material (50 g) was placed separately in an Erlenmeyer flask (1 L). A mixture of dichloromethane and methanol (200 ml) in a 1:1 ratio was added to the flask. The plant material was extracted continuously by shaking using a magnetic stirrer for 8 h. The extraction process was repeated three times. The dried leaf material of *X. undulatum* (200 g) was separately placed in an Erlenmeyer flask (2 L) and extracted with the dichloromethane and methanol (1000 ml) in a 1:1 ratio. The plant material was extracted 3 times under similar conditions.

All the plant extracts were then filtered separately through a Whatman No.1 (GE Healthcare, Life Sciences) using a Buchner funnel connected to a vacuum pump. All the filtrates were combined separately and placed in a round bottom flask. The plant extract was then concentrated using a rotary evaporator to dryness and the extracts weighed to calculate the extraction yield. The plant residue was discarded.

5.3. Chromatographic techniques

5.3.1. Thin layer chromatography

TLC was performed on precoated Kieselgel 60 with florescent indicator UV₂₅₄ (MACHEREY-NAGEL), 0.20 mm thick, with glass support. The TLC plates were examined under a short wavelength, 254 nm UV light and subsequently by using a specific spray reagent. The following mixture of solvents were used as eluting solvents for the TLC analysis and column chromatography for the plant extracts of *X. undulatum* in order to attain the improved separation: Hexane:EtOAc (95:5), Hexane:EtOAc (90:10), Hexane:EtOAc (80:20),

DCM:MeOH (95:5), DCM:MeOH (90:10), DCM:MeOH (80:20), DCM:EtOAc (95:5), DCM:EtOAc (90:10), DCM:EtOAc (80:20), EtOAc:MeOH:HCOOH:H₂O (8:1:1:2). Vanillin staining was used to reveal the presence of compounds. The staining reagent was prepared by adding 6 g vanillin in a 100ml container. Ethanol (95 ml, 96%) was added to the vanillin to give a clear solution followed by careful addition of 1.5 ml of concentrated sulfuric acid. The solution was stored in the dark. TLC plates were dipped into the vanillin stain and dried with a hair dryer for visualization.

5.3.2. UPLC-QTOF-MS analysis

The crude extracts and fractions were analyzed using a Waters Acquity UPLC system (Waters corp., MA USA), equipped with a binary solvent delivery system and an autosampler. The instrument was centrally operated by MassLynx 4.19 software (Waters Inc., Milford, Massachusetts, USA) for data acquisition. The sample (1mg/ml) was prepared by dissolving dried extract, fraction or pure compounds in 100% methanol. The solution was centrifuged at 10 000 g for 10 minutes to remove the particulates. The separation of compounds was achieved on Waters BEH C18 1.7 μ m particle size (2.1 mm \times 100 mm). The column temperature was maintained at 40°C. The mobile phase consisted of solvent A: water with 0.1% formic acid and solvent B: methanol with 0.1% formic acid. The gradient elution was optimized as follows: 3% B (0-0.1 min), 100% B (0.3-14.00 min), 100% B (14.00-16.00 min), 3% B (16.00-16.50 min), 3% B (16.50-20.00 min). The flow rate was 0.3 ml/min for the entire run, giving a total run time of 20 minutes and the injection volume was 5 μ l. A Waters Synapt G2 high definition QTOF mass spectrometer equipped with an ESI source was used to acquire negative ion data. The instrument was calibrated by direct infusion of 5 nM sodium formate solution at a flow rate of 10 μ l/min over a mass range of 50-1200 Da.

The following MS source parameters were set for both positive and negative mode: source temperature 100 °C, sampling cone 15 V, extraction cone 4.0 V, desolvation temperature 400 °C, cone gas flow 10.0 L/h, desolvation gas flow 700 L/h, capillary 2.0 kV. Both positive and

negative modes were obtained, but the results were analyzed using the negative mode as the higher intensity peaks were obtained in the negative mode. A 2 ng/μl leucine enkephalin solution was used as lockspray solution throughout all the acquisitions. It was constantly infused at a rate of 3 μl/min through a separate orthogonal ESI probe to compensate for experimental drift in mass accuracy. The trap collision energy was 28 V.

Compounds were tentatively identified based on their accurate mass generated from MassLynx V 4.1, iFit value, MS/MS fragmentations (product ions). The electronic databases Metlin, Metfusion, Pubchem, ChEMBL, Chemical Entities of Biological Interest (ChEBI) Massbank, Reaxys, Scifinder and Scopus were used for the tentative identification of compounds. The compounds were identified by comparing the mass fragmentation pattern generated by the instrument with that in electronic data bases and literature.

A novel software, integrated MassLynx with UNIFI (Waters UNIFI Scientific Library Component Listing version: 01) informatics was also used for identifying the compounds present in crude extract. This software analyses the chromatographic and mass spectral components of the extract or samples by using the different libraries including UNIFI Toxicology Library, UNIFI Pesticide Library, UNIFI Traditional Medicine Library (Library type, Green Tea and Chinese Traditional Medicine) incorporated in UNIFI software. The Chinese Traditional Medicine library contains more than 6000 compounds isolated from 600 herbs [1]. In order to identify the compounds, the raw data from the MassLynx was imported to UNIFI software where the software integrates the data acquisition, data mining and generates a report through library searching. The UNIFI data analysis automatically matches the compound information collected from MassLynx with the Chinese Traditional Medicinal library and gives possible chemical identifications. The software provides the elemental composition to generate the possible formula for accurate mass peak based on fragmentation pattern matching. It also provides the i-Fit confidence score and a formula with a score greater than 0.8 given as a priority in the list of possible identifications. Further, the identified compounds with a mass difference of less than 5 mDa between the theoretical and instrument exact mass reading were selected for further identification and are provided with the detailed information including

compound name, molecular formula, average molecular mass, monoisotopic retention time, exact mass error in mDa and ionization mode.

5.3.3. Silica gel column chromatography

For the column chromatography, a glass silica column of 107 cm length and 7 cm internal diameter with a reservoir of 2 L capacity was used. The column was packed with a slurry suspension formed by mixing 1500 g of dried silica gel (0.063-0.2 mm/70-230 mesh) with hexane. The dried DCM:MeOH (1:1) *X. undulatum* leaf extract (17 g) was re-dissolved in the same solvent system and mixed with 12g of silica gel 60. The mixture was stirred gently with the help of glass rod to prepare a suspension. Then the solvent was gently evaporated by using a rotary evaporator until the sample was dry. The resulting powdered sample was poured slowly onto the bed of the silica gel layer and allowed to settle. Initially DCM:Hexane (50:50 to 80:20) solvent system was used as the eluent. Later, the polarity was increased to DCM (100%) and DCM:MeOH (95:5 to 70:30) as the eluants with the solvent flow rate of 2 ml/min. The separation provided a total of 46 fractions. These fractions were further analysed by TLC using silica gel plates (60 F₂₅₄) and visualised under UV light at 254nm. Pasteur pipettes were used for spotting the samples. The TLC plates were also dipped in the vanillin stain to detect the compounds. The TLC plates were dried by using a hair dryer after staining. After visualisation, the similar fractions were combined based on their R_f values. After combining the similar fractions, 15 fractions (AT-1-49A to AT-1-49O) were obtained and two fractions (AT-1-49N and AT-1-49O) were selected for further purification using prep HPLC-MS based on their promising biological activity.

5.3.4. Isolation of compounds using preparative HPLC-MS

Isolation of compounds from the active fractions (AT-1-49N and AT-1-49O) of *X. undulatum* DCM:MeOH leaf extract was performed on Waters chromatographic system with Waters PDA (2998) and MS detector (Waters, Milford, MA, USA). For the sample preparation, two active fractions (70 mg each) were separately dissolved in HPLC grade methanol (100 ml). The

solution was centrifuged for 10 min and then filtered through a 0.22 μm (Millipore) membrane syringe filter unit. Six major peaks labelled as XU01 to XU06 from sample AT-1-49N and two major peaks labelled as XU07 and XU08 from AT-1-49O were targeted for isolation. The injection volume was 150 μl per injection.

The chromatographic conditions were optimized on a preparative column to achieve well resolved chromatograms. The separation was achieved on an X-Bridge preparative C18 column (19 \times 250 mm, i.d., 5 μm particle size, Waters). The mobile phase consisted of 0.1% ammonia in water (solvent A) and acetonitrile (solvent B) at a flow rate of 20ml/min. Gradient elution was applied as follows: 95% A : 5% B for 1 min, 50% A : 50% B (1:00-8:00 min), 15% A : 85% B (8:00-13:00 min), 95% A : 5% B (13:00-13:20 min). Injection volume was 150 μl . The total run time was 20 minutes. Data were collected using MassLynx4.1TM (Waters, USA) software. The preparative HPLC system was interfaced with QDa mass spectrometer (Waters, USA) and the negative ion mode selected. The probe temperature and source temperature were set at 600 $^{\circ}\text{C}$ and 120 $^{\circ}\text{C}$, respectively. The capillary and cone voltage were set to 800 and 10 V, respectively. Data were collected between 100 and 650 m/z . The pure compounds were collected using a fraction collector and subsequently combined and concentrated speed vacuum. The compounds were later analyzed by UPLC-MS and their structures are confirmed by NMR.

5.4. Bioassay screenings

5.4.1. APP^{sw}-transfected HeLa cell culture

HeLa cells stably transfected with Swedish mutant form of APP (APP^{sw}) were cultured in Dulbecco's Modified Eagle Medium (DMEM), supplemented with 10% heat-inactivated fetal bovine serum (FBS), 1% penicillin, 100 $\mu\text{g/ml}$ streptomycin, 260 $\mu\text{g/ml}$ Zeocin and 400 $\mu\text{g/ml}$ G418 at 37 $^{\circ}\text{C}$ in a humidified atmosphere of 5% CO₂ and 95% air.

Fetal bovine serum is vitally important as a source of growth and adhesion factors, hormones, lipids and minerals for the cell culture in basal media. It also regulates the cell membrane

permeability and serve as a carrier. Penicillin/ Streptomycin is used to avoid contamination. Antibiotics Zeocin and G418 are used for the selection of APPsw-transfected HeLa cell.

5.4.2. Cell culture protocol

Spent cell culture media was removed and discarded from culture vessel. Cells were washed using PBS (5 ml). Wash solution was gently added to the side of vessel opposite to the attached cell layer to avoid disturbing the cell layer. The wash step was used to remove any traces of serum, calcium and magnesium that would inhibit the action of the dissociation reagent. The wash solution was removed from the cell culture. The pre-warmed dissociation reagent such as trypsin-EDTA (1 ml) was added to the side of flask. Culture vessel was then incubated at room temperature for approximately 2-3 min. The actual incubation time varies with the cell line used. The cells were observed under the microscope for detachment. An equivalent of 5 ml of pre-warmed complete growth medium was added to inactivate trypsin-EDTA. The growth medium was dispersed by pipetting over the cell layer surface several times. The cells were transferred to a 15 ml conical tube and centrifuged at 1000 g for 3 min. The centrifuge speed and time also vary with the cell type used. The cell culture media was removed. The cell pellet was resuspended in 5 ml of pre-warmed complete growth medium and a sample was removed for counting. To subculture after 3 days, the cells (0.5 ml) were transferred to a T75 flask with cell culture media 10-12 ml. To subculture after 4 days, the cells (0.3 ml) were transferred to a T75 flask with 10-12 ml culture media. The total number of cells were determined by using a hemocytometer.

5.4.3. Sample preparation

The samples were prepared by dissolving the stock solution of the compound in DMSO, kept at -20°C. Before each experiment, the solution was diluted to the final concentration in fresh media. In order not to effect cell growth, the final DMSO concentration did not exceed 0.5 % in all experiments.

5.4.4. Cell Viability Measurement

Cell viability was measured by MTT assay using an EZ-Cytox kit (Daeil Lab Co, Ltd, Republic of Korea) according to the manufacturer's instructions. APPsw-transfected HeLa cells in a 96 well plate were cultured with sample solubilized in DMSO for 8 h, and then incubated with EZ-Cytox solution for 1 h at 37°C. The absorbance was measured on a microplate reader (BIO-TEK® DowerWave XS, Winooski, VT, USA) with a software capable of measurement at or near 450 nm. The results were expressed as the percentage of MTT reduction relative to the absorbance of control cells.

5.4.5. sAPP α , sAPP β , A β peptide assay

APPsw-transfected HeLa cells at 80% confluence in a 35 mm dish were cultured with the samples solubilized in dimethyl sulphoxide (DMSO) for 8 h in serum free medium. The conditioned medium was analyzed by a sandwich ELISA (Invitrogen) for detection of A β 42 and A β 40 according to the supplier's instruction. The levels of sAPP α and sAPP β -sw were measured from the conditioned medium using the specific ELISA (IBL) according to the supplier's instruction. The APPsw-transfected HeLa cells with DMSO (vehicle) was used as a negative control.

5.4.6. Statistical analysis

Data was expressed as mean \pm SEM. Statistical comparisons between controls and treated experimental groups were performed using the Student's *t*-test. *P* < 0.05 was considered statistically significant.

5.5. Structure elucidation using NMR

¹H and ¹³C NMR spectra were acquired at room temperature on a Bruker Avarice III 400 spectrometer (¹H at 400.21 MHz and ¹³C at 100.63 MHz) and 500 spectrometer (¹H at 500.10 MHz and ¹³C at 125.75 MHz). Deuterated CD₃OD (Merck, Switzerland) was used to dissolve

the compounds. Chemical shifts were reported in ppm, referenced to residual solvent resonances (methanol-*d*₄ δH 3.31, δC 49.0 ppm). Bruker TopSpin 4.0.6 software was used to process the spectra. The coupling constant “*J*” was given in Hertz (Hz) while the signal multiplicity was reported as: *s* = singlet, *d* = doublet, *dd* = doublet of doublet, *q* = quartet, *m* = multiplet.

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Chapter 6

Conclusion

Alzheimer's disease is an irreversible, slow and progressive brain disorder that destroys memory and other thinking abilities. It is the most common type of dementia. The abnormal cleavage of APP (Amyloid precursor protein), an integral membrane protein produces the accumulated forms of beta amyloid ($A\beta_{42}$), to form the plaques in brain which are the main causes of Alzheimer's disease.

The traditional plants play an important role in the treatment of many diseases like cancer, HIV-AIDS, malaria and Alzheimer's disease. One of the FDA approved drug for the treatment of Alzheimer's disease is galanthamine, an alkaloid (natural product). South Africa has a wide plant diversity. 15% of these plant species are used for medicinal purposes [1] and about 80% of South African people use traditional medicines for their health care [2]. The goal of this study was to identify, characterize and develop new natural ingredients from the South African medicinal plants for the treatment of Alzheimer's disease by targeting the reduction of $A\beta_{42}$ resulting in decreased levels of plaques.

For the selection of the plants in this study, different database searches Google Scholar, Scifinder, Reaxys, Chemspider, Pubchem, Scopus were used for the selection of plant species by using the keywords; African medicinal plants and traditional medicinal plants in combination with the memory loss, mental illness, depression, Alzheimer's disease, dementia, anxiety, epilepsy, forgetfulness, convulsion, hysteria, sedative effects, acetylcholinesterase inhibition which resulted in the selection of 51 plant species. Due to the broad search and wide search terms, a scoring system was developed and used to prioritize, which resulted in 21 plant species being selected for further studies. The different plant parts of the plant species were collected and extracted to give a total of 34 extracts which were screened for $A\beta_{42}$ reduction. Among them, four plant extracts, stems of *H. arborescens*, leaves of *X. undulatum*, leaves of *C. paniculata* and leaves of *S. brachypetala*, potently decreased the levels of $A\beta_{42}$ by $68.6 \pm$

0.2, $76.9 \pm 1\%$, $57.5 \pm 1.3\%$ and $44.8 \pm 0.1\%$, respectively at 50 $\mu\text{g/ml}$. The stem extract of *H. arborescens*, leaf extracts of *X. undulatum* and *C. paniculata* were for the first times screened for A β 42 reduction. In the case for *H. arborescens* and *C. paniculata* which are traditionally used for the treatment of mental disorders, the reduction in A β 42 levels provided some scientific evidence authenticating their traditional uses and further exploration of their potential to be further developed into anti-Alzheimer's ingredients. The ability for the reduction of A β 42 for *S. brachypetala* was demonstrated in other assays in published literature while *X. undulatum* has been previously reported for acetylcholinesterase inhibition [3], [4]. These four extracts were selected for the further study to identify and or isolate the active ingredients responsible for A β 42 reduction.

The three plants, *S. brachypetala* leaf extract, *C. paniculata* leaf extract and *H. arborescens* stem extract were assayed for their A β 42 reduction in dose dependent studies. The plant extracts were further analyzed on UPLC-QTOF-MS to identify the active ingredients which could be responsible for the decreased levels of A β 42. ESI negative mode was used to identify the active ingredients as the ionization was poor for the positive mode for further analysis

S. bracypetala leaf extract showed significant reduction in A β 42 production only at higher concentrations i.e. $29.69 \pm 4\%$ at 50 $\mu\text{g/ml}$ and by $62.17 \pm 7\%$ at 100 $\mu\text{g/ml}$. A total of four compounds belong to the class flavonoids, myricetin-3-O-alpha-L-rhamnopyranoside (**25**), isoquercetin (**26**), quercetin-3-O-rhamnoside (**27**), quercetin (**28**) were identified through UPLC-MS analysis. These compounds were previously identified in the same plant and their potential to treat Alzheimer's disease through A β reduction was published in different bioassays [4]. Based on these findings, the compounds are in all likelihood responsible for the A β 42 reduction in *S. bracypetala*.

The extract of leaves of *C. paniculata* significantly reduced the A β 42 levels in a dose dependent manner by $29.03 \pm 1.4\%$, $42.40 \pm 1.7\%$, $47.50 \pm 5.8\%$ and $61.14 \pm 9.3\%$ at concentrations 10, 25, 50 and 100 $\mu\text{g/ml}$, respectively. This is the first report for the reduction of A β 42 and thus provides scientific evidence which substantiates its traditional use related to

treatment of nervous disorder treatments. The UPLC-QTOF-MS analysis led to the identification of nine compounds; quinic acid (**29**), 3,5-dicaffeoylquinic acid (**30**), rutin (**31**), valeriananoid E (**32**), acuminoside (**33**), dictamnocide D (**34**), 2''-O- β -D-glucopyranosylsaikosaponin B2 (**35**), clinoposaponin C (**36**), spinasaponin C (**37**). The presence of rutin was confirmed by using pure standard for the UPLC-QTOF-MS analysis. Rutin is a well-researched compound for the Alzheimer's disease in previous studies [5], [6]. 3,5-dicaffeoylquinic acid was previously examined for AChE inhibition [7]. Based on these studies rutin and/or 3,5-dicaffeoylquinic acid could be responsible for the activity in *C. paniculata*.

The stem extract of *H. arborescens* significantly reduced A β 42 levels only at high test concentrations by $75.78 \pm 0.1\%$ at 50 $\mu\text{g/ml}$ and by $31.37 \pm 0.2\%$ at 25 $\mu\text{g/ml}$. A total of four compounds were tentatively identified as cynarin (1,5-dicaffeoylquinic acid) (**38**) and three triterpenoid saponin alternoside IX (**39**), alternoside I (**40**) and saikogenin B4 (**41**) using the UPLC-QTOF-MS. Cynarin was previously reported for the reduction of A β cytotoxicity while the other identified compounds, triterpenoid saponins were not examined for Alzheimer's disease. In this study cynarin could be the active ingredient from *H. arborescens* leaf extract for the reduction of A β 42.

X. undulatum leaf extract was highly potent against A β 42 reduction in the present study based on the data, it was further explored for its potential to treat Alzheimer's disease through the reduction of A β 42. The plant is commercialised in Europe for the treatment of diarrhoea and traded as the commercial brand Uzara. The plant is also used traditionally for treatment of many diseases such as dysentery, diarrhoea, headache, malaria, typhoid, skin diseases, infected sores and wounds. Although the plant has been widely researched and well-known for its antidepressant effect, acetylcholinesterase inhibitory effects and selective serotonin reuptake inhibitor (SSRI) activity, there is limited published data available showing its inhibitory effects for the reduction of A β 42, an important target in the treatment of Alzheimer's disease. In the present study, the DCM:MeOH leaf extract significantly reduced the A β 42 production

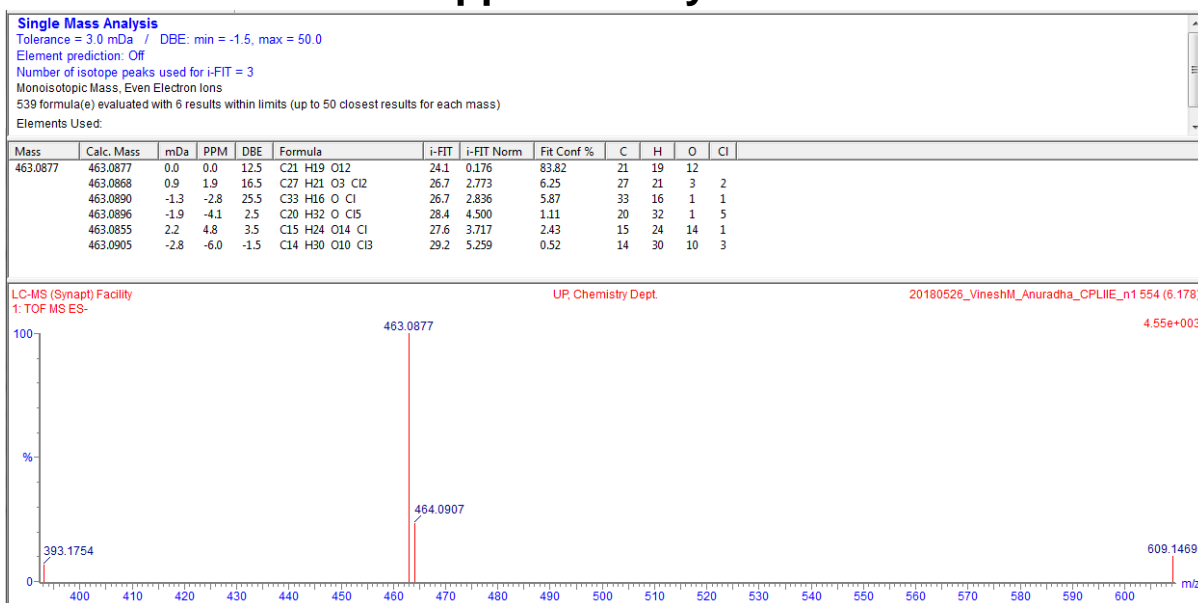
by $49.8 \pm 4.0\%$, $71.4 \pm 1.6\%$ and $72.9 \pm 1.1\%$ at concentrations of 1, 5 and 10 $\mu\text{g/ml}$, respectively. To the best of our knowledge, this is the first study examining the plant extracts for its A β 42 production as a potential for treating Alzheimer's diseases. The main class of compounds in the extract was identified as cardenolide glycosides through the analysis of the samples UPLC-QTOF-MS confirming other previous findings [8]. In order to identify the active ingredients between the several cardenolides present in the crude extract, bioassay-guided fraction was undertaken. The crude DCM:MeOH extract was fractionated using silica gel column to produce a total of 46 fractions which were combined based on their similar TLC profiles to give a final 15 combined fractions. These fractions were then tested for A β 42 levels and only two fractions (AT-1-49N and AT-1-49O) were found to significantly exhibit A β 42 reduction. These fractions were once again analysed on UPLC-QTOF-MS to identify the compounds and the results revealed the presence of cardenolide glycosides in the two active fractions. These fractions were further purified using mass directed purification using preparative HPLC which resulted in the isolation of five pure compounds. The structure elucidation of the compounds was undertaken using mass spectral and NMR analysis which resulted in the identification of acetylated glycosydated crotoxogenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), crotoxigenin-3-O-glucopyranoside (**59**), desglucouzarin (**60**). These compounds were tested for A β 42 reduction and four cardenolide glycosides, acetylated glycosydated crotoxogenin (**56**), xysmalogenin-3, β -D-glucopyranoside (**57**), crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**), desglucouzarin (**60**) were found to significantly decreased the A β 42 levels at 10 μM by $75.04 \pm 0.85\%$, $74.66 \pm 0.1\%$, $65.55 \pm 0.19\%$ and $69.15 \pm 0.25\%$, respectively as compared to the negative control. Although, the compound crotoxigenin-3-O-glucopyranoside (**59**) was found to be inactive, structurally the characteristic lactone ring was present. This pointed to the probability that the lactone is not responsible for the activity towards the A β 42 reduction. As it is known that the lactone ring is the main moiety responsible for the cardiotoxic effects [9], any analogues synthetically prepared lacking the lactone ring may still retain A β 42 reduction while also decreasing

cardiotoxicity. Further dose response studies of the 4 cardenolides identified crotoxinigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**) as the most potent compound as it significantly reduced A β 42 levels at the low concentration of 0.025 μ M, indicating that this is the most active compound responsible for A β 42 reduction in the plant and exhibits the potency to develop into anti-Alzheimer's drug. The bioassay-guided approach of identifying the active ingredient in *X. undulatum* responsible for the reduction of A β 42 level was highly successful and resulted in identification of 4 cardenolides active active ingredient, with one being the most active at nanomolar levels, which compares favourably to conventional pharmaceutical drugs. Often the approach of bioassay-guided fraction is a challenge to natural product chemists as bioactivity is lost during fractionation for various reasons, however in this case potency of the fractions and the final compounds increased during the purification process. The future work will involve the prediction of binding sites in A β 42 protein by docking studies and the removal of the butenolide ring and testing the glycosidated aglycones for their potential to reduce A β 42.

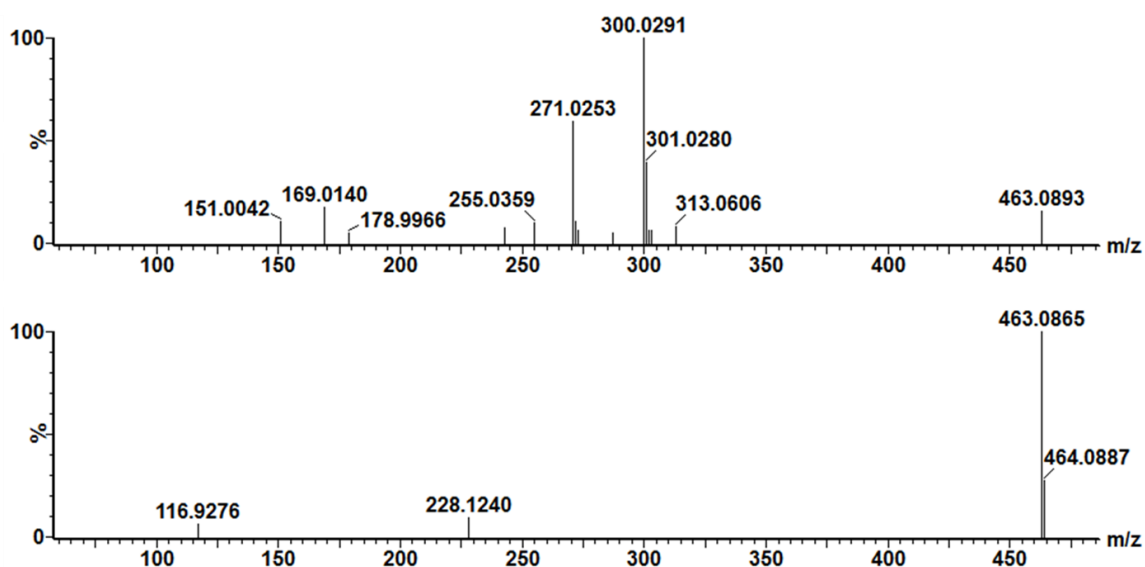
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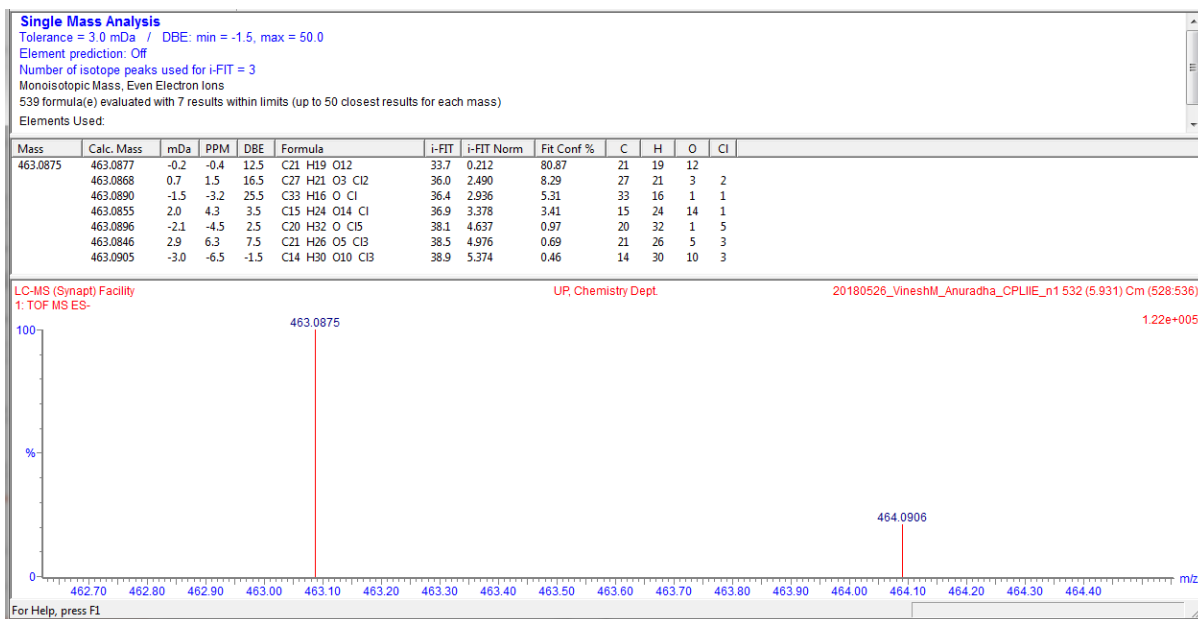
Supplementary Data



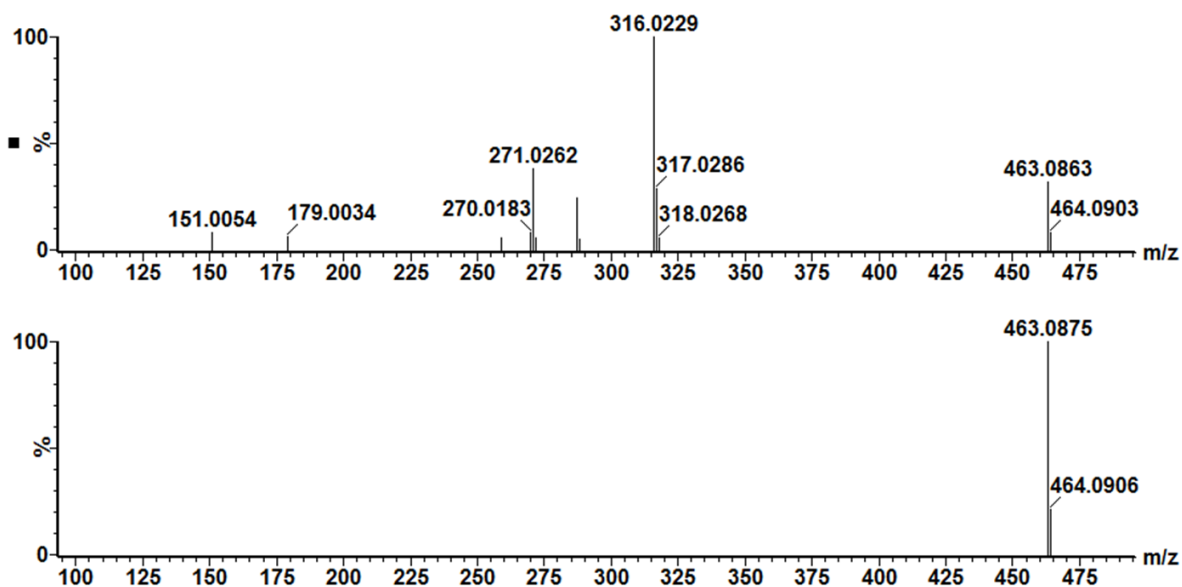
Supplementary data 1: iFit value of isoquercetin (25) in *S.brachypetala* leaves extracted with DCM:MeOH



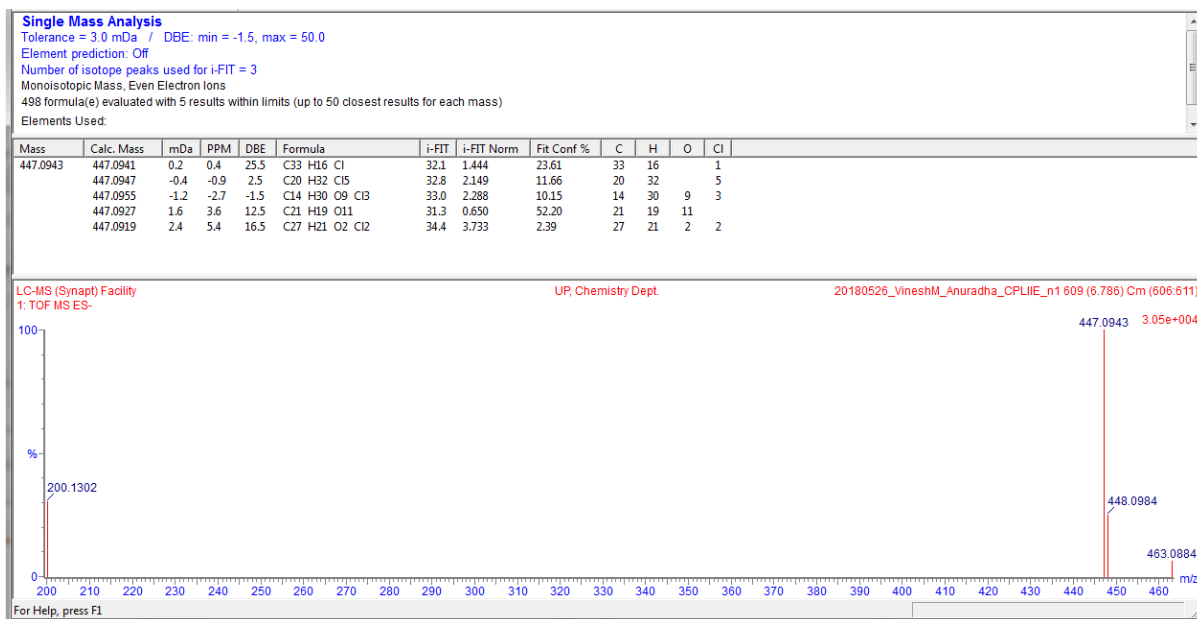
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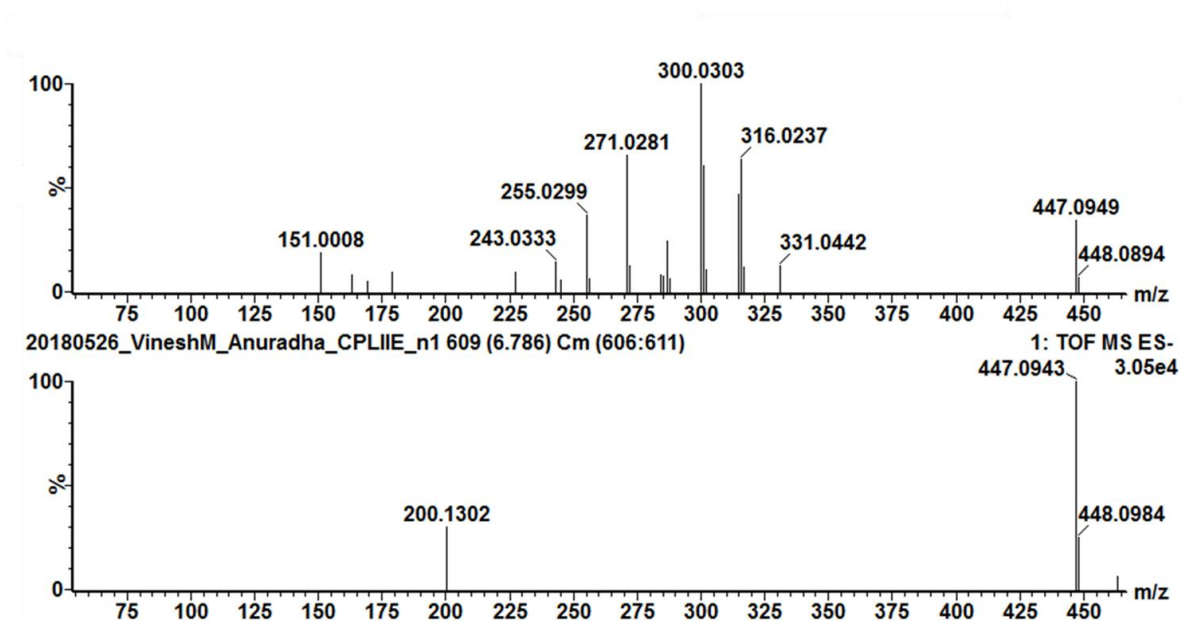
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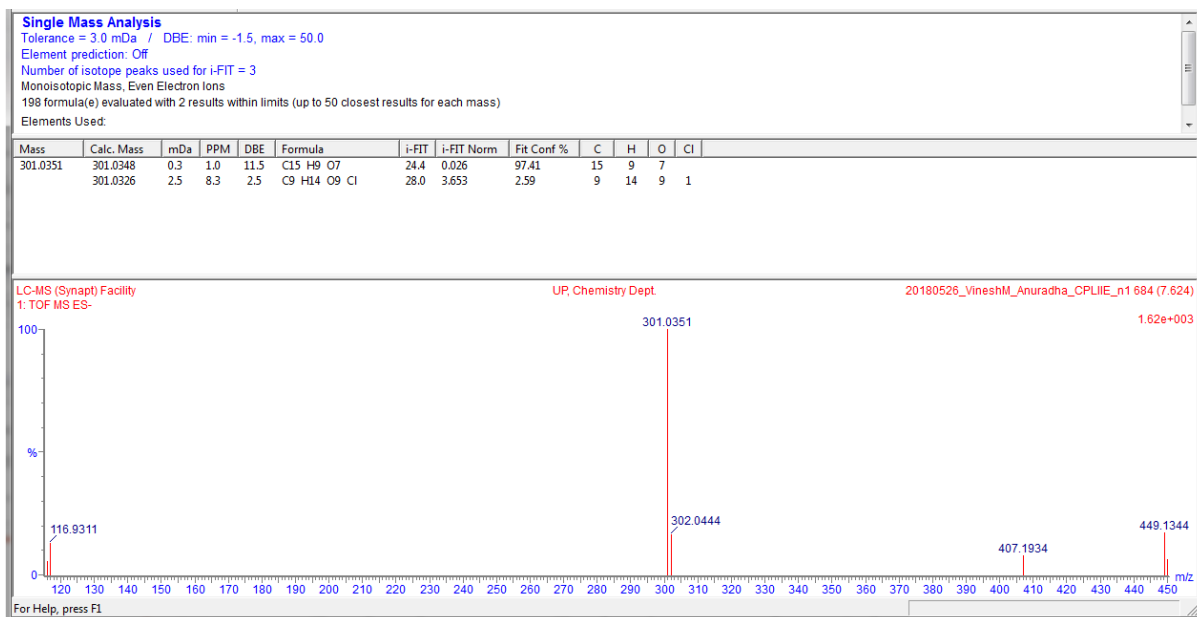
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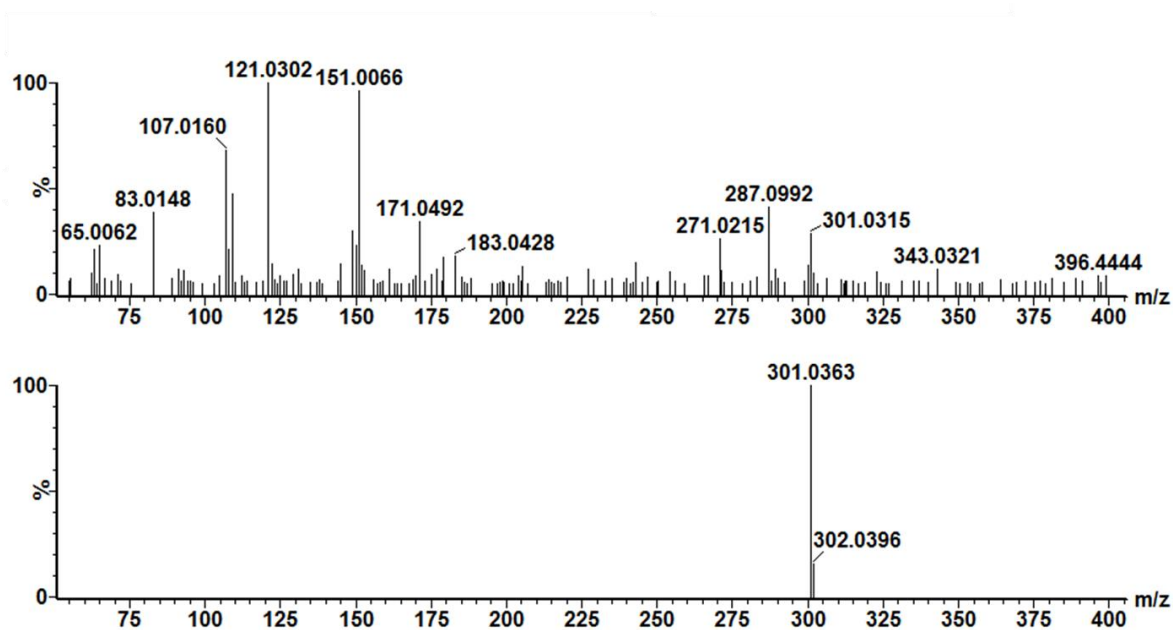
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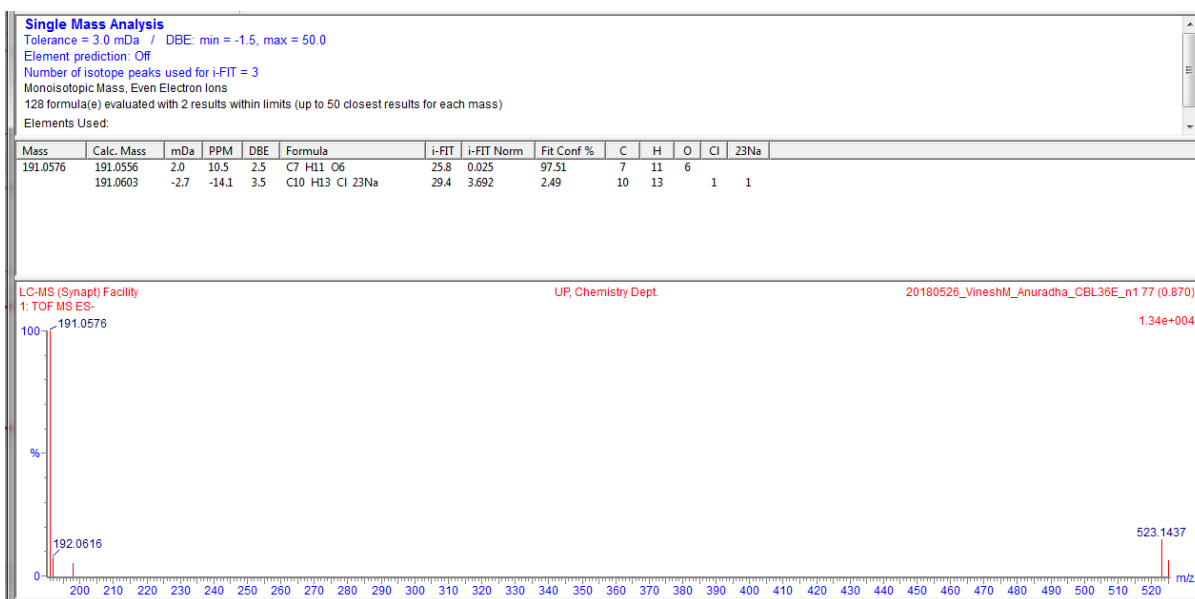
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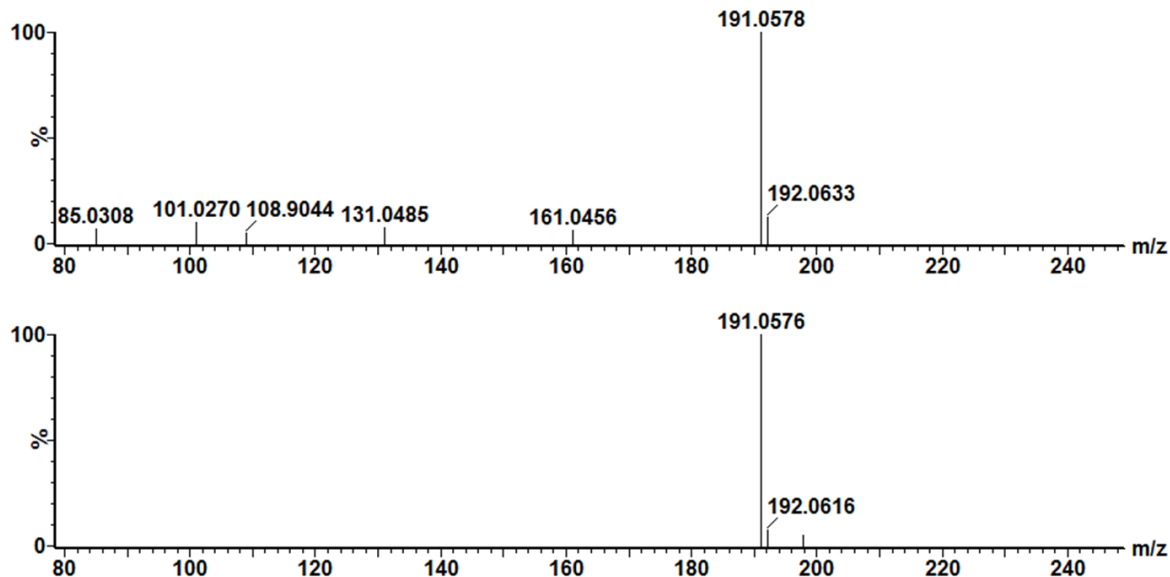
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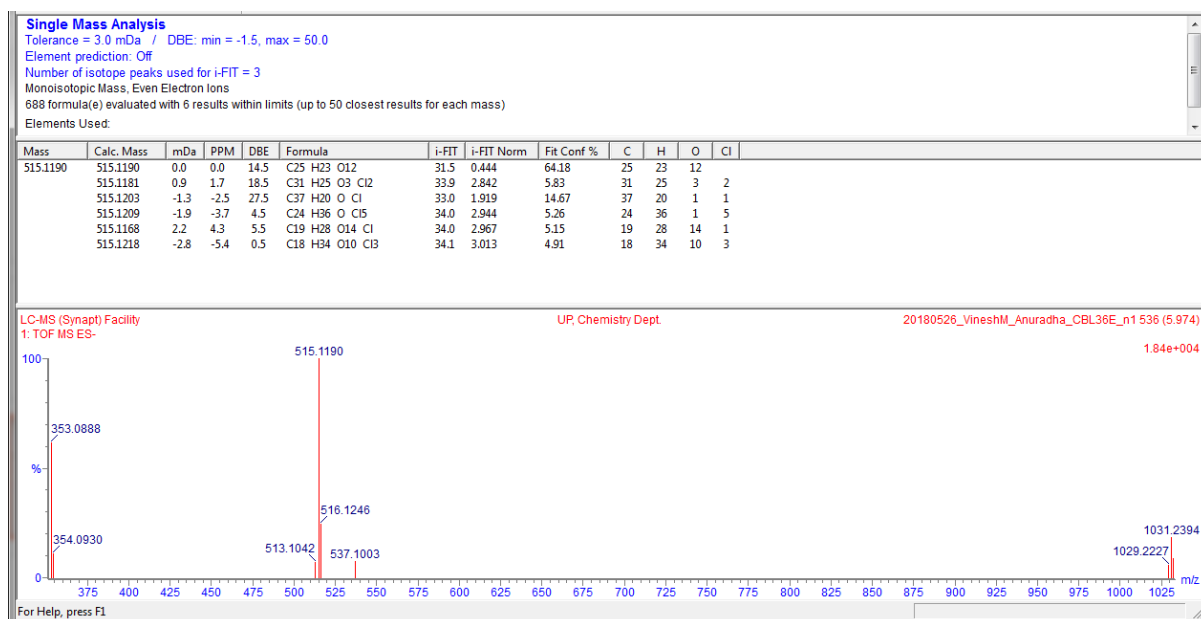
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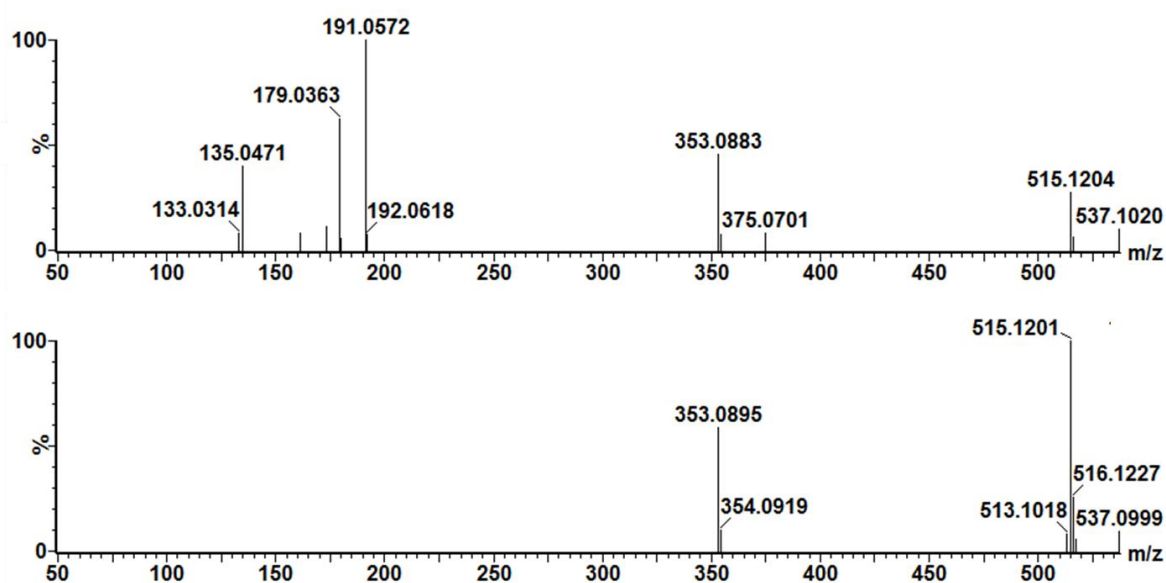
Supplementary data 9: iFit value of quinic acid (**29**) in *C. paniculata* leaves extracted with DCM:MeOH



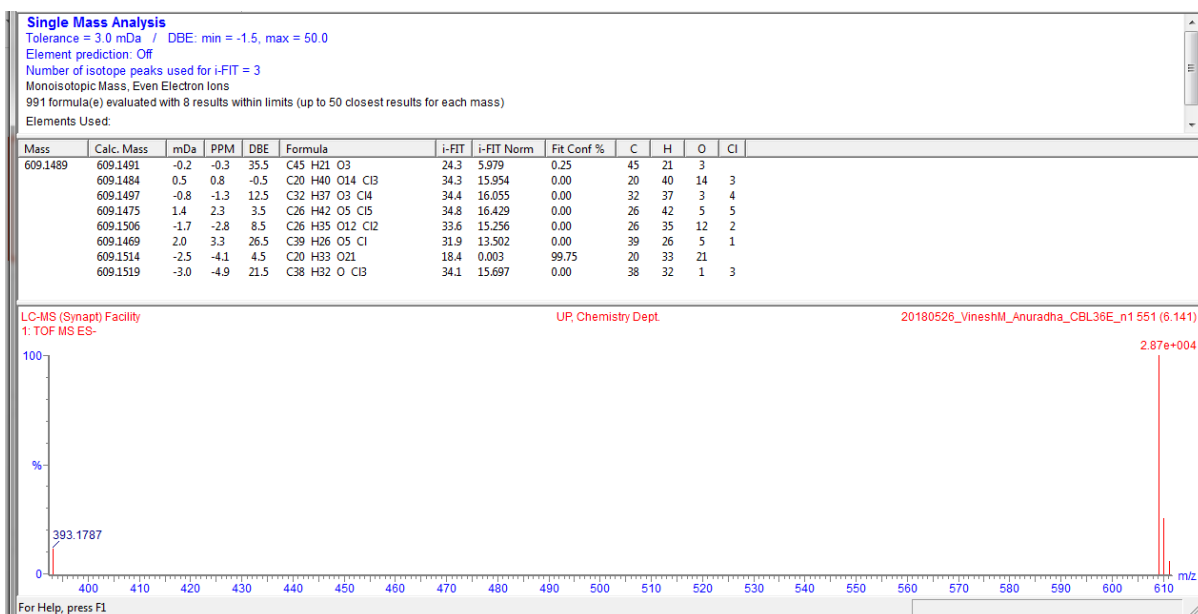
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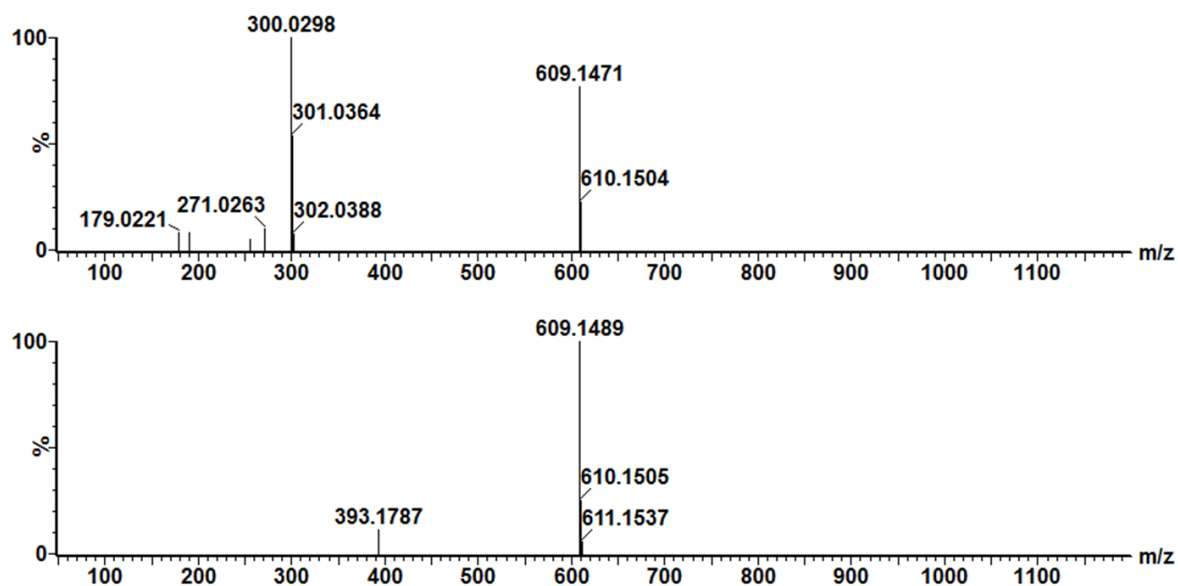
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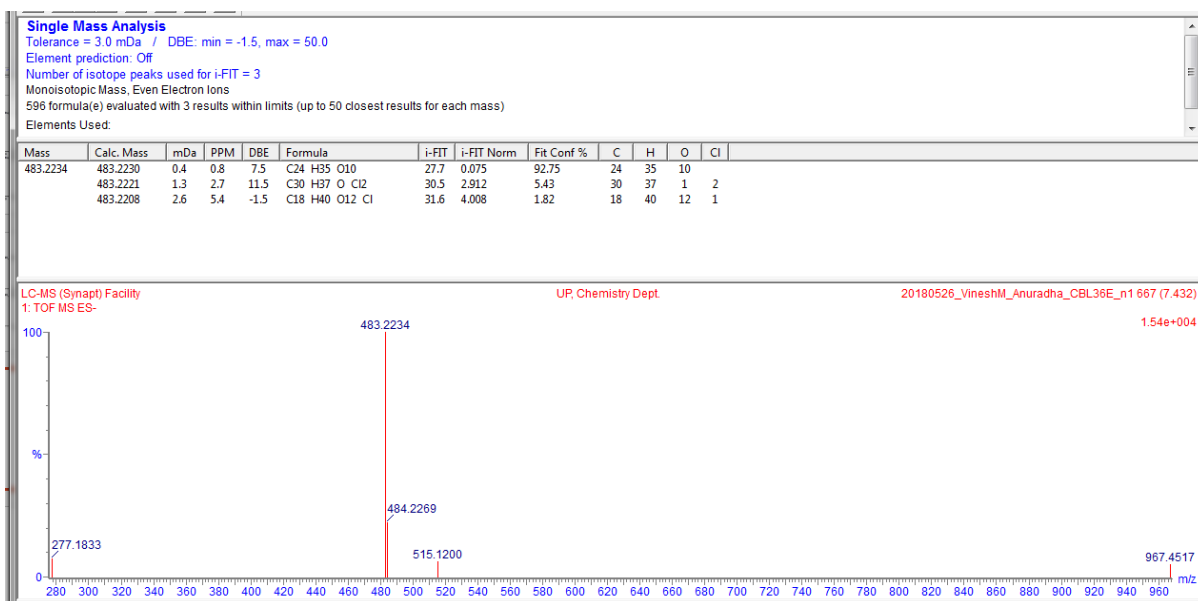
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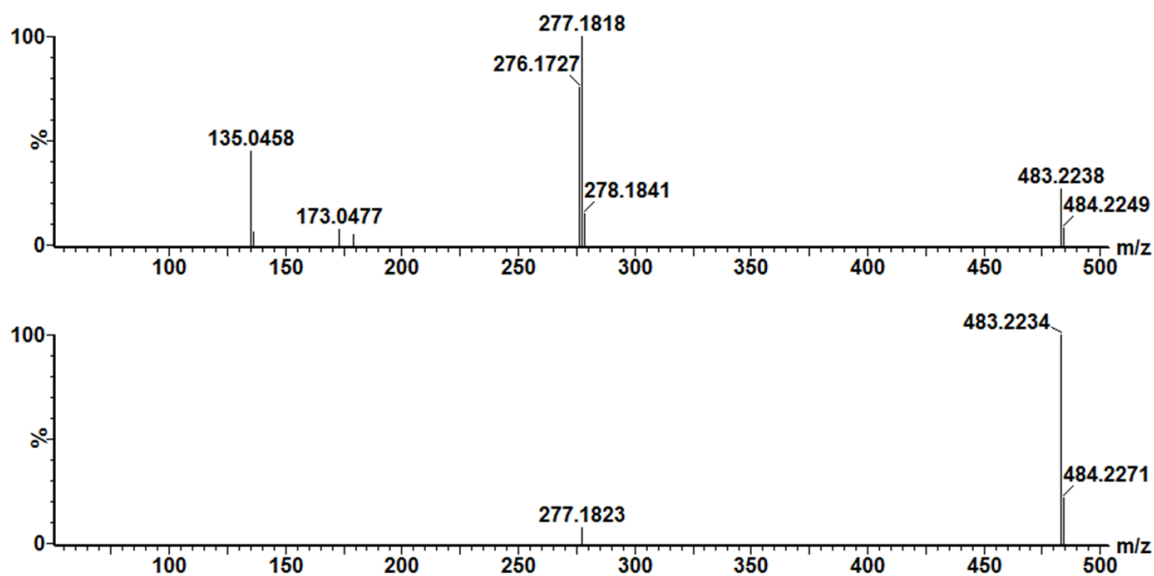
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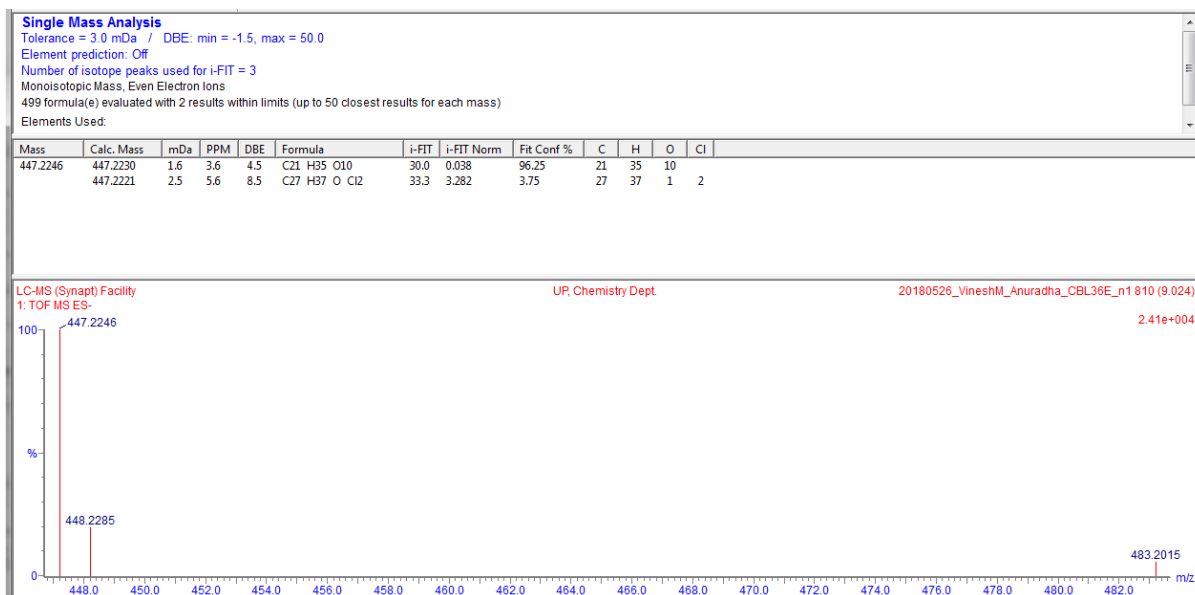
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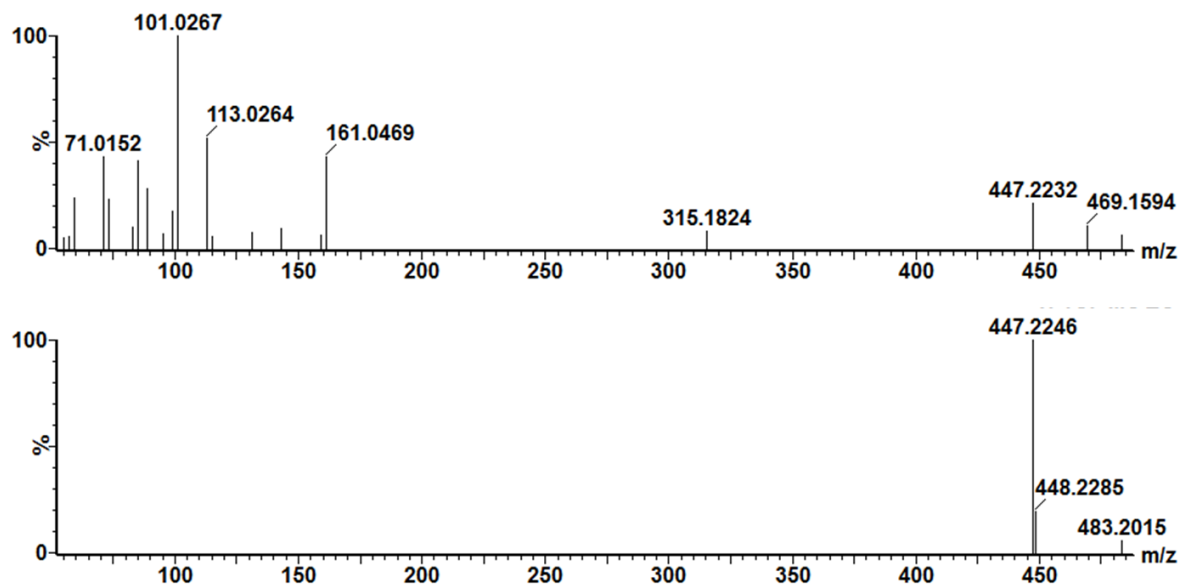
Supplementary data 15: iFit value of valeriananoid E (32) in *C. paniculata* leaves extracted with DCM:MeOH



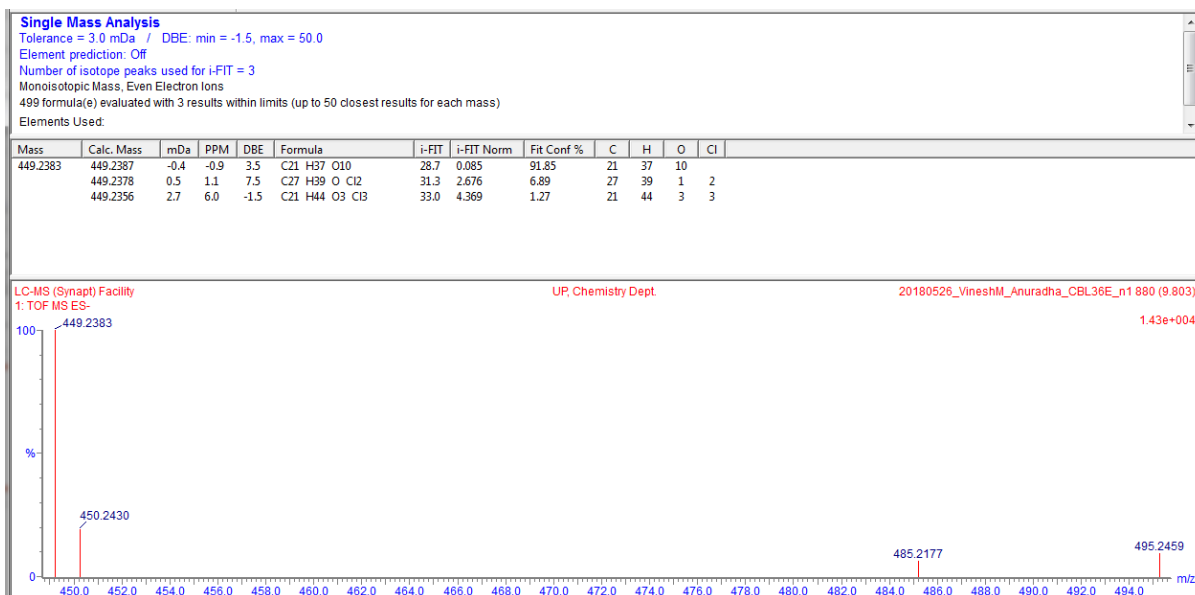
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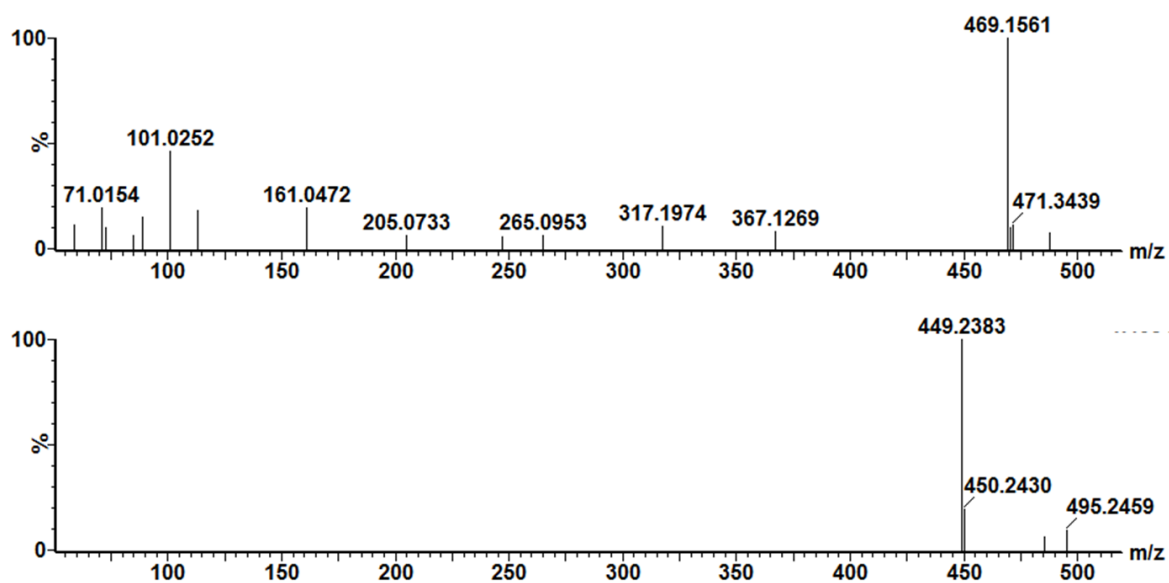
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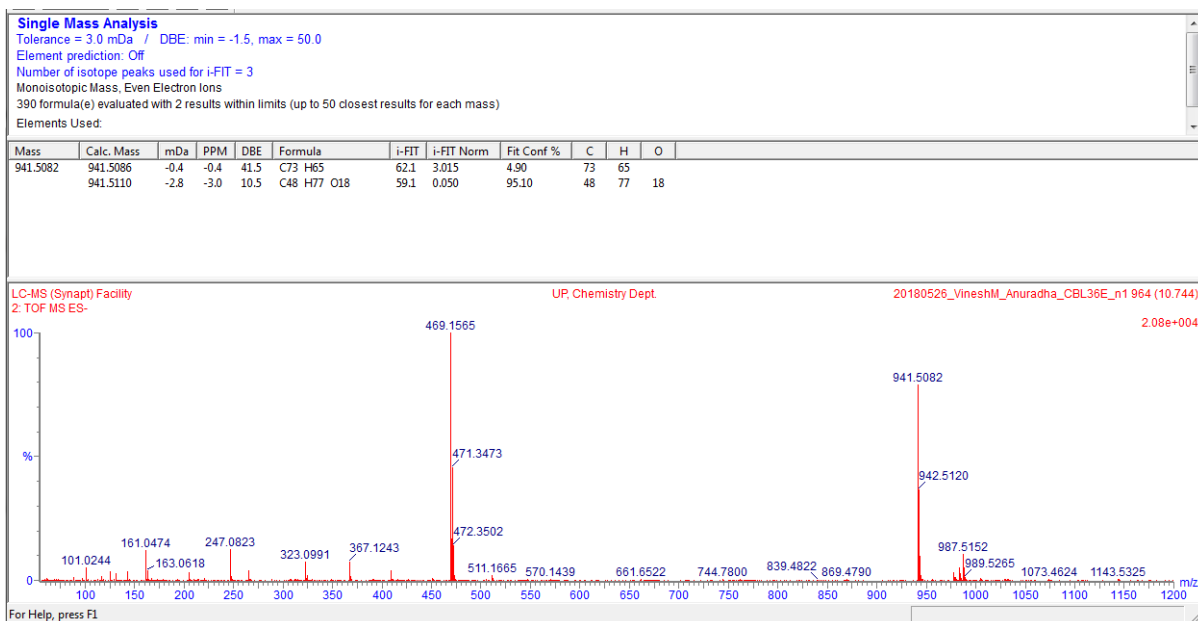
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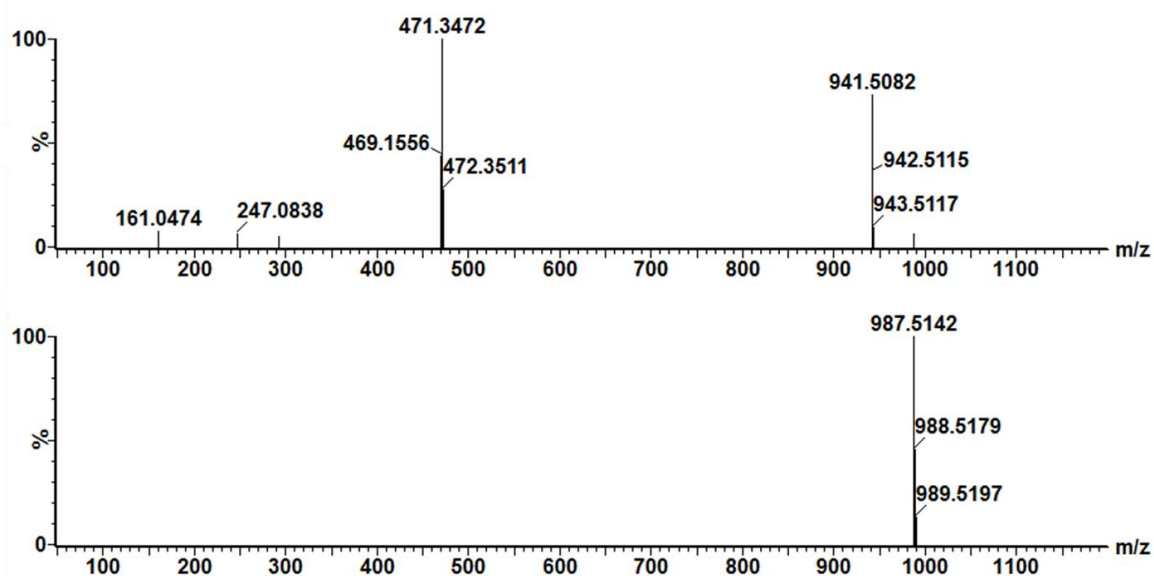
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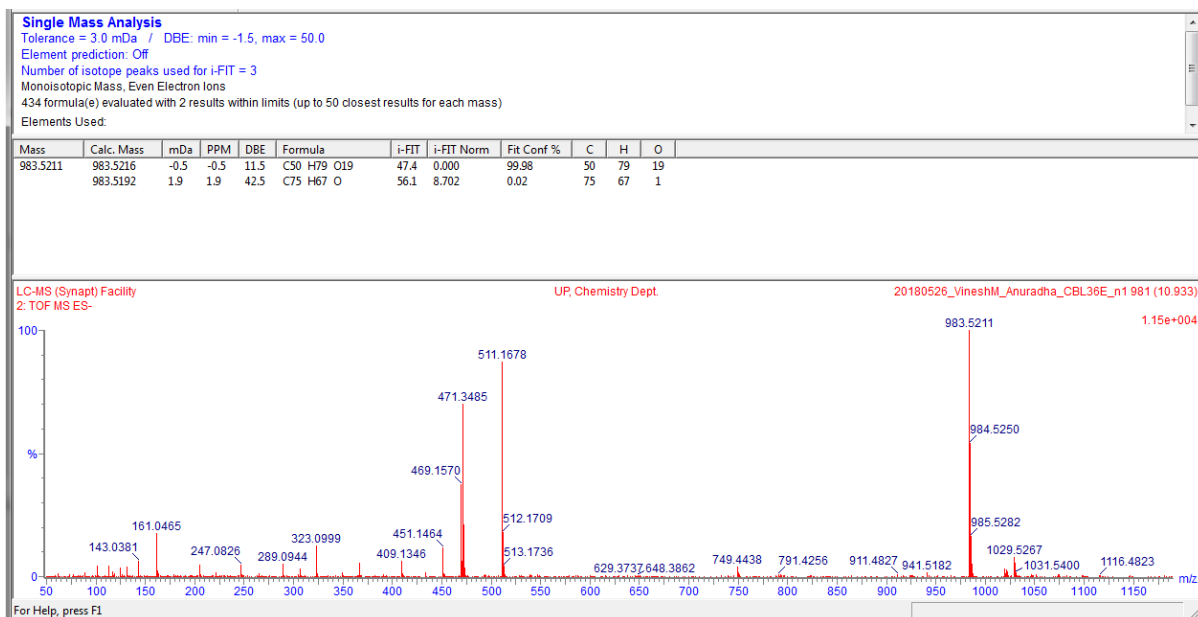
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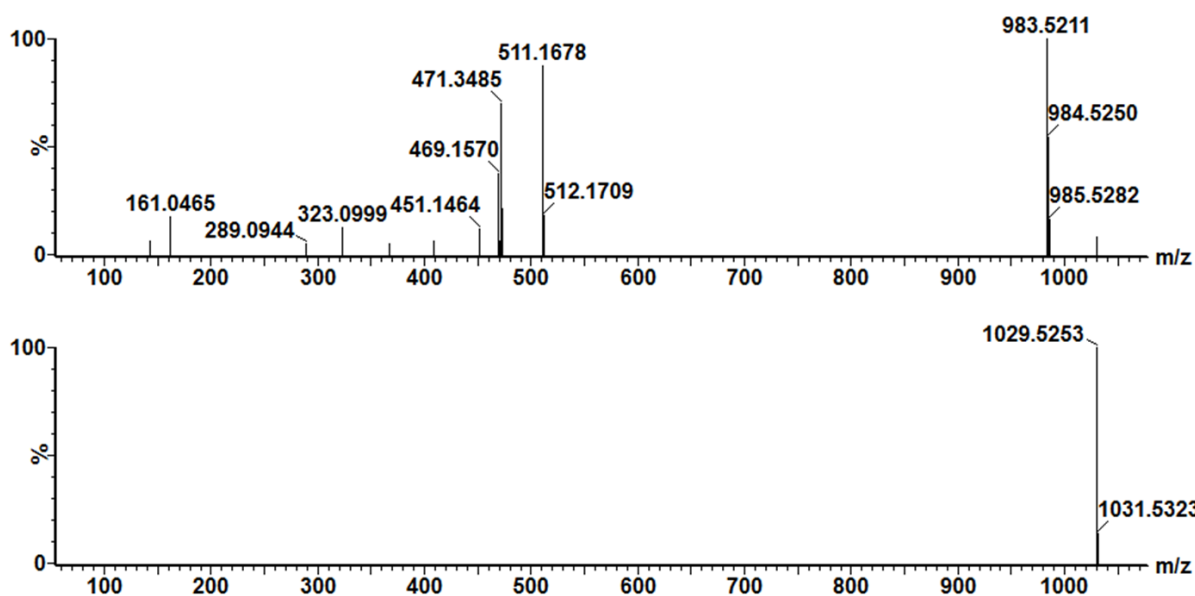
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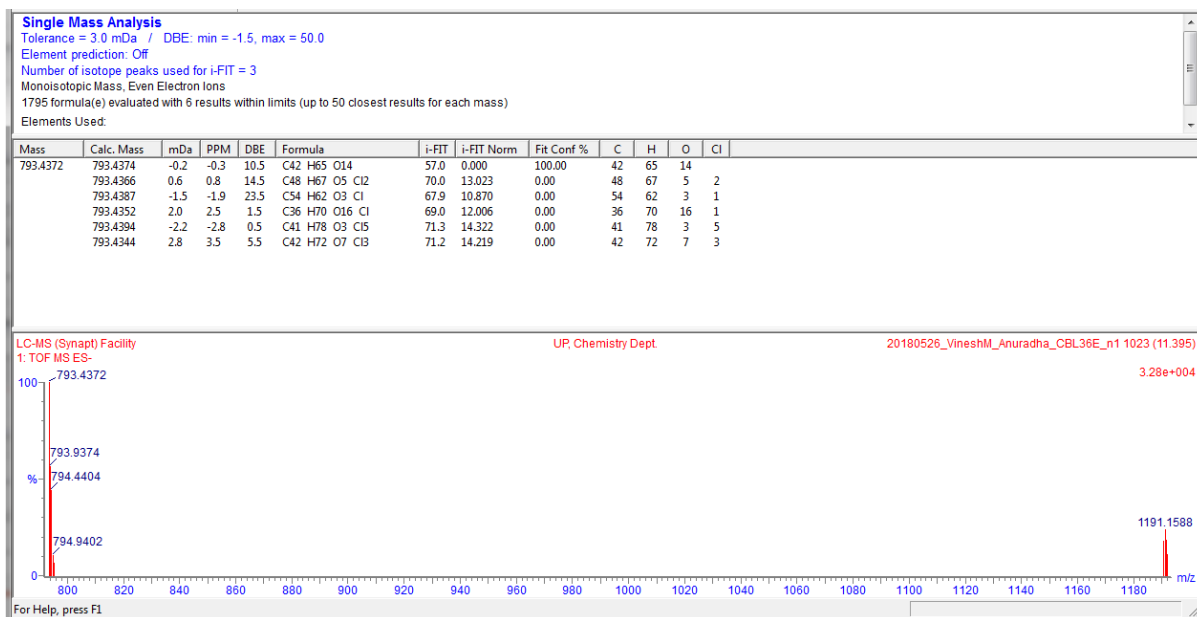
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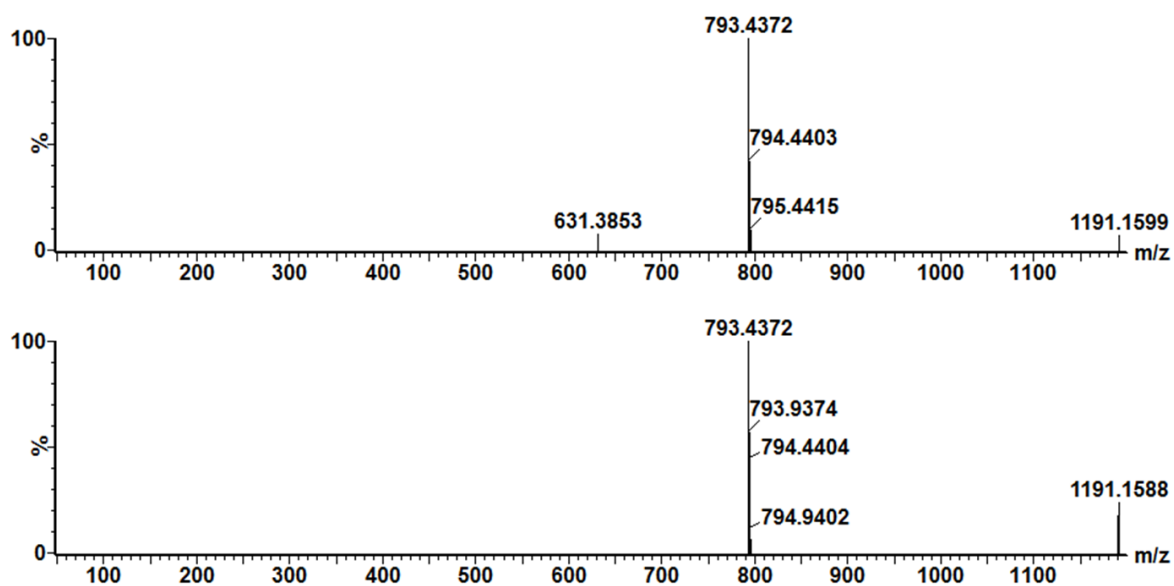
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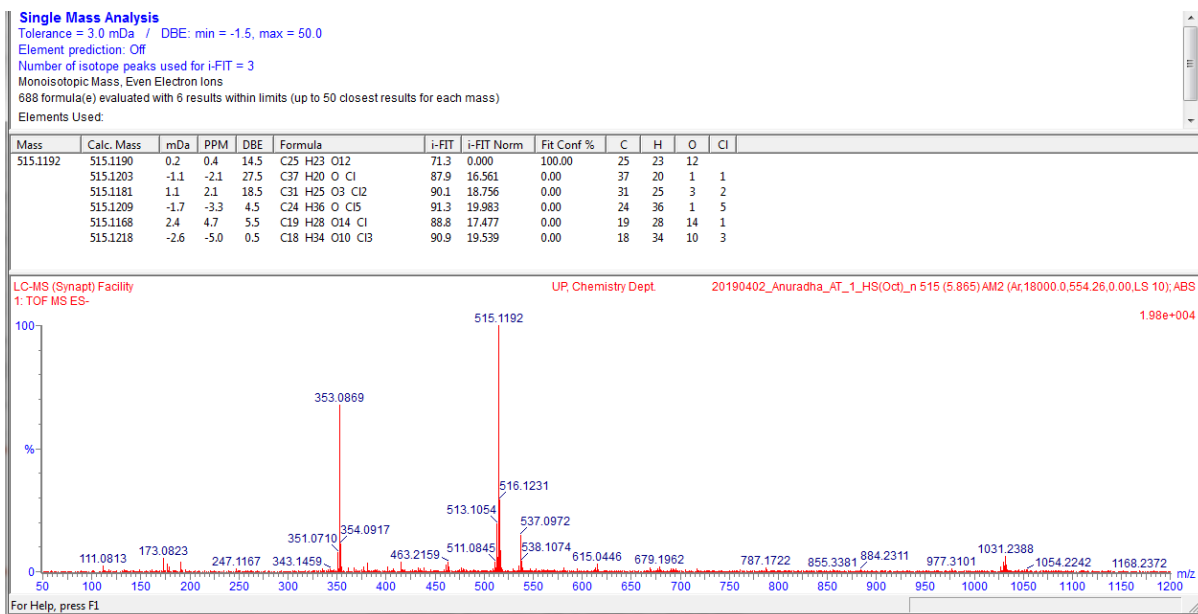
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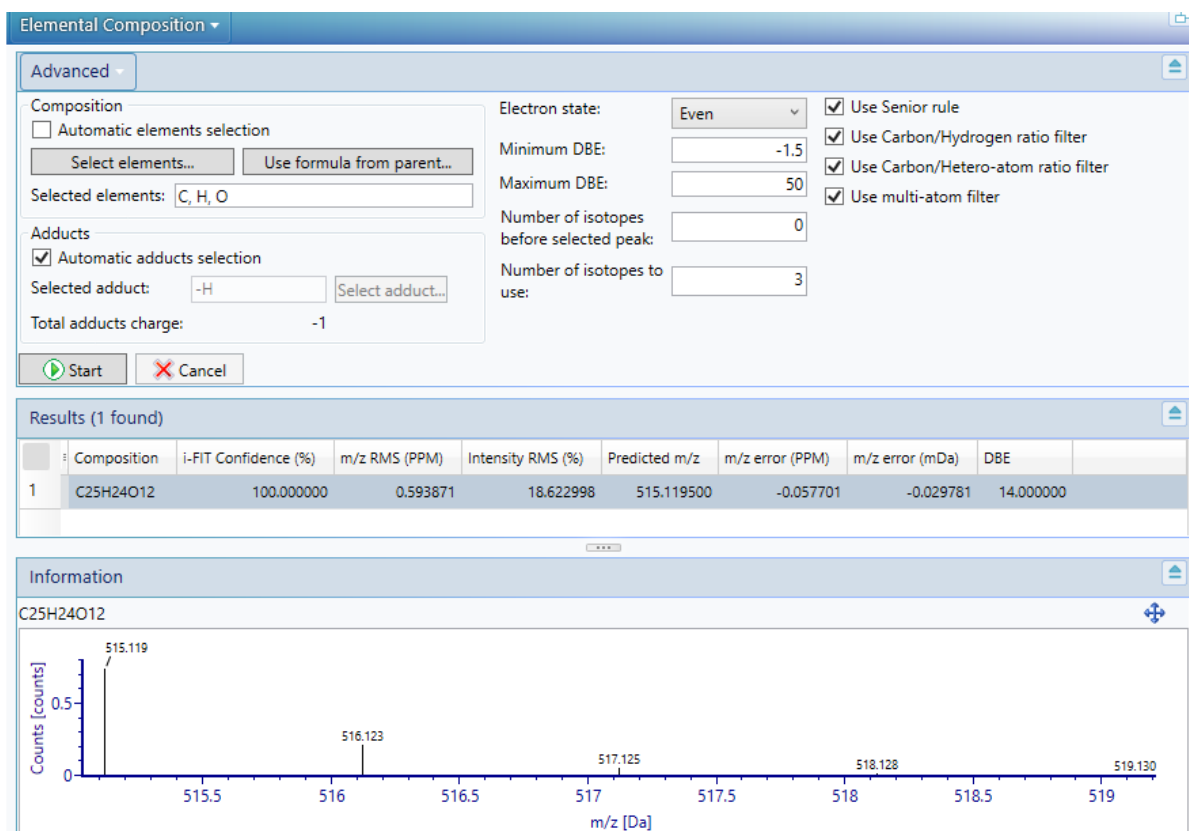
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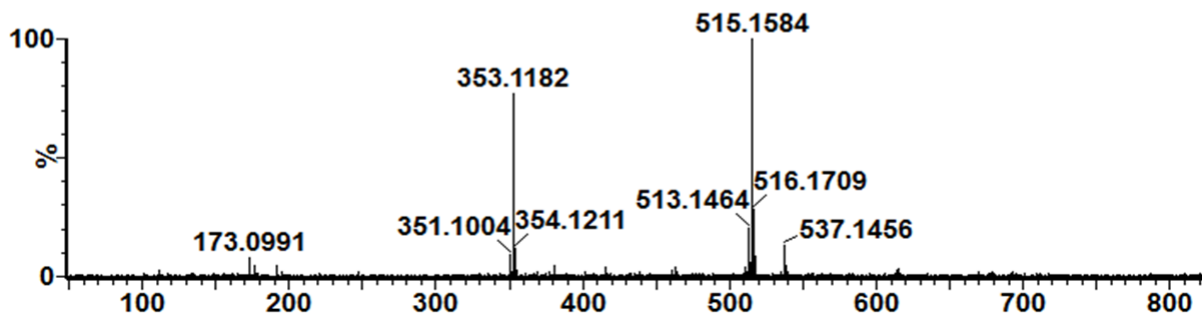
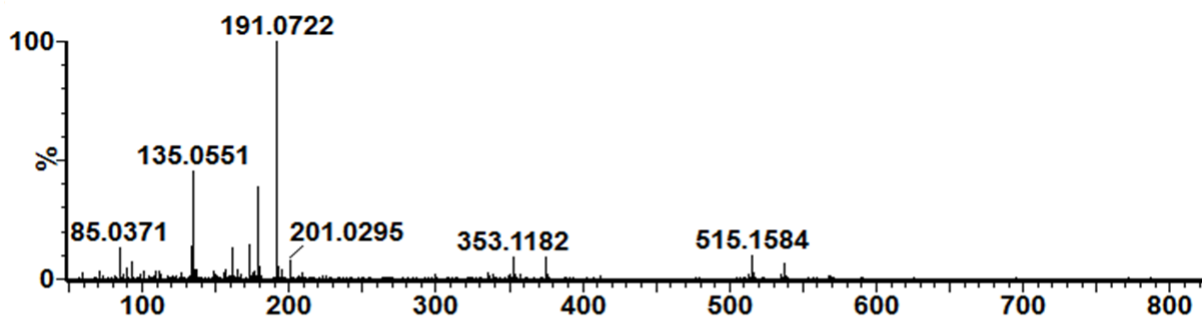
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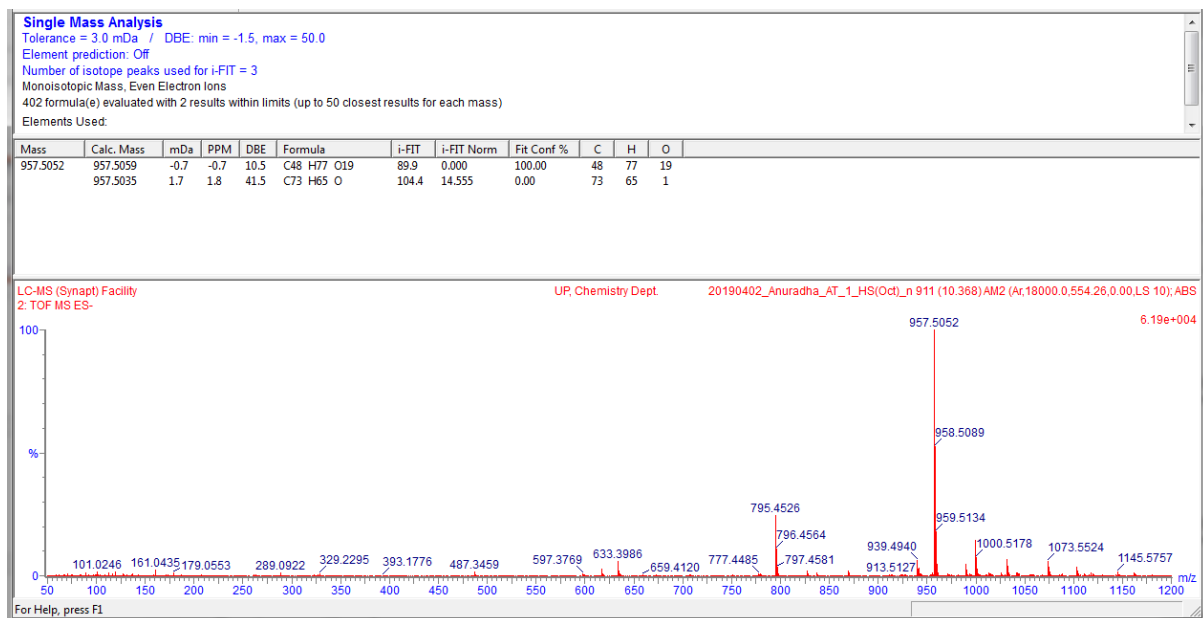
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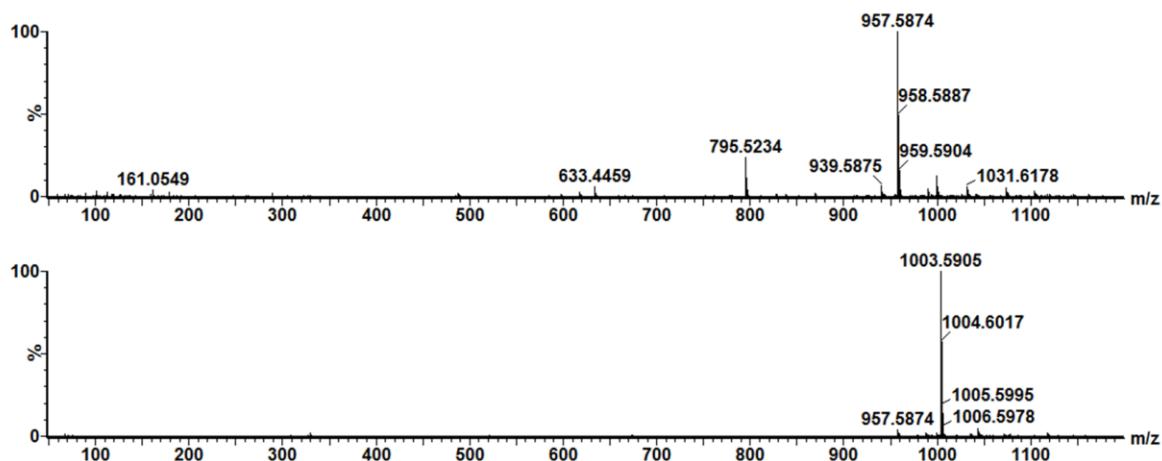
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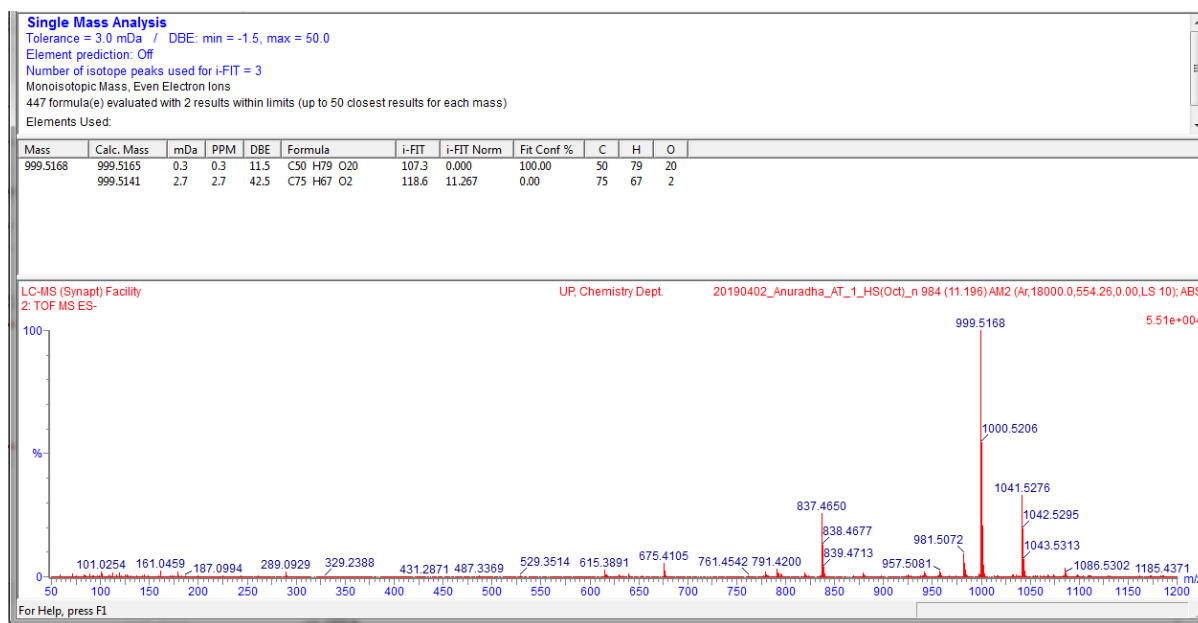
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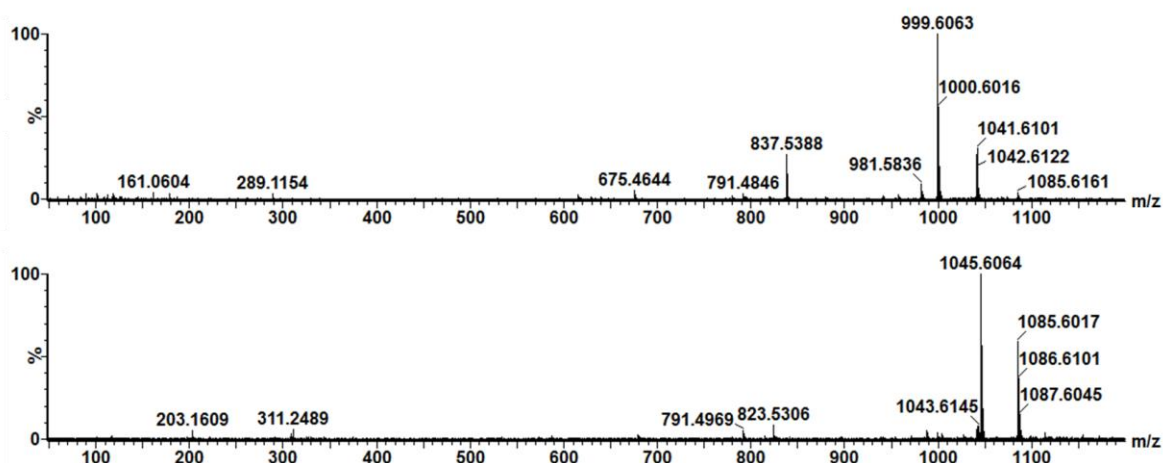
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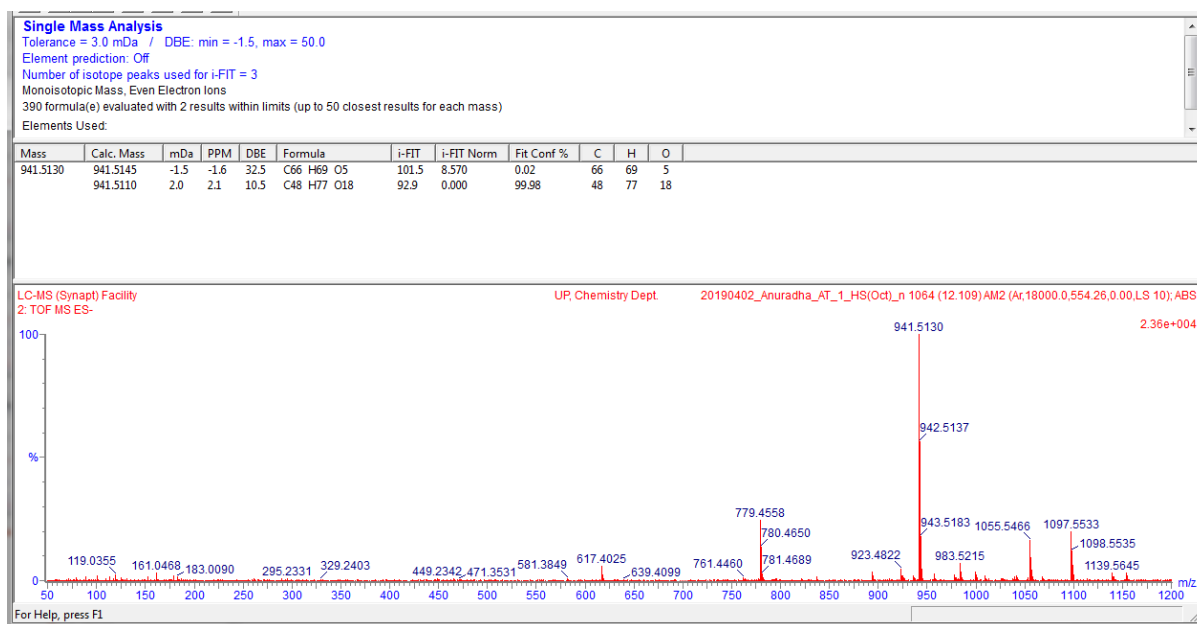
Supplementary data 31: MS and MS/MS data of alternoside IX (**39**) in the DCM:MeOH extract of leaves of *H. arborescens*



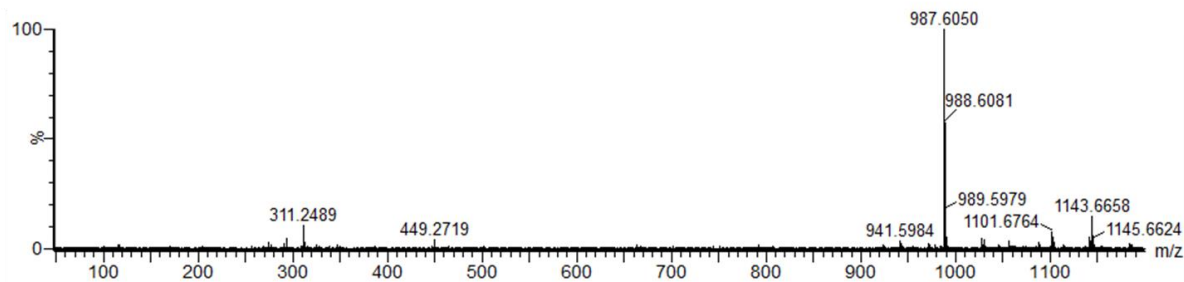
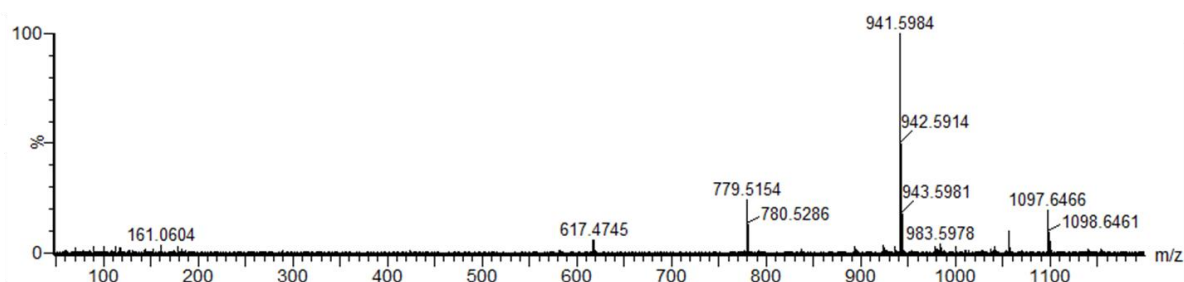
Supplementary data 32: iFit value of alternoside I (**40**) in *H. arborescens* stems extracted with DCM:MeOH



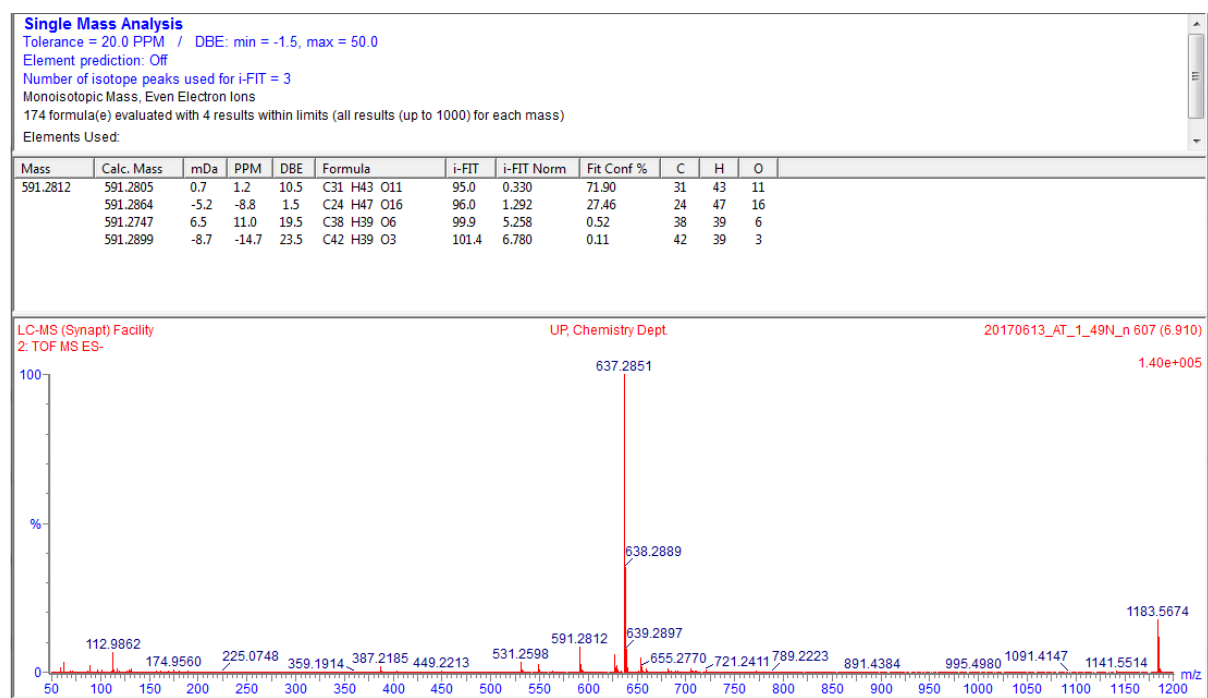
Supplementary data 33: MS and MS/MS data of alternoside I (**40**) in the DCM:MeOH extract of leaves of *H. arborescens*



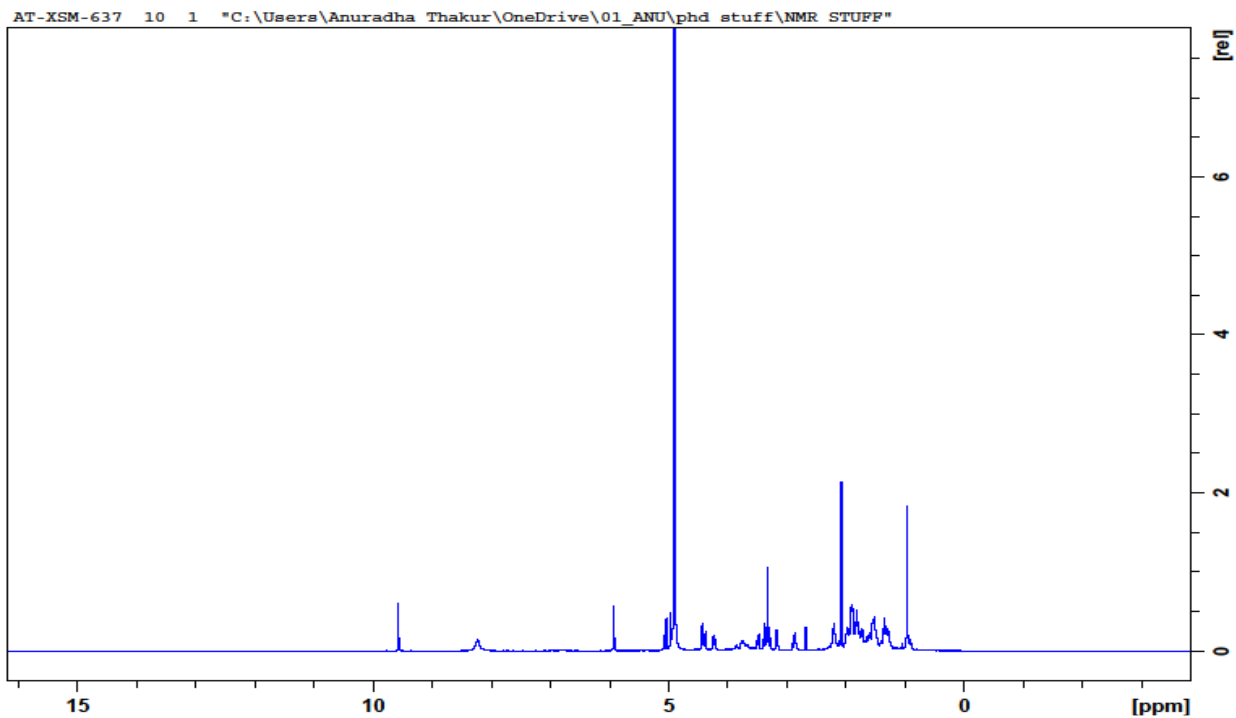
Supplementary data 34: iFit value of saikogenin B₄ (**41**) in *H. arborescens* stems extracted with DCM:MeOH



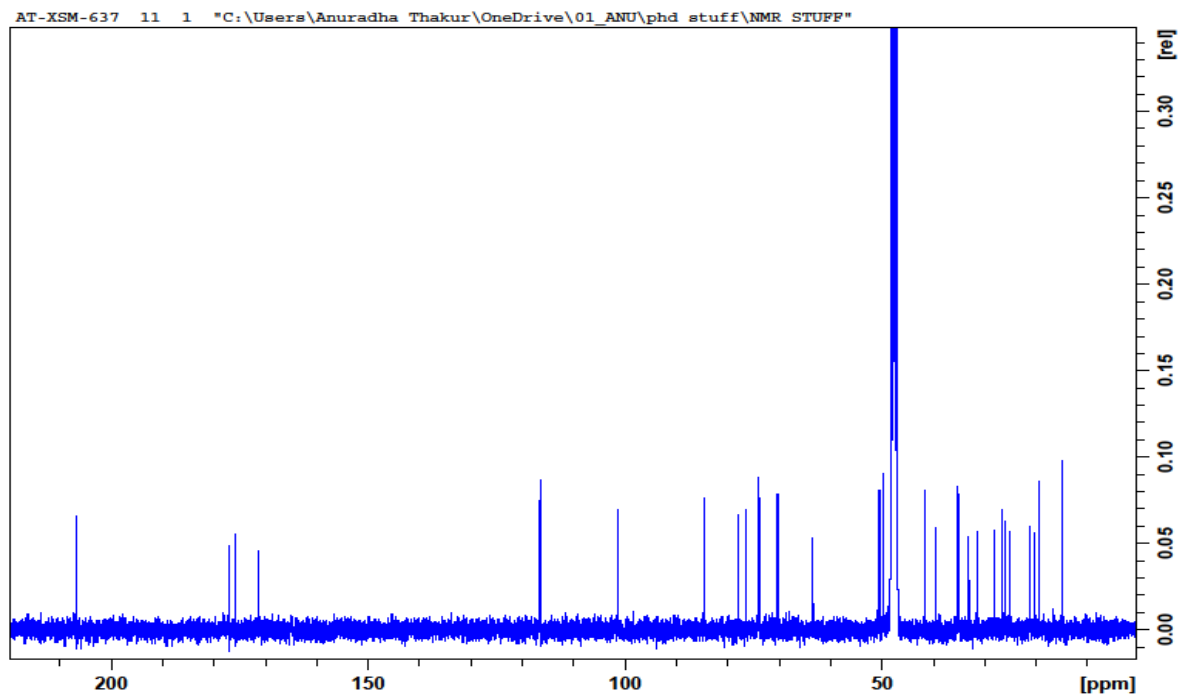
Supplementary data 35: MS and MS/MS data of saikogenin B₄ (41) in the DCM:MeOH extract of leaves of *H. arborescens*



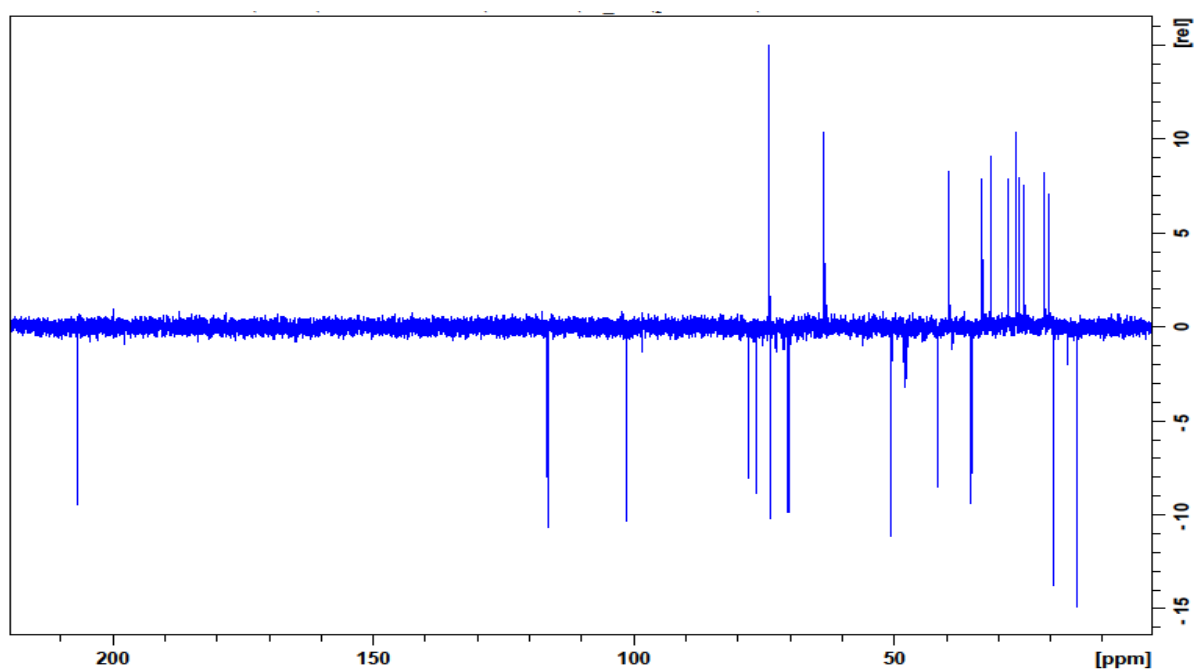
Supplementary data 36: iFit value of acetylated glycosydated crotoxinigenin (56) in *X. undulatum* leaves extracted with DCM:MeOH



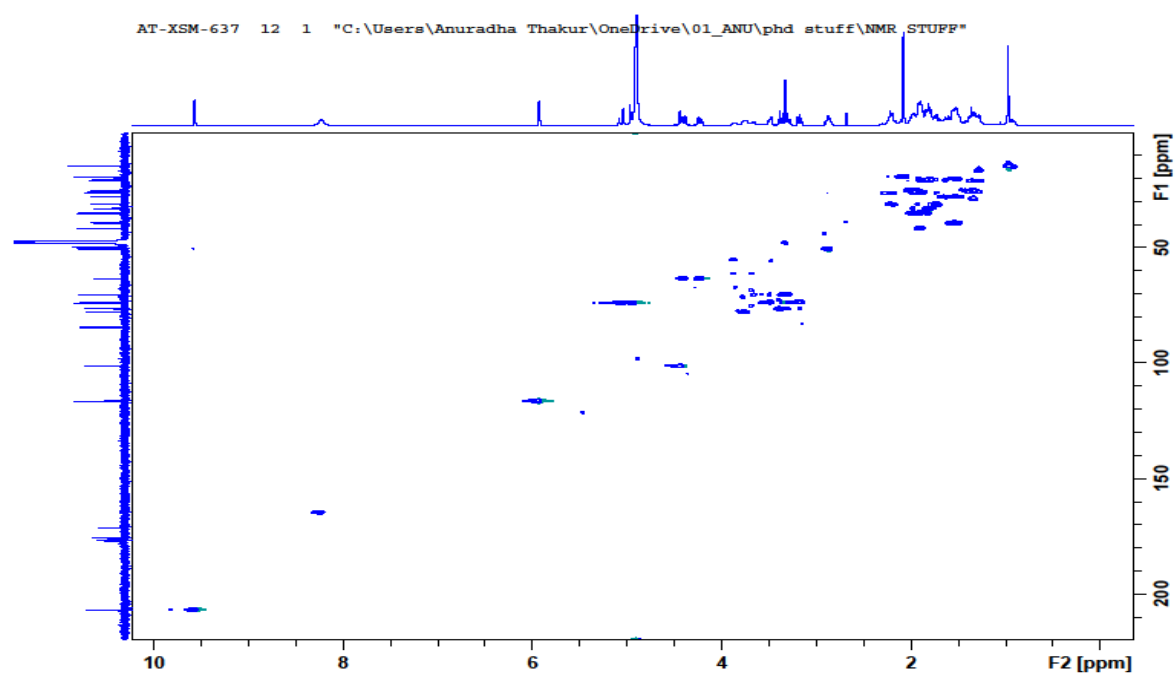
Supplementary data 37: ^1H NMR for compound 3 acetylated glycosydated crotoxigenin (**56**) in MeOD- d_4



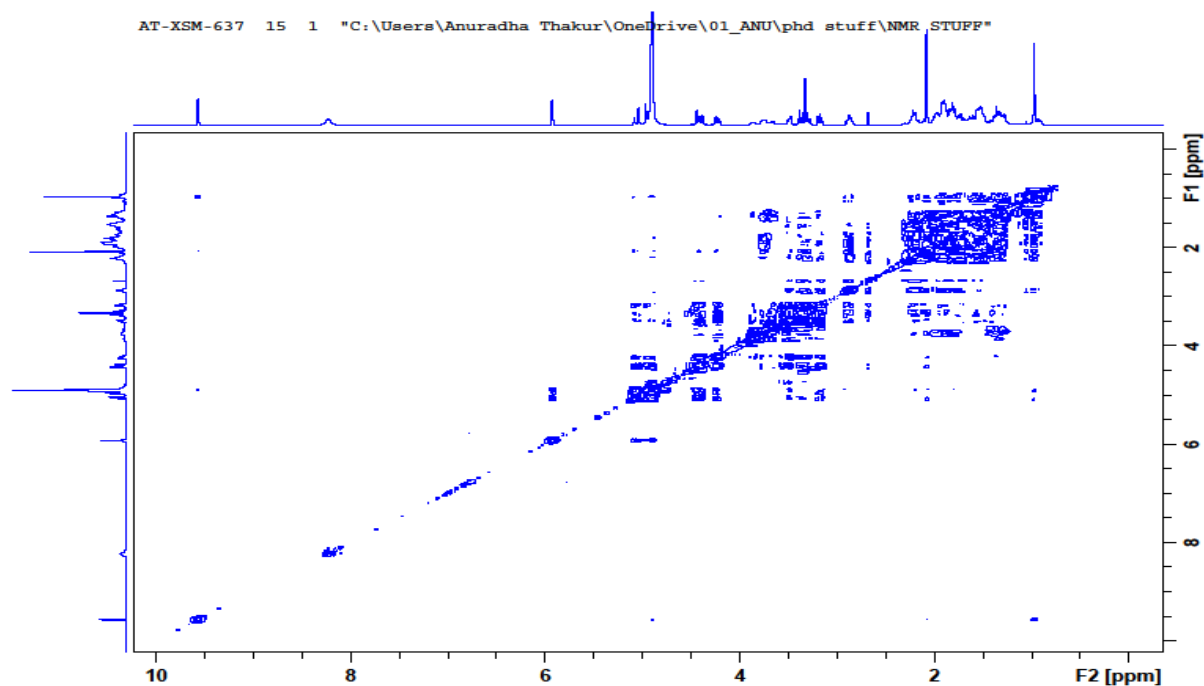
Supplementary data 38: ^{13}C NMR for compound 3 acetylated glycosydated crotoxigenin (**56**) in MeOD- d_4



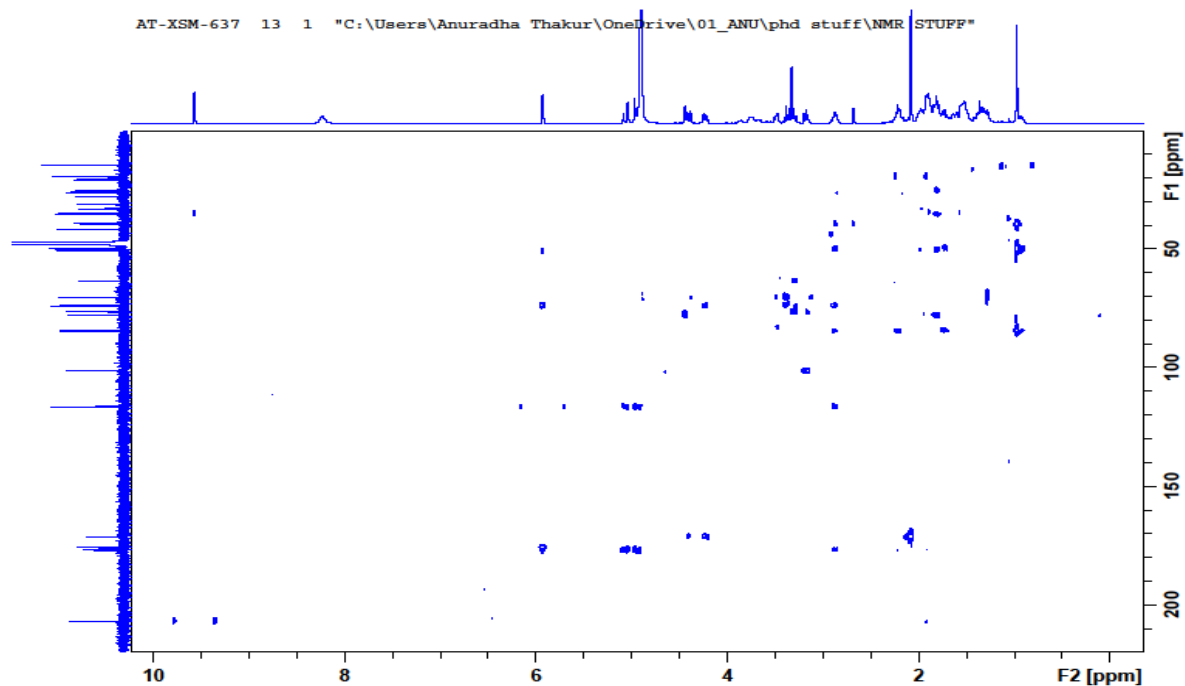
Supplementary data 39:DEPT 135 for acetylated glycosydated crotoxigenin (**56**) in MeOD-*d*₄



Supplementary data 40:HSQC spectrum for acetylated glycosydated crotoxigenin (**56**) in MeOD-*d*₄



Supplementary data 41: COSY spectrum for acetylated glycosydated crotoxigenin (**56**) in MeOD-*d*₄



Supplementary data 42: HMBC for acetylated glycosydated crotoxigenin (**56**) in MeOD-*d*₄

Single Mass Analysis

Tolerance = 20.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

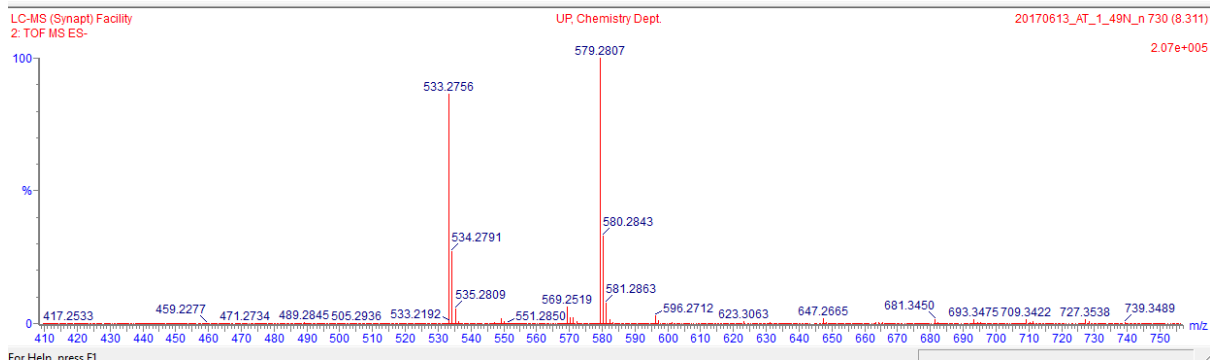
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

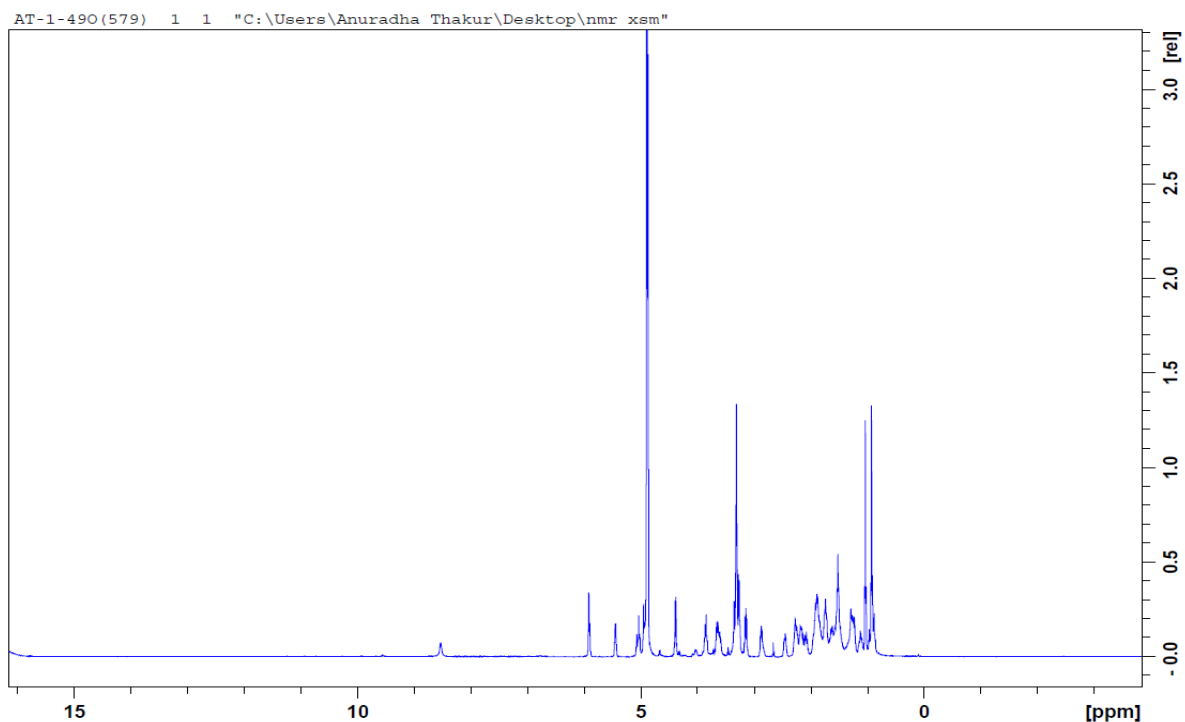
141 formula(e) evaluated with 4 results within limits (all results (up to 1000) for each mass)

Elements Used:

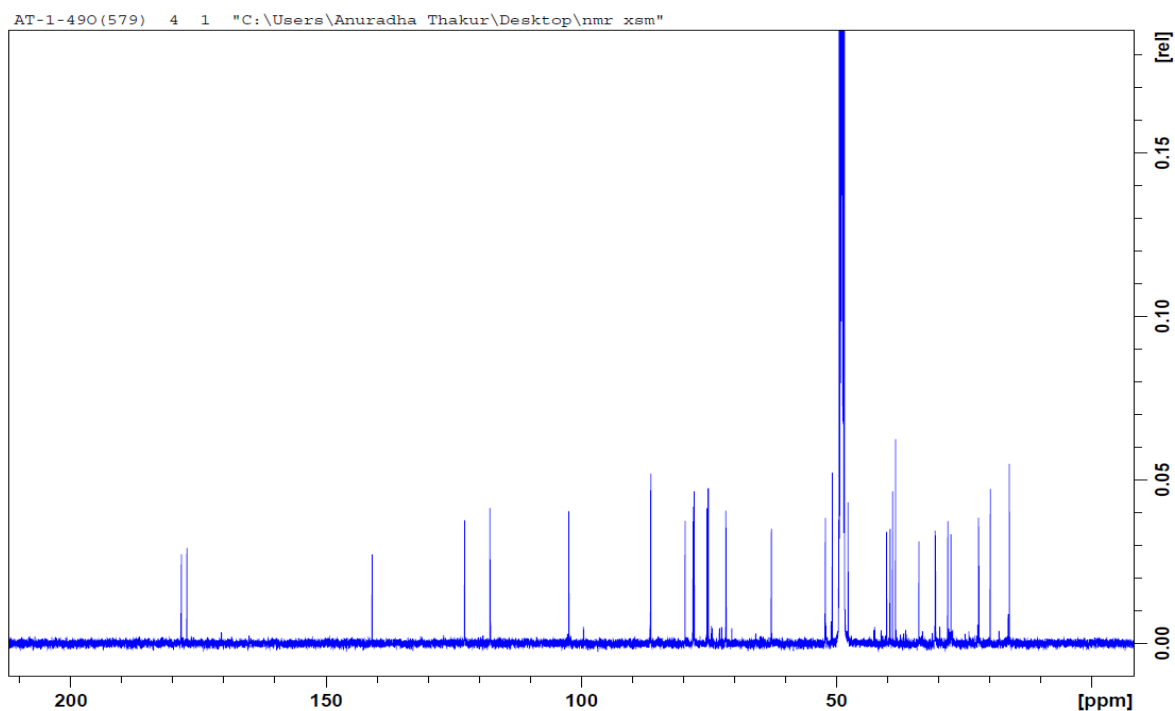
Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FIT	i-FIT Norm	Fit Conf %	C	H	O
533.2756	533.2751	0.5	0.9	9.5	C29 H41 O9	286.5	0.001	99.94	29	41	9
533.2809	533.2809	-5.3	-9.9	0.5	C22 H45 O14	294.0	7.476	0.06	22	45	14
533.2692	533.2692	6.4	12.0	18.5	C36 H37 O4	296.5	9.998	0.00	36	37	4
533.2844	533.2844	-8.8	-16.5	22.5	C40 H37 O	299.0	12.519	0.00	40	37	1



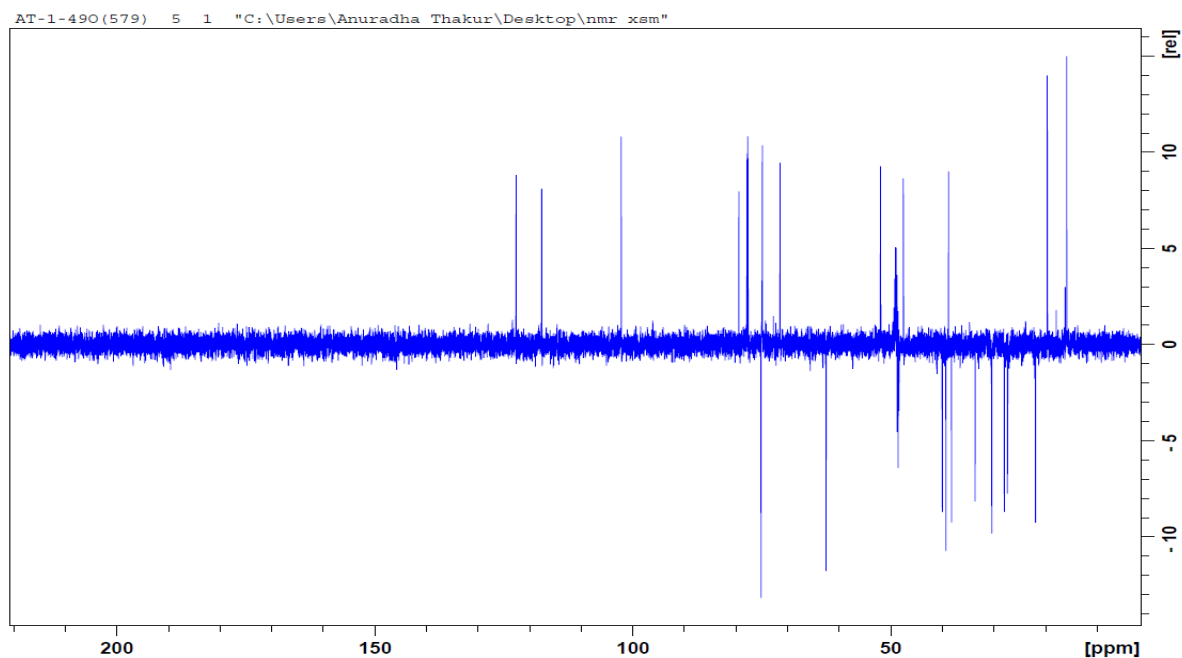
Supplementary data 43: iFit value of Xysmalogenin-3, β -D-glucopyranoside (**57**) in *X. undulatum* leaves extracted with DCM:MeOH



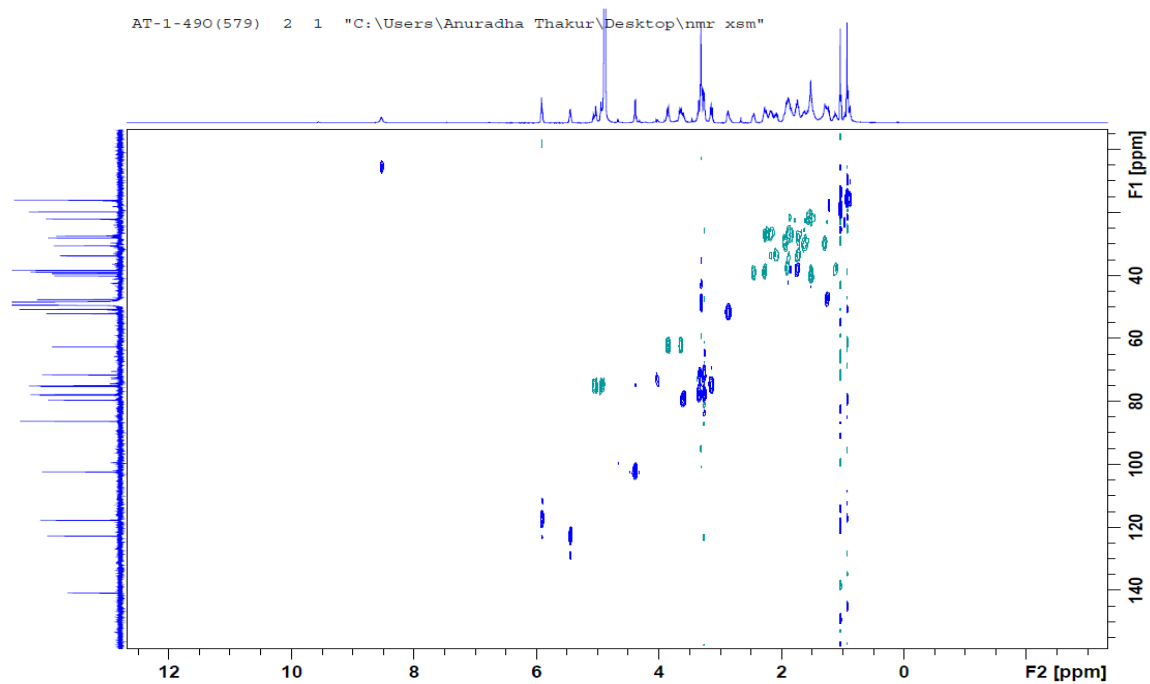
Supplementary data 44: ^1H NMR for Xysmalogenin-3, β -D-glucopyranoside (**57**) in $\text{MeOD-}d_4$



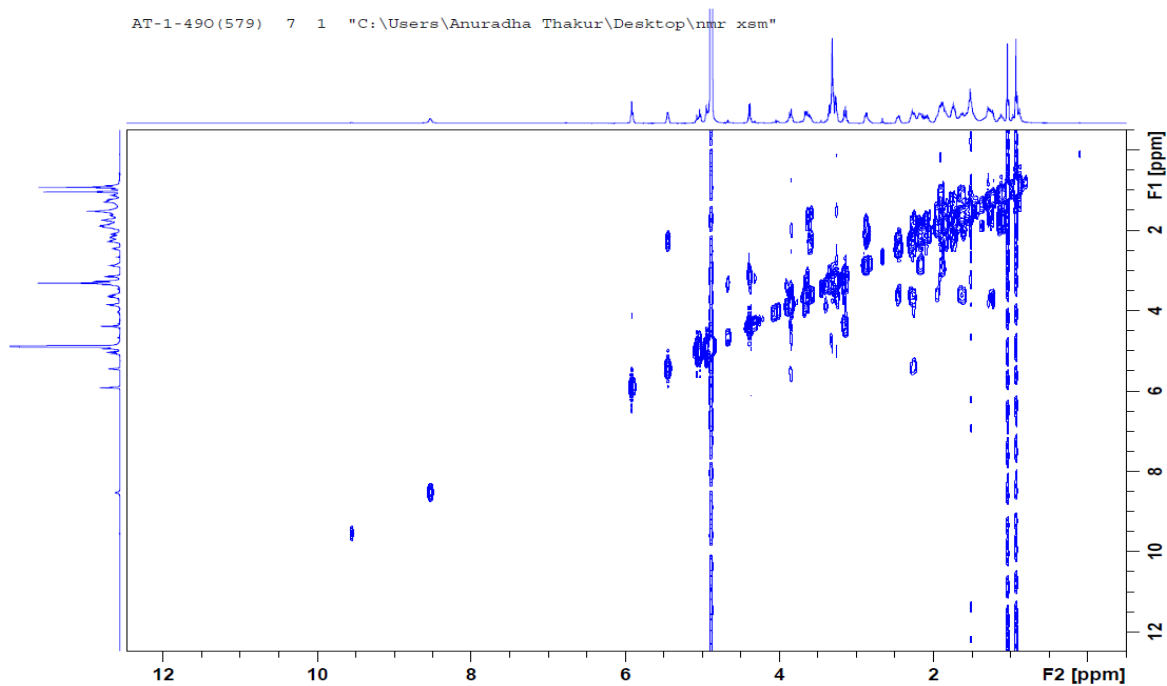
Supplementary data 45: ^{13}C NMR for compound Xysmalogenin-3, β -D-glucopyranoside (**57**) in MeOD- d_4



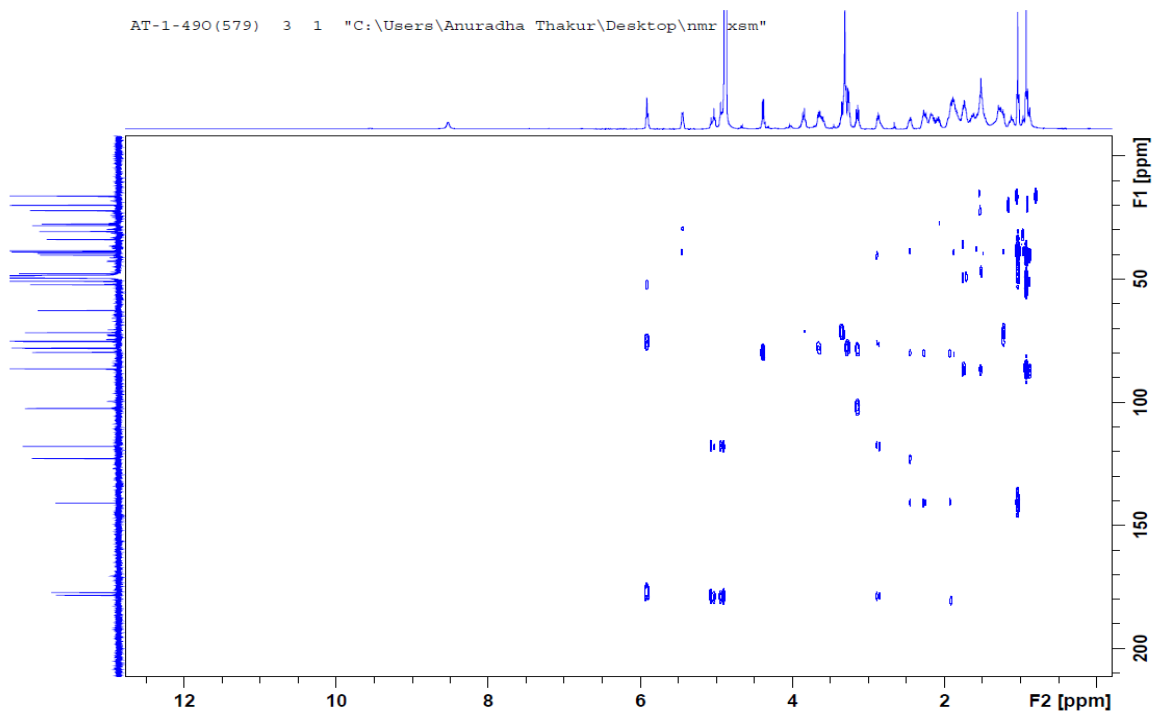
Supplementary data 46: DEPT 135 for Xysmalogenin-3, β -D-glucopyranoside (**57**) in MeOD- d_4



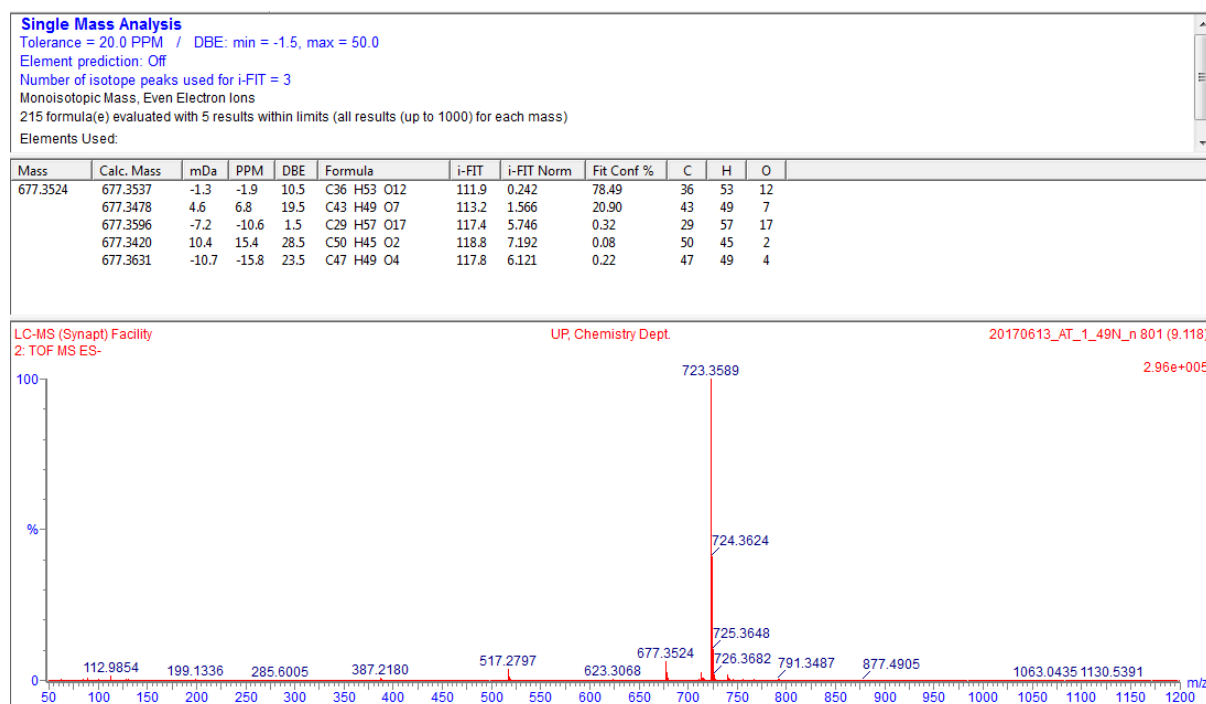
Supplementary data 47: HSQC for Xysmalogenin-3, β -D-glucopyranoside (**57**) in MeOD-*d*₄



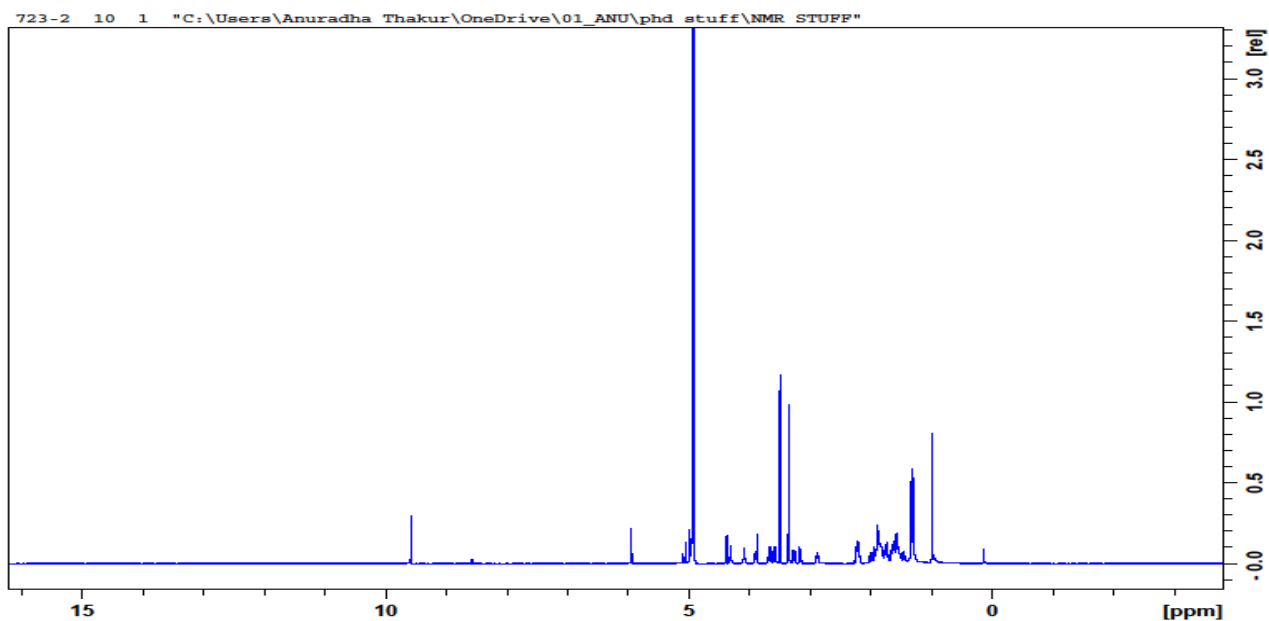
Supplementary data 48: COSY spectrum for Xysmalogenin-3, β -D-glucopyranoside (**57**) in MeOD-*d*₄



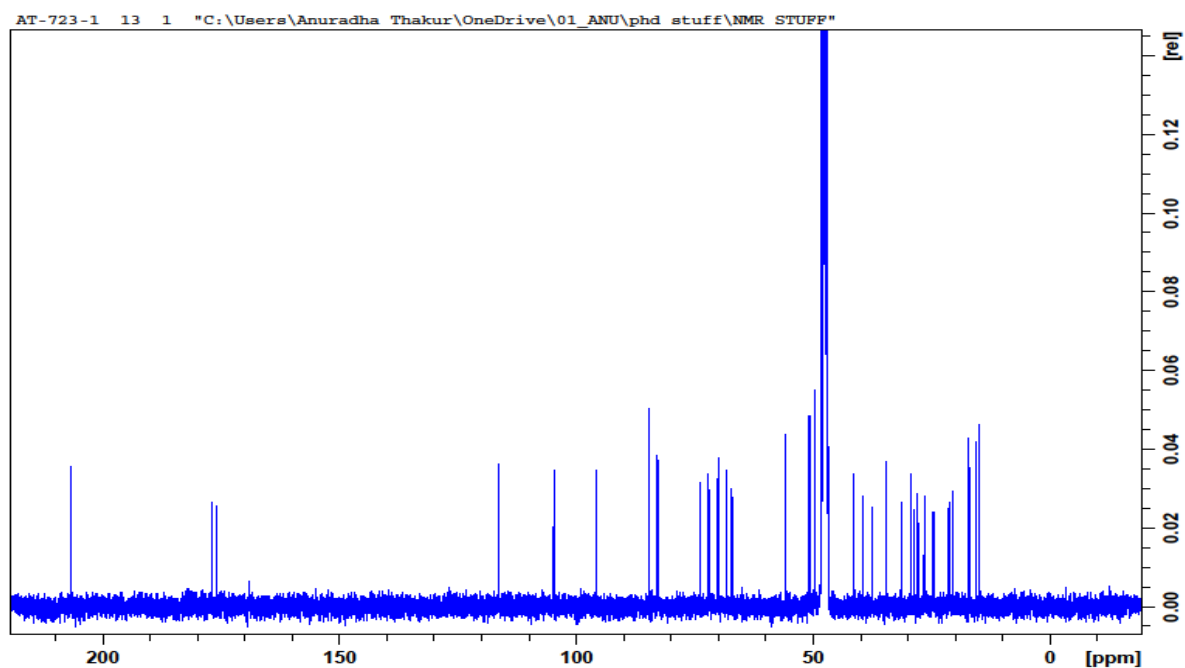
Supplementary data 49: COSY spectrum for Xysmalogenin-3, β -D-glucopyranoside (**57**) in MeOD- d_4



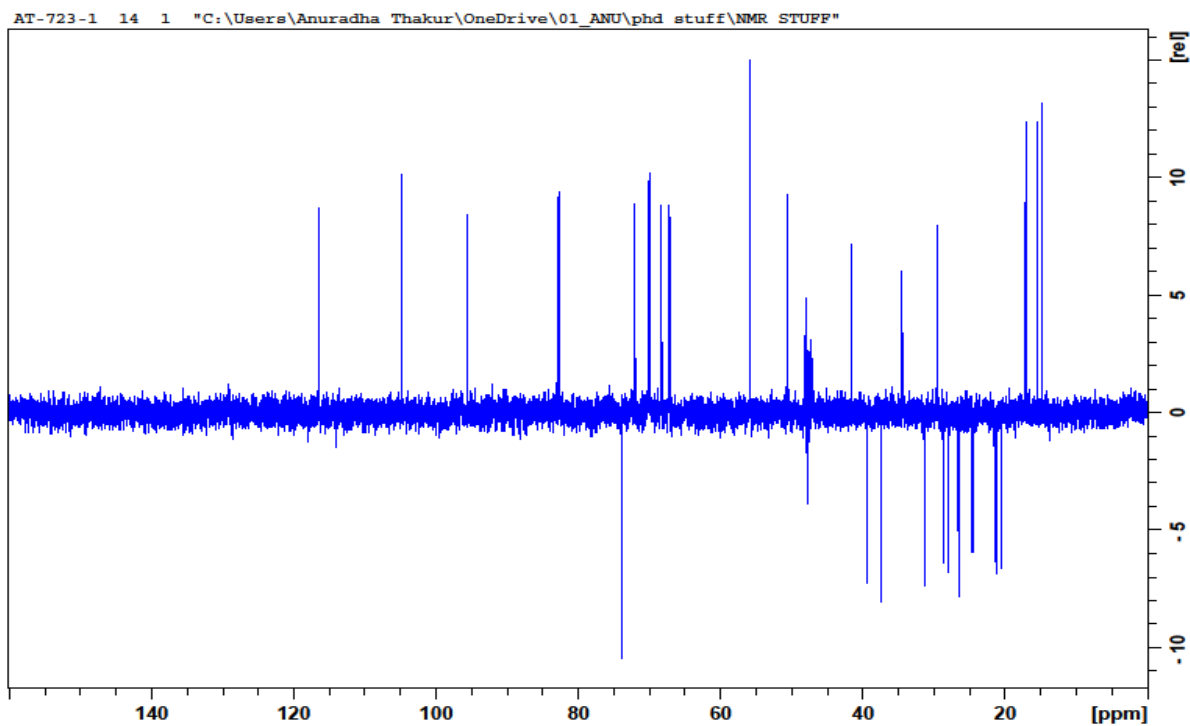
Supplementary data 50: iFit value of Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyranoside (**58**) in *X. undulatum* leaves extracted with DCM:MeOH



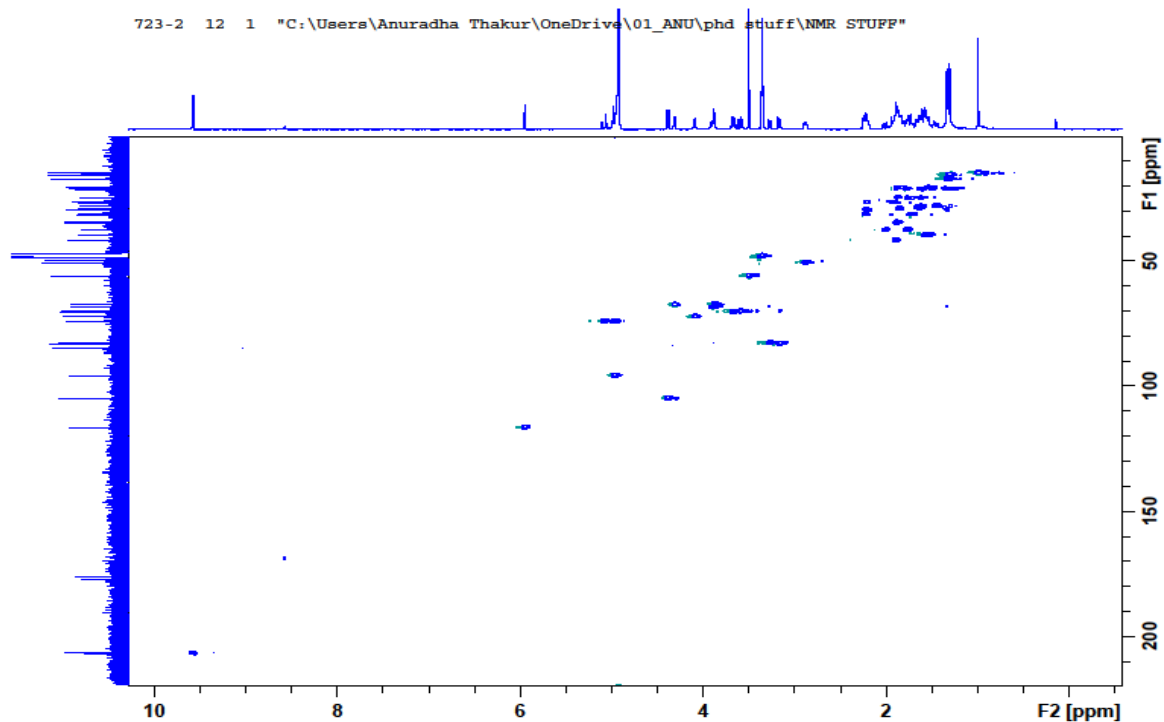
Supplementary data 51: ^1H NMR for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD- d_4



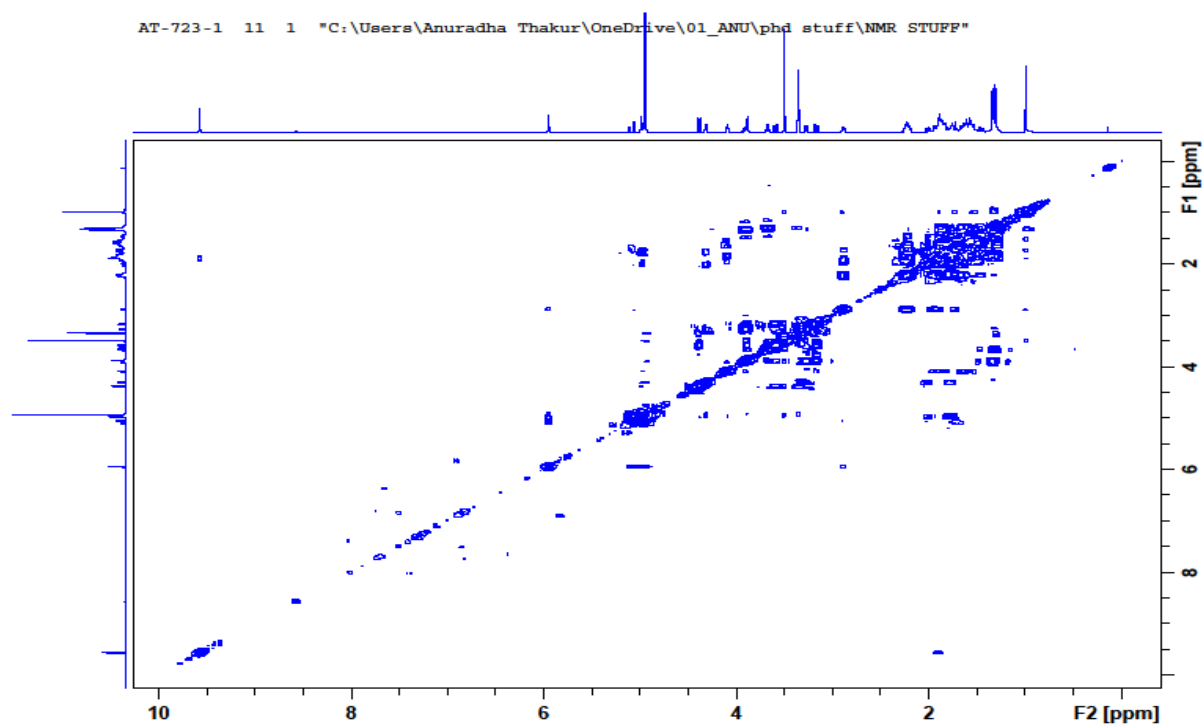
Supplementary data 52: ^{13}C NMR for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD- d_4



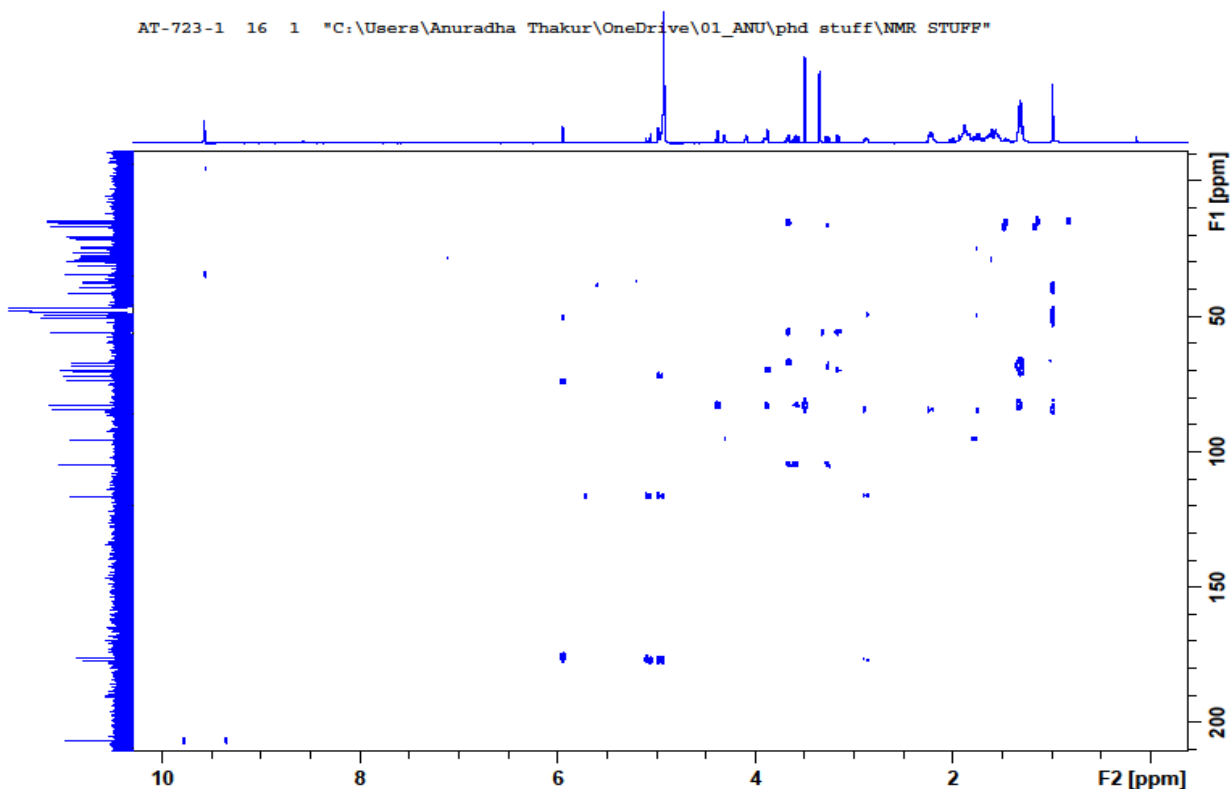
Supplementary data 53: DEPT 135 for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD-*d*₄



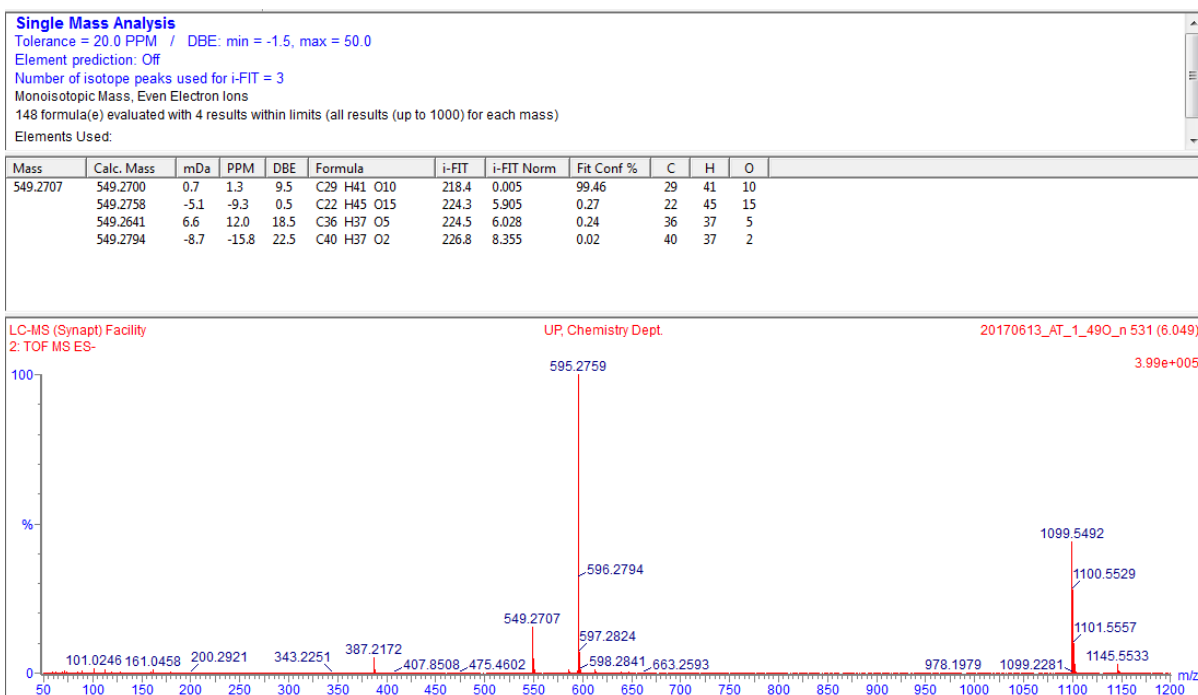
Supplementary data 54: HSQC spectrum for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD-*d*₄



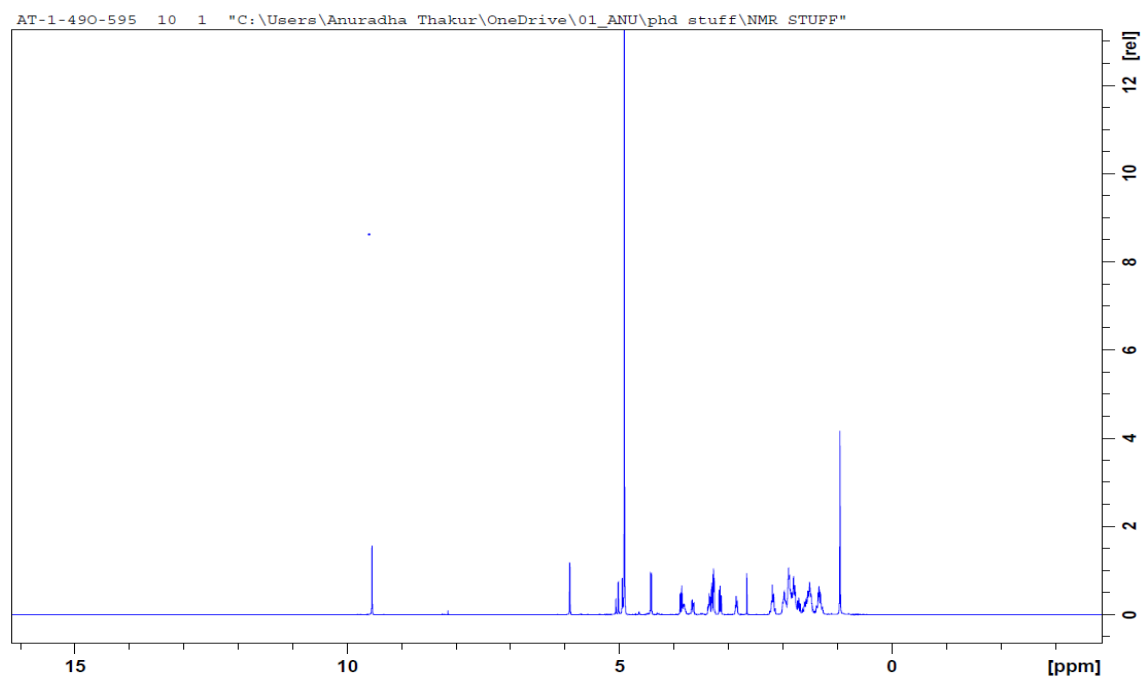
Supplementary data 55: COSY spectrum for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD-*d*4



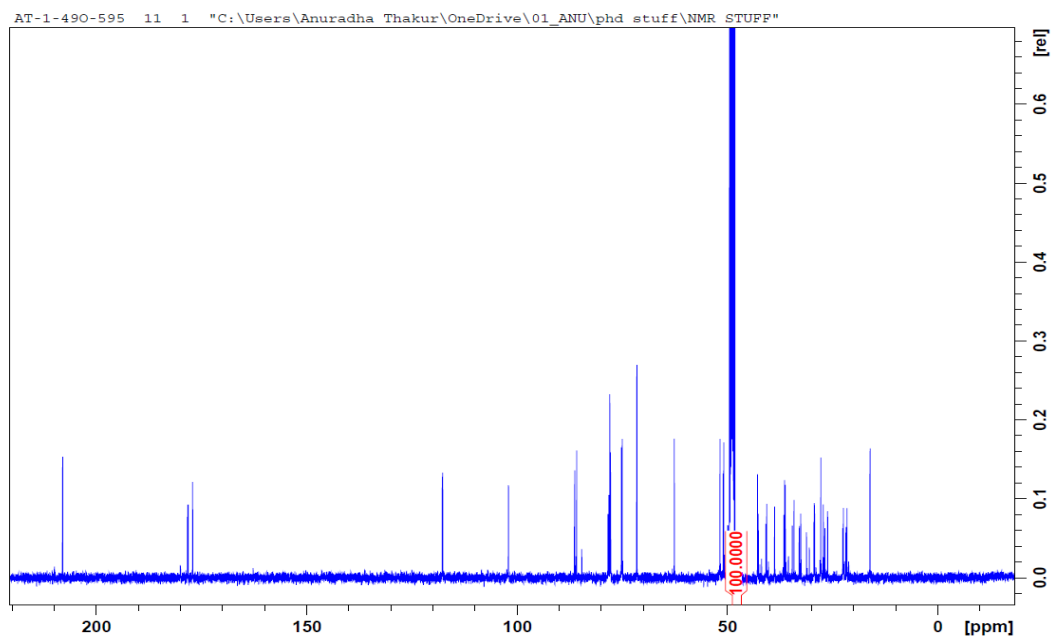
Supplementary data 56: HMBC for Crotoxigenin-3-O- β -digitalopyranosyl-(1-4)-O- β -digitoxopyanoside (**58**) in MeOD-*d*4



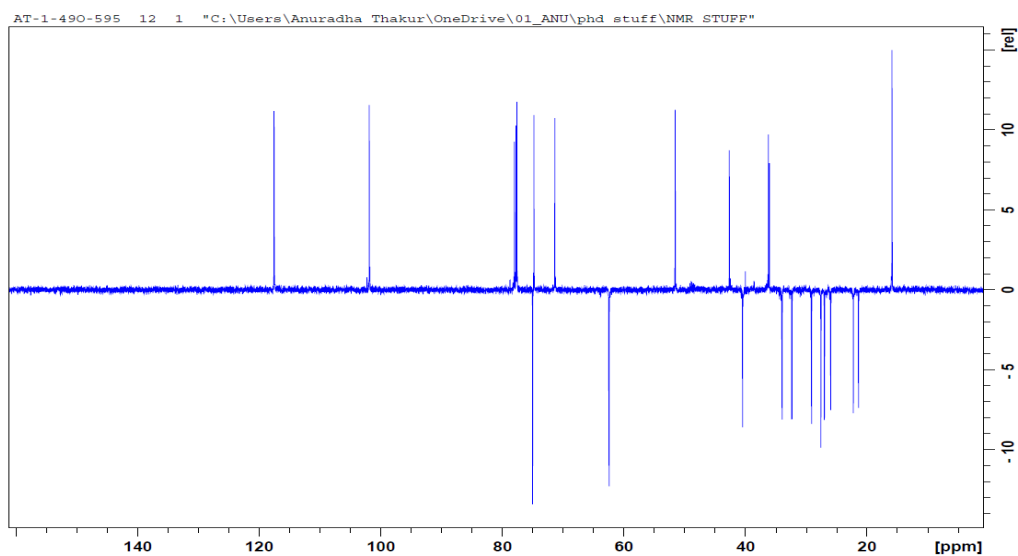
Supplementary data 57: iFit value of crotoxigenin 3-O-glucopyranoside (**59**) in *X. undulatum* leaves extracted with DCM:MeOH



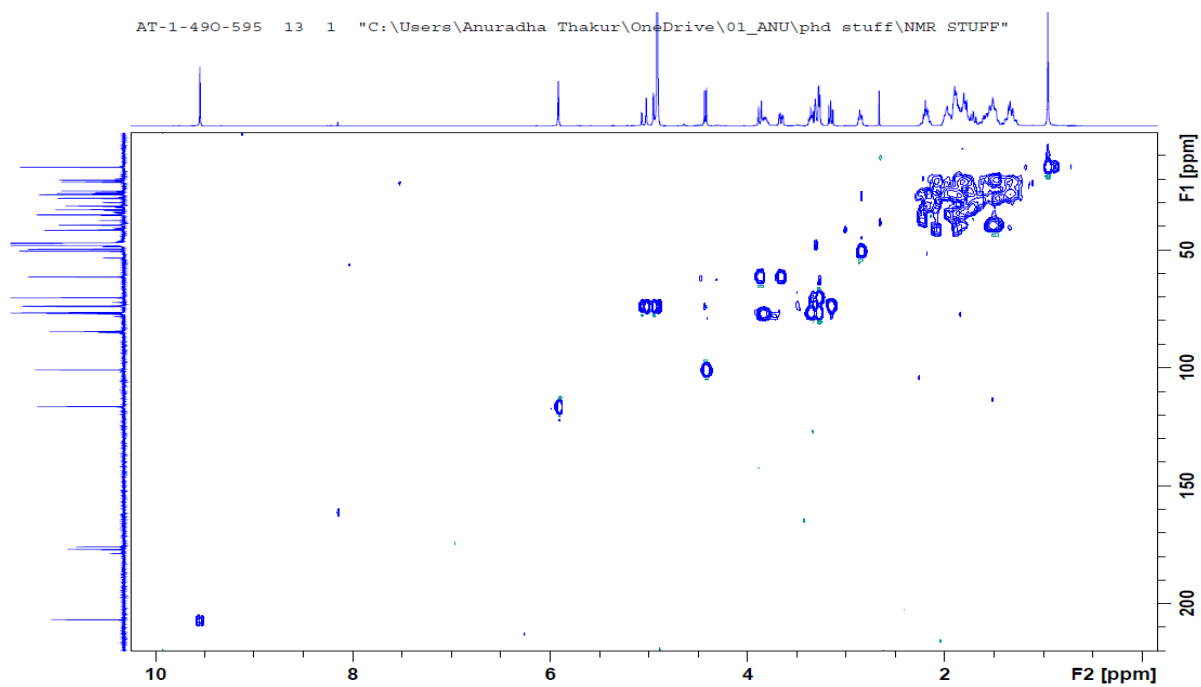
Supplementary data 58: ¹H NMR for crotoxigenin 3-O-glucopyranoside (**59**) in MeOD-*d*₄



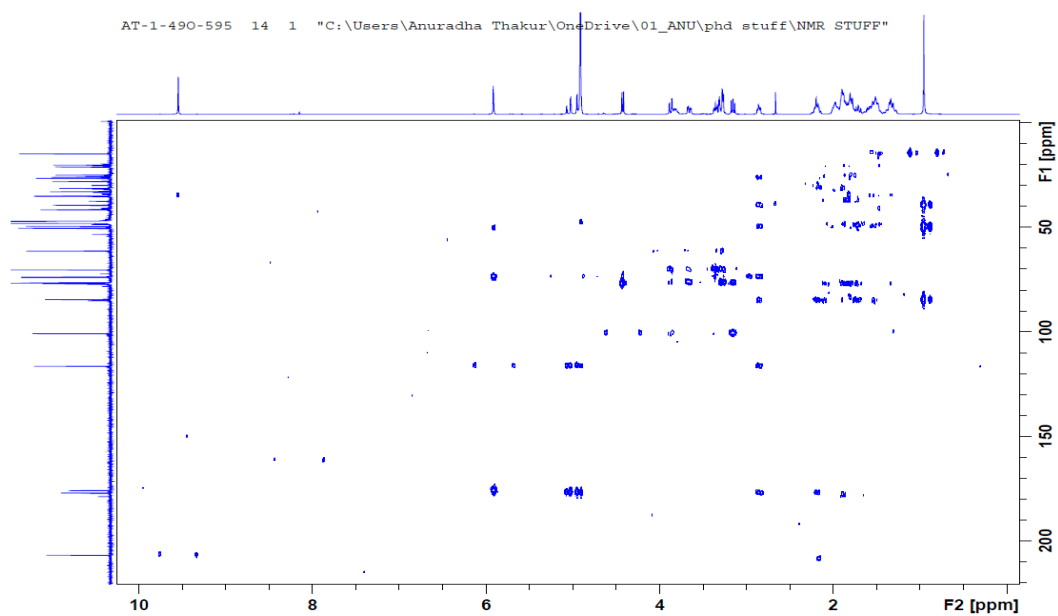
Supplementary data 59: ^{13}C NMR for crotoxinigenin 3-O-glucopyranoside (**59**) in MeOD- d_4



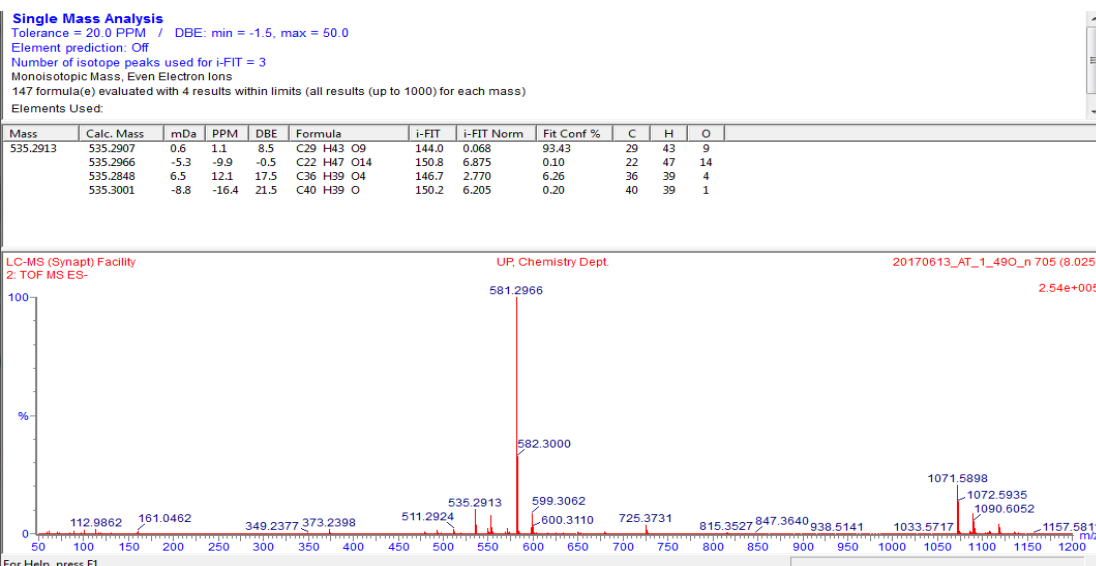
Supplementary data 60: DEPT 135 for crotoxinigenin 3-O-glucopyranoside (**59**) in MeOD- d_4



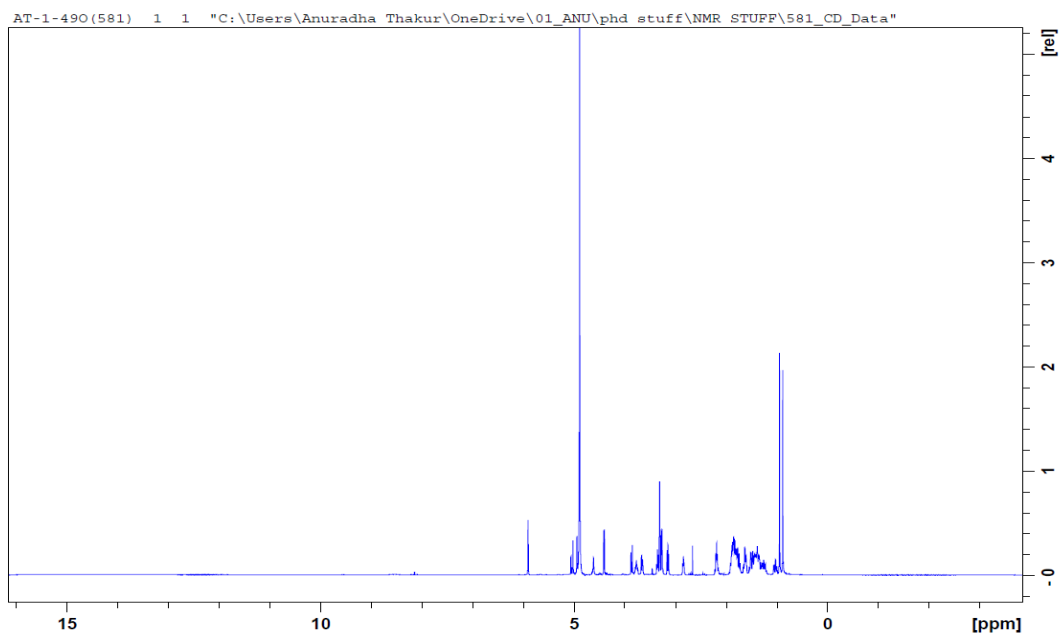
Supplementary data 61: HSQC spectrum for crotoxigenin 3-O-glucopyranoside (**59**) in MeOD-*d*4



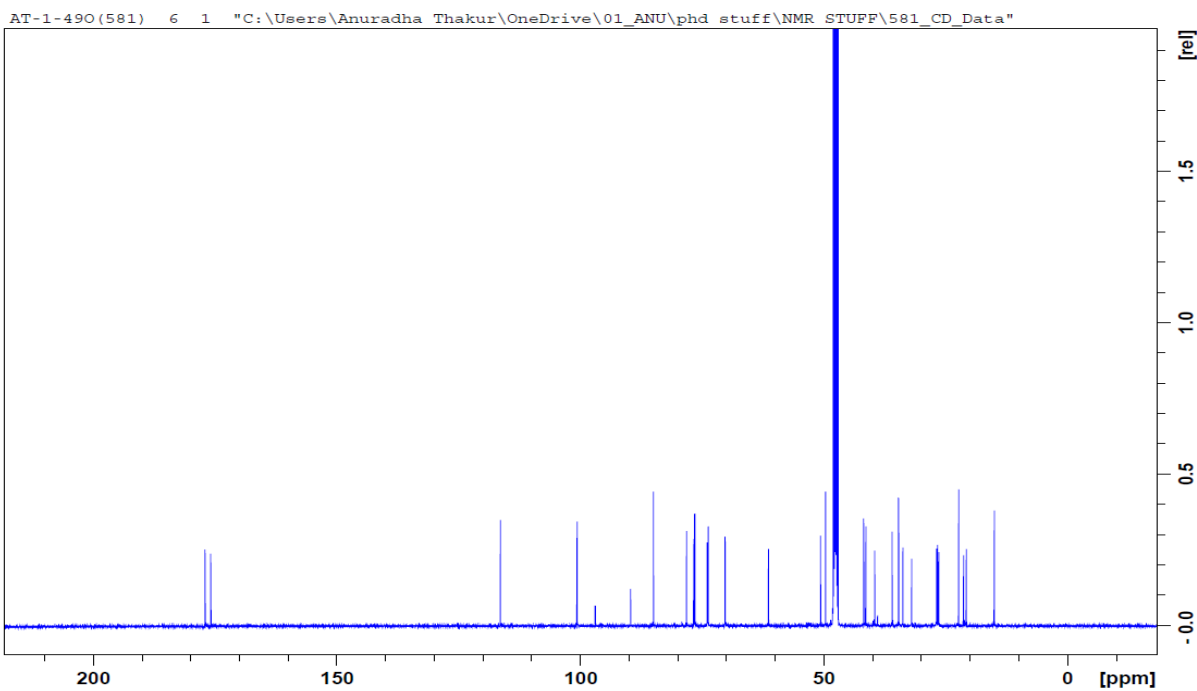
Supplementary data 62: HMBC spectrum for crotoxigenin 3-O-glucopyranoside (**59**) in MeOD-*d*4



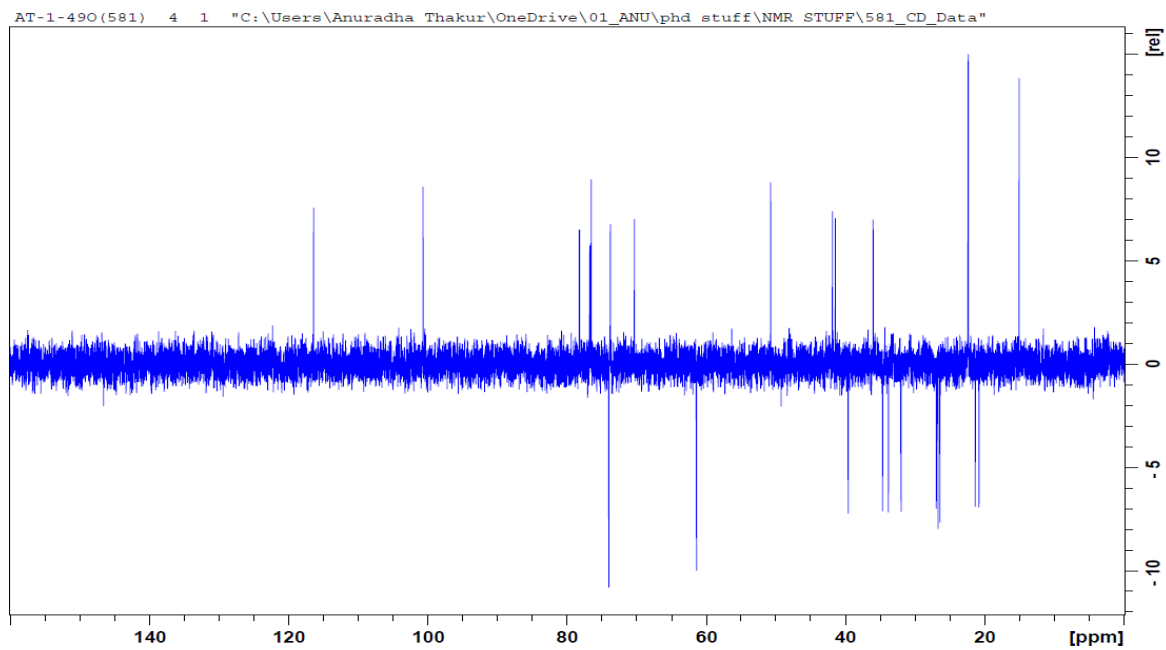
Supplementary data 63: iFit value of desglucouzarin (**60**) in *X. undulatum* leaves extracted with DCM:MeOH



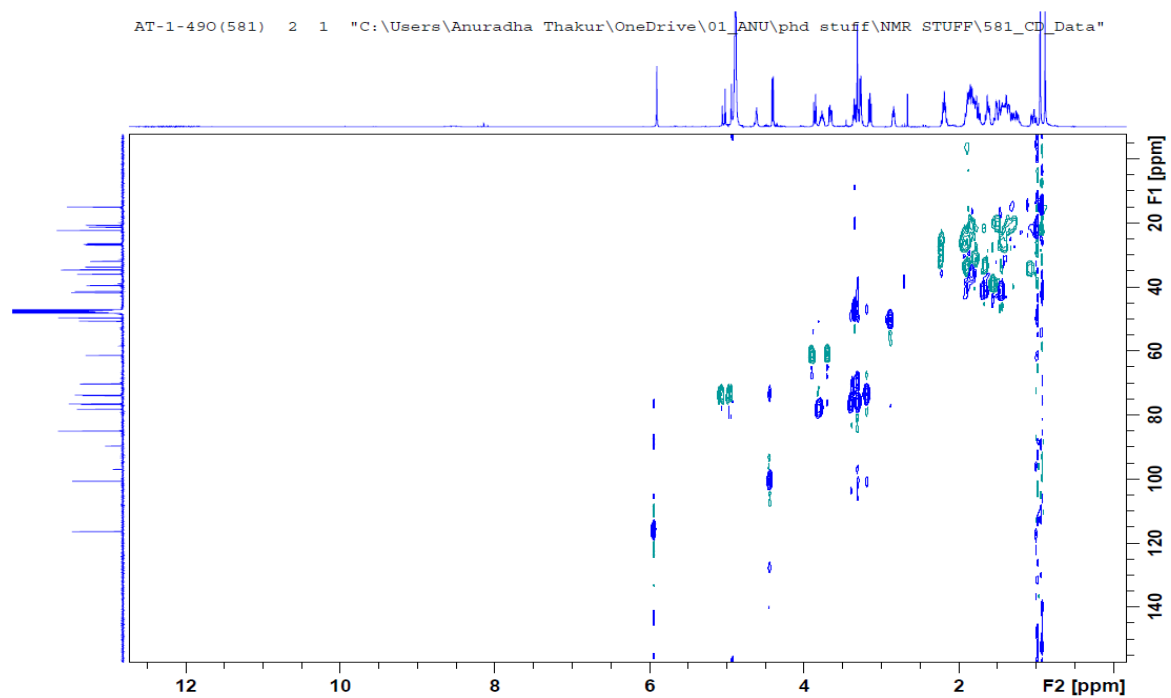
Supplementary data 64: ¹H NMR for desglucouzarin (**60**) in MeOD-d₄



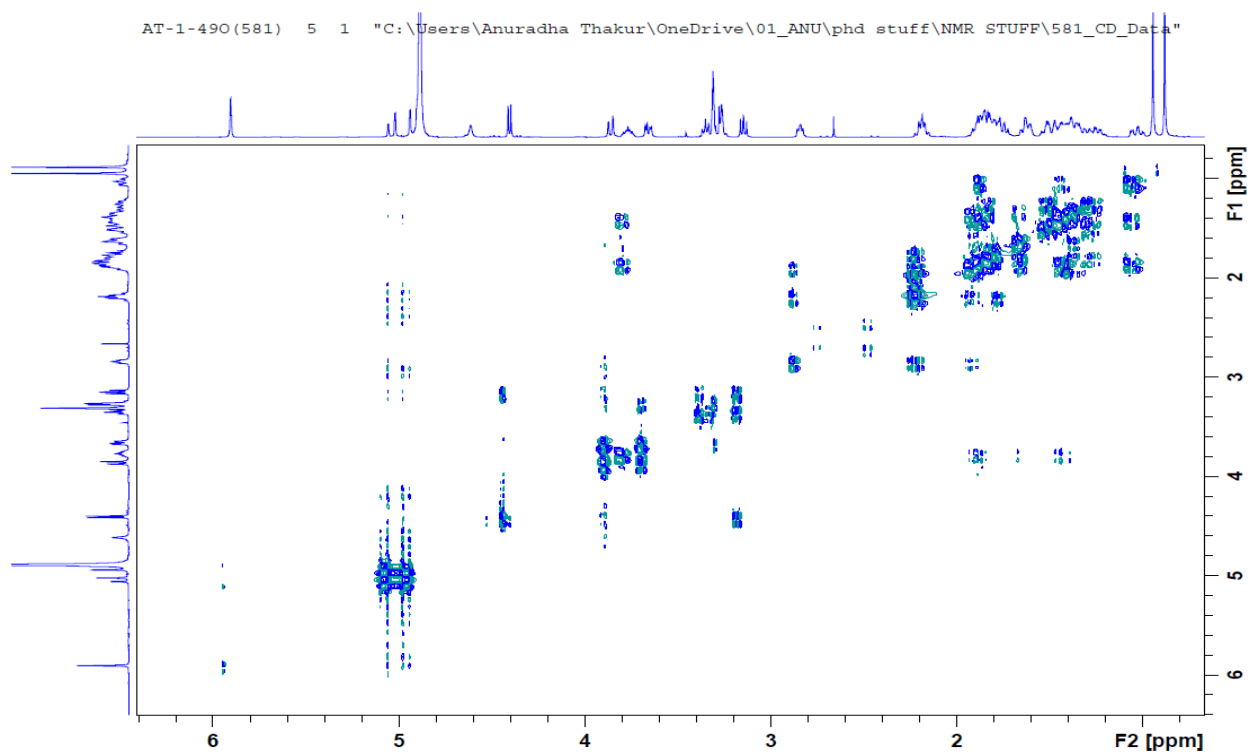
Supplementary data 65: ¹³C NMR for desglucouzarin (**60**) in MeOD-*d*₄



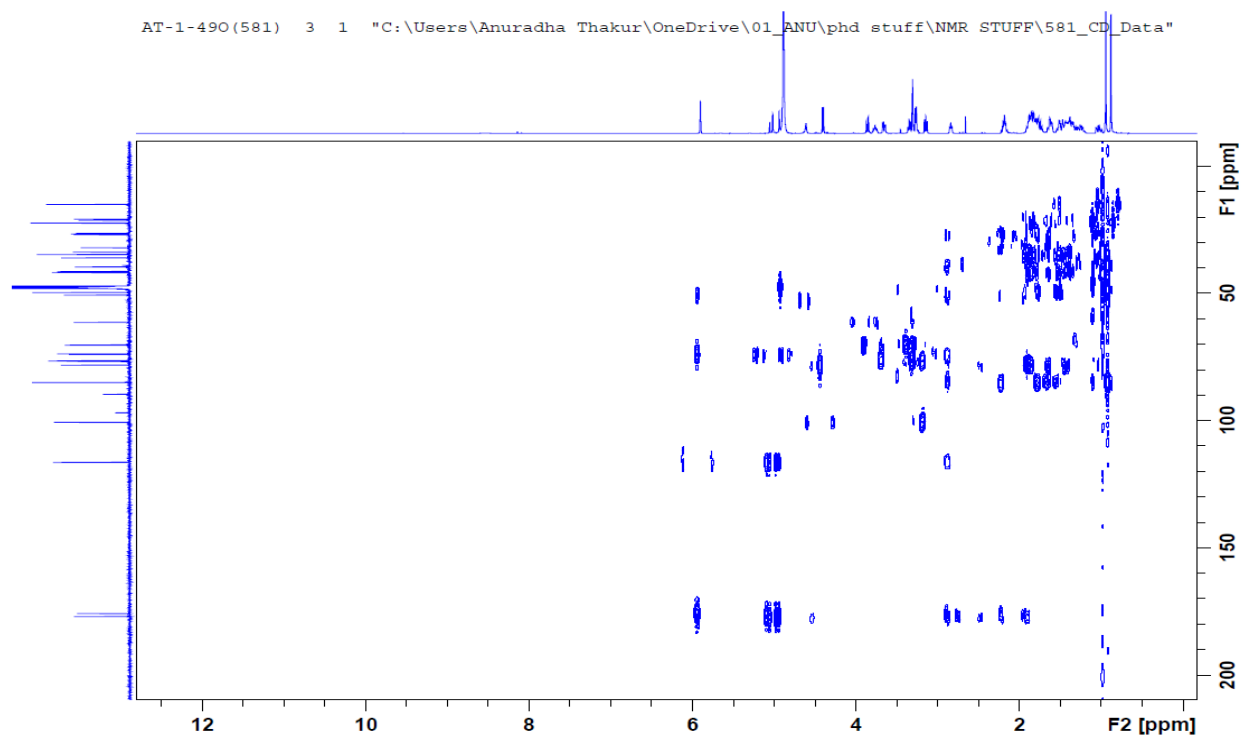
Supplementary data 66: DEPT 135 for desglucouzarin (**60**) in MeOD-*d*₄



Supplementary data 67: HSQC spectrum for desglucouzarin (**60**) in MeOD-*d*4



Supplementary data 68: ^1H NMR for desglucouzarin (**60**) in MeOD-*d*4



Supplementary data 69: ^1H NMR desglucouzarin (**60**) in $\text{MeOD-}d_4$