## PHYTOCHEMICAL, CHEMOPREVENTIVE AND ANTIMALARIAL ACTIVITY EVALUATION OF FIVE SELECTED MEDICINAL PLANTS FROM THE CAMEROONIAN FLORA

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i

Dedicated

70

Brienne and Ethan

"If you want to shine like a sun, first burn like a sun"

A.P.J. Abdul Kalam

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#### Abstract

Screening of ethnomedicinal plants for antimalarial and chemopreventive activities among plants used in Cameroonian folklore medicine to treat fevers, malaria and tumour was conducted on Croton oligandrus (Euphorbiaceae), Justicia hypocrateriformis (Acanthanceae), Pseudospondias microcarpa (Anacardiaceae), Zanthoxylum lepreurii and Zanthoxylum zanthoxyloides (Rutaceae). The selection of plants was based on ethnomedicinal use and literature review. Bioassay-guided isolation of active components from the active plants was performed with the aim to scientifically validate their folklore usage. The plant materials were dried, ground, and Soxhlet-extracted, successively, with n-hexane, dichloromethane (DCM) and methanol (MeOH). The antimalarial and chemopreventive potential of the plants were evaluated using haem polymerisation and the luciferase assays, respectively. The *n*-hexane and the DCM extracts of *C. oligandrus*, and the DCM and MeOH extracts of Z. zanthoxyloides were the most active in the luciferase assay causing 18, 21, 34 and 36-fold induction of the level of luciferase in AREc32 cells, respectively. *Pseudospondias microcarpa* was the most active antimalarial plant with IC<sub>50</sub> values of 73.9  $\pm$  25.8, 2.5  $\pm$  1.5 and 4.0  $\pm$  1.7  $\mu$ M for the stem bark *n*hexane, DCM and MeOH extracts, respectively, and  $13.0 \pm 9.0 \,\mu$ M for the leaves MeOH extract. Fifty-one compounds including eleven novel ones were isolated from active fractions using column chromatography, thin layer chromatography and reversed-phase high-pressure liquid chromatography. Structural elucidation was carried out by spectroscopic means including 1D and 2D NMR and mass spectrometry. Crotonolins A-D, skimmianine, hesperidin and myrtopsine were identified as potent chemopreventive compounds with fold induction greater than 2. None of the isolated compounds demonstrated any inhibition of haem polymerisation. This study generated the first phytochemical report on *P. microcarpa*, the second of *C. oligandrus* and the third of *J.* 

hypocrateriformis. The biogenesis and chemotaxonomy of isolated compounds have also

been discussed.

## Table of contents

Dedicatio	)n	ii
Acknowl	edgementsi	ii
Abstract	i	v
Table of	contents	⁄i
List of al	obreviationsx	ii
List of fig	guresx	v
List of ta	blesx	X
Chapter	1 Introduction	1
1.1 Ca	incer	2
1.1.1	General overview	2
1.1.2	Mechanism of cancer development	2
1.1.3	Cancer occurrence and mortality rates	4
1.1.4	Cancer chemotherapy	4
1.1.5	Treatment of cancer by traditional medicine	6
1.1.6	Cancer chemoprevention	6
1.1.7	Chemoprevention by dietary phytochemicals	8
1.1.8	Phase II detoxification enzyme1	0
1.1.9	Nrf2 as a target for cancer chemoprevention1	1
1.2 Ma	alaria1	2
1.2.1	General overview	2
1.2.2	Malaria occurrence and mortality rate1	3
1.2.3	Life cycle of malaria <i>Plasmodium</i> 1	4
1.2.4	Current antimalarial drugs and prevention1	7
1.2.4	.1 Quinine and its derivatives	7
1.2.4	.2 Antifolates	8
1.2.4	.3 Artemisinin and related drugs1	9

1.2.4.4	Other used antimalarial drugs	20	
1.2.5	Malaria prevention	21	
1.2.6	Malaria drug resistance	21	
1.2.6.1	Resistance to quinoline antimalarials	22	
1.2.6.2	2 Resistance to artemisinin and its derivatives	23	
1.2.6.3	Resistance to antifolates	23	
1.2.7	Mechanism of action of chloroquine: hemozoin as target for new		
antimala	arials development	24	
1.2.8	Malaria and natural products	25	
1.3 Nat	ural products drug discovery	26	
1.3.1	Random screening	27	
1.3.2	Ethnobotanical screening	28	
1.3.3	Taxonomic and chemotaxonomic screening	30	
1.3.4	Virtual screening	30	
1.4 Can	neroonian medicinal plants and their potential	31	
1.5 Sele	cted Cameroonian medicinal plants for this study	32	
1.5.1	Croton oligandrus Pierre ex Hutch	32	
1.5.2	Ruspolia hypocrateriformis (Vahl) Milne-Redh	35	
1.5.3	Pseudospondias microcarpa (A. Rich.) Engl	37	
1.5.4	Zanthoxylum lepreurii Guill. and Perr.	40	
1.5.5	Zanthoxylum zanthoxyloides (Lam.) Zepern. and Timler	48	
1.6 Aim ar	nd objectives	59	
Chapter 2	Materials and Methods	60	
2.1 Plant r	naterials	62	
2.2 Chemi	2.2 Chemicals and reagents		
2.2.1 Ch	emicals	63	
2.2.2 Ce	2.2.2 Cell lines, cell culture media and reagents		

2.3 F	2.3 Phytochemical work		. 65
2.3	3.1 Pla	nt preparation and Soxhlet extraction	. 65
2.3	2.3.2 Chromatographic techniques		
	2.3.2.1	Vacuum liquid chromatography	. 66
	2.3.2.2	Solid-phase extraction	. 67
	2.3.2.3	Thin layer chromatography	. 68
	2.3.2.4	Column chromatography	. 70
	2.3.2.5	High-performance liquid chromatography	.71
2.3	3.3	Isolation of compounds	.72
	2.3.3.1	Isolation of compounds from Croton oligandrus	.72
	2.3.3.2	Isolation of compounds from Justicia hypocrateriformis	.75
	2.3.3.3	Isolation of compounds from <i>Pseudospondias microcarpa</i>	.77
	2.3.3.4	Isolation of compounds from Zanthoxylum leprieurii	. 79
	2.3.3.5	Isolation of compounds from Zanthoxylum zanthoxyloides	. 81
2.3	3.4	Identification and characterisation of isolated compounds	. 84
	2.3.4.1	Nuclear magnetic resonance	. 84
	2.3.4.2	Ultraviolet-visible and Fourier transform infrared spectroscopies	. 85
	2.3.4.3	Optical rotation	. 85
	2.3.4.4	Mass spectrometry	. 86
2.4	Bioa	etivity studies	. 86
2.4	4.1	Samples preparation for screening	. 86
2.4	4.2	Cell culture	. 87
2.4	4.3	Cell viability assays	. 88
2.4	4.4	Luciferase reporter gene assay	. 89
2.4	4.5	Haem polymerisation assay	. 90
2.5 S	Statisti	cal analysis	.91

Chapter 3 Results and Discussion
<b>3.1 Yield of extraction</b>
<b>3.2 Preliminary screening</b>
<b>3.2.1 Luciferase activity</b>
3.2.1.1 AREc32 cells viability94
3.2.1.2 Luciferase assay
<b>3.2.2 Haem polymerisation assay</b>
<b>3.3 Characterisation and structure elucidation of isolated compounds</b> 101
3.3.1 Phytochemistry of Croton oligandrus101
3.3.1.1 Structure elucidation of $3-\beta$ -O-acetyl aleuritolic acid (7)
3.3.1.2 Structure elucidation of cluytyl ferulate (149) and hexacosanoyl ferulate
( <b>150</b> ) as a mixture
3.3.1.3 Structure elucidation of crotocorylifuran (151)109
3.3.1.4 Structure elucidation of megalocarpoidolide D (153) 111
3.3.1.5 Structure elucidation of 12- <i>epi</i> -megalocarpoloide D ( <b>154</b> )116
3.3.1.6 Structure elucidation of crotonolin A $(156)$ and crotonolin B $(157)$ as a
mixture
3.3.1.7 Structure elucidation of crotonolin C (158) and crotonolin D (159) as a
mixture121
3.3.1.8 Structure elucidation of crotonolin F (160) 123
3.3.1.9 Structure elucidation of 7-Oxodehydroabietic acid (161) 126
<b>3.3.2 Phytochemistry of</b> <i>Justicia hypocrateriformis</i>
3.3.2.1 Structure elucidation of luteolin-7- <i>O</i> - $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-
xylopyranoside (163)
3.3.2.2 Structure elucidation of chrysoeriol-7- <i>O</i> - $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-
xylopyranoside (164)
3.3.2.3 Structure elucidation of chrysoeriol-7- $O$ -[4'''- $O$ -acetyl- $\beta$ -D-apiofuranosyl-
$(1\rightarrow 2)$ ]- $\beta$ -D-xylopyranoside (165)

	3.3.2.4 Structure elucidation of luteolin-7- <i>O</i> - $\alpha$ -D-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-	
	xylopyranoside (166)	. 137
	3.3.2.5 Structure elucidation of chrysoeriol-7- <i>O</i> - $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -I	D-
	xylopyranoside (167)	. 141
	3.3.2.6 Structure elucidation of luteolin 7- <i>O</i> -[ $\beta$ -D-glucopyranosyl-(1 $\rightarrow$ 2)- $\alpha$ -L-	
	rhamnosyl- $(1\rightarrow 6)$ ]- $\beta$ -D-glucopyranoside (168)	. 142
	3.3.2.7 Structure elucidation of secundallerones B and C (170-171) as a mixture	2144
3	3.3 Phytochemistry of Pseudospondias microcarpa	. 149
	3.3.3.1 Structure elucidation of scopoletin (103)	. 150
	3.3.3.2 Structure elucidation of isovetexin (172)	. 151
	3.3.3.3 Structure elucidation of apigenin 7- $O$ - $\beta$ -D-neohesperidoside (173)	. 154
	3.3.3.4 Structure elucidation of pithecellobiumol B (174)	. 155
3	3.3.4 Phytochemistry of Zanthoxylum leprieurii	. 158
	3.3.4.1 Structure elucidation of kaurenoic acid (175)	. 159
	3.3.4.2 Structure elucidation of xylopic acid (176)	162
	3.3.4.3 Structure elucidation of <i>ent</i> -kauran-16 $\beta$ -ol-19-oic acid ( <b>177</b> )	. 164
	3.3.4.4 Structure elucidation of <i>ent</i> -kauran-16 $\beta$ -ol ( <b>179</b> )	. 165
	3.3.4.5 Structure elucidation of dulcisflavane (180)	. 170
	3.3.4.6 Structure elucidation of icariside D2 (182)	. 170
3	3.3.5 Phytochemistry of Zanthoxylum zanthoxyloides	. 176
	3.3.5.1 Structure elucidation of skimmianine (55)	. 176
	3.3.5.2 Structure elucidation of <i>N</i> -methylplatydesminium cation ( <b>183</b> )	. 179
	3.3.5.3 Structure elucidation of isoplatydesmine (184)	. 181
	3.3.5.4 Structure elucidation of ribalinine (186)	. 182
	3.3.5.5 Structure elucidation of atanine (128)	. 184
	3.3.5.6 Structure elucidation of <i>N</i> -methylatanine ( <b>187</b> )	. 185
	3.3.5.7 Structure elucidation of zanthoamides G (188)	. 188
	3.3.5.8 Structure elucidation of zanthoamides H (189)	. 192

3.3.5.9 Structure elucidation of zanthoamides I (190)	93
3.3.5.10 Structure elucidation of hesperetin (191)	97
3.4 Biogenesis and chemotaxonomic significance of isolated compounds	00
3.4.1 Alkaloids	00
<b>3.4.2 Diterpenes</b>	01
<b>3.4.3 Flavonoids</b>	03
3.5 Bio-activity of the isolated compounds	05
3.5.1 Cytotoxicity of isolated compounds	05
3.5.2 Chemopreventive activity of isolated compounds	07
3.5.2.1 Cytotoxicity of the compounds against AREc32 cells	07
3.5.2.2 Luciferase activity	08
3.5.3 Haem polymerisation assay of the isolated compounds	09
Chapter 4 Conclusion and Future Prospects	11
4. Conclusion and future prospects	12
References	15
Appendix I MS, <sup>1</sup> H and <sup>13</sup> C NMR spectra of crotonolin E (155)	45
Appendix II ESI-MS spectrum of justivialoside B (167)24	46
Appendix III ESI-MS spectrum of scopoletin (103)24	47
Appendix IV <sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) spectrum of myrtopsine (185)	47
Appendix V List of publications and presentations	48

## List of abbreviations

δ	Chemical shift (ppm)
<sup>1</sup> H-NMR	Proton nuclear magnetic resonance
<sup>13</sup> C-NMR	Carbon nuclear magnetic resonance
1D, 2D	One dimensional, two dimensional
Acetone-d <sub>6</sub>	Deuterated acetone
ACT	Artemisinin combined therapy
ADP:	Adenosine di-phosphate
AMU	Atomic mass units
ATP	Adenosine triphosphate
ARE	Antioxidant response element
B.C.	Before christ
Ca.	Around
CC	Open column chromatography
CD <sub>3</sub> OD	Methanol deuterated
CDCl <sub>3</sub>	Deuterated chloroform
cm	Centimeter
CNH	Cameroon national herbarium
COB	Croton oligandrus bark
conc	Concentration
COSY	Correlation spectroscopy
Cpd (s)	Compound (s)
CPP	Copalyl diphosphate
CR-UK	Cancer research United Kingdom
d	Doublet
DBE	Double bond equivalence
DCM	Dichloromethane
DMEM:	Dulbecco's modified eagle medium
DEPT	Distortionless Enhancement by Polarisation Transfer
DMSO	Dimethyl sulfoxide
DMSO-d <sub>6</sub>	Deuterated dimethyl sulfoxide
DNA	Deoxyribonucleic acid
Doxy	Doxorubicin
ESI	Electrospray ionisation

EtOAc	Ethyl acetate
FBS:	Fetal bovine serum
Fig.	Figure
FT-IR:	Fourier transform infra-red
g	Gram
GBP	Great British pound
GGPP	Geranyl geranyl diphosphate
h	Hour
HMBC	Heteronuclear multiple bond coherence
HPLC	High performance liquid chromatography
HR	High resolution
HSQC	Heteronuclear single quantum coherence
Hz	Hertz
IC <sub>50</sub>	Concentration needed to produce 50% of haem polymerisation
	inhibition
Inj. Vol.	Injection volume
J	Coupling constant
JHL	Justicia hypocrateriformis leaves
Keapl	Kelch-like ECH-associated protein 1
L	Liter
LC <sub>10</sub>	Concentration needed to produce 10% of cells death
LC <sub>50</sub> (LD <sub>50</sub> )	Concentration (or dosage) needed to produce 50% of cells death
LC/MS	Liquid chromatography-mass spectrometry
LCT	Liquid Chromatography Time of Flight
m	Multiplet (for NMR spectrum)
Μ	Molar (concentration)
m/z	Mass to charge ratio
MeOH	Methanol
mg	Milligram
MHz	Mega hertz
min	Minute
mL	Millilitre
mm	Millimetre
MS	Mass spectroscopy

MTT	3-(4,5-Dimethyithiazo-2-yl)-2,5-diphenyltetrazolium
nm	Nanometer
NMR	Nuclear magnetic resonance
NOESY	Nuclear Overhauser Enhancement spectroscopy
NQOI	Quinone reductase
Nrf2	Nuclear factor erythroid 2-related factor
OD	Optical density
OV	overlapping
PBS	Phosphate buffer saline
PMB	Pseudospondias microcarpa bark
PML	Pseudospondias microcarpa leaves
ppm	Parts per million
Pyr-d5	Deuterated pyridine
RBC	Red blood cell
RP	Reversed phase
S	Singlet
SDS	Sodium dodecyl sulfate
SEM	Standard error of mean
SPE	Solid phase extraction
t	Triplet
tBHQ	Tert-Butylhydroquinone
$t_R$ , $\mathbf{R}_{\mathrm{f}}$	Retention time, Retention factor
TFA	Trifluoro acetic acid
TLC	Thin layer chromatography
TOCSY	Total Correlation spectroscopy
TOF	Time of Flight
μg	Microgram
μL	Microliter
μm	Micromole
UV-DAD	Ultraviolet-Diode array detector
VLC	Vacuum liquid chromatography
WHO	World health organisation
ZLF	Zanthoxylum leprieurii fruits
ZZF	Zanthoxylum zanthoxyloides fruits

## List of figures

Fig.	Title	Page
1.1	Different stages of cancer development	3
1.2	Some anticancer dugs	5
1.3	Some chemopreventive agents	8
1.4	Multistage carcinogenesis and strategies for cancer chemoprevention	9
1.5	Geographic distribution of malaria	13
1.6	Life cycle of malaria cycle infection	16
1.7	Some quinoline antimalarials	18
1.8	Some antifolates used in malaria therapy	19
1.9	Artemisin and derivatives	20
1.10	Other antimalarials: clindamycin, lumefantrine and atovaquone	20
1.11	Quinoline antimalarials mode of action	25
1.12	Recent antiplasmodial compounds from natural source	26
1.13	Picture of a) Ptericarp and b) seeds of C. oligandrus	33
1.14	Isolated compounds from the bark of C. oligandrus	34
1.15	Young flourishing plant of R. hypocrateriformis	36
1.16	Isolated compounds from the roots of R. hypocrateriformis	36
1.17	Fruits and ptericarps of P. microcarpa	38
1.18	Dried Fruits of Z. leprieurii	40
1.19	Isolated compounds from Z. leprieurii	47
1.20	Dried Fruits of Z. zanthoxyloides	49
1.21	Identified and isolated compounds from Z. zanthoxyloides	57
2.1	Map showing Cameroon geographical position and the different	62
	regions where the plants were collected	
2.2	Soxhlet apparatus	66
2.3	VLC system	67
2.4	Solid Phase Extraction system	68
2.5	TLC plates of xylopic acid and b) skimmianine	69
2.6	CC system operated by manual fraction collection	70
2.7	Chromatogram of C. oligandrus fractions D4+D6	73
2.8	Chromatogram of J. hypocrateriformis fractions F2 and F3	76
2.9	Chromatogram of <i>P. microcarpa</i> leaves fractions F2 + F3	78

2.10	Chromatogram of Z. zanthoxyloides DCM fractions D4-D6	81
2.11	Chromatogram of Z. zanthoxyloides a) F2, and b) F3	82
2.12	MTT transformation to formazan	88
2.13	Formation of oxyluciferin	89
3.1	Pictures showing the different plant parts collected	93
3.2	Effect of tested extracts on the viability of AREc32 cells	96
3.3	Luciferase activity of the screened extracts	97
3.4	Luciferase activity of the fractions	99
3.5	Structures of isolated compounds from the bark of C. oligandrus	102
3.6	ESI-MS spectrum of 7	104
3.7	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of <b>7</b>	104
3.8	<sup>13</sup> C NMR (150 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>7</b>	105
3.9	ESI-MS spectrum of 149 and 150	107
3.10	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of <b>149</b> and <b>150</b>	107
3.11	$^{13}$ C NMR (150 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>149</b> and	108
	150	
3.12	ESI-MS spectrum of 151	110
3.13	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of compound <b>151</b>	110
3.14	$^{13}$ C NMR (150 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>151</b>	111
3.15	ESI-MS spectrum of 153	113
3.16	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> ) of <b>153</b>	113
3.17	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> ) and key <sup>1</sup> H- <sup>1</sup> H NOESY ( $\leftrightarrow$ ) and HMBC	114
	$(\rightarrow)$ correlations of <b>153</b>	
3.18	HSQC-DEPT (600 MHz, CDCl <sub>3</sub> ) of 153	114
3.19	<sup>1</sup> H- <sup>1</sup> H COSY (600 MHz, CDCl <sub>3</sub> ) of <b>153</b>	115
3.20	HMBC (300 MHz, CDCl <sub>3</sub> ) of <b>153</b>	115
3.21	<sup>1</sup> H- <sup>1</sup> H NOESY spectrum (300 MHz, CDCl <sub>3</sub> ) of <b>153</b>	116
3.22	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> ) of <b>154</b>	117
3.23	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> ) and key <sup>1</sup> H- <sup>1</sup> H NOESY ( $\leftrightarrow$ ) and HMBC	117
	$(\rightarrow)$ correlations of <b>154</b>	
3.24	<sup>1</sup> H- <sup>1</sup> H NOESY spectrum of <b>154</b>	118
3.25	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) of <b>156</b> and <b>157</b>	120
3.26	$^{13}\text{C}$ NMR (150 MHz, CD <sub>3</sub> OD) and key $^{1}\text{H-}^{1}\text{H}$ NOESY ( $\leftrightarrow$ ) and	120
	HMBC $(\rightarrow)$ correlations of <b>156</b> and <b>157</b>	

3.27	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of <b>158</b> and <b>159</b>	122
3.28	<sup>13</sup> C NMR (150 MHz, CDCl <sub>3</sub> ) and key <sup>1</sup> H- <sup>1</sup> H NOESY ( $\leftrightarrow$ ) and HMBC	122
	$(\rightarrow)$ correlations of <b>158</b> and <b>159</b>	
3.29	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of <b>160</b>	124
3.30	<sup>13</sup> C NMR (150 MHz, CDCl <sub>3</sub> ) and key <sup>1</sup> H- <sup>1</sup> H NOESY ( $\leftrightarrow$ ) and HMBC	124
	$(\rightarrow)$ correlations of <b>160</b>	
3.31	HSQC NMR spectrum (600 MHz, CDCl <sub>3</sub> ) of 160	125
3.32	HMBC NMR spectrum (600 MHz, CDCl <sub>3</sub> ) of <b>160</b>	125
3.33	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) of <b>161</b>	127
3.34	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>161</b>	127
3.35	Structures of isolated compounds from the leaves of J.	131
	hypocrateriformis	
3.36	ESI-MS spectrum of 163	133
3.37	Key HMBC correlations of 163	133
3.38	<sup>1</sup> H NMR (600 MHz, Pyr-d <sub>5</sub> ) of <b>163</b>	134
3.39	<sup>1</sup> H- <sup>1</sup> H COSY (600 MHz, Pyr-d <sub>5</sub> ) of 1 <b>63</b>	134
3.40	HSQC-DEPT (600 MHz, Pyr-d <sub>5</sub> ) of <b>163</b>	135
3.41	HMBC (600 MHz, Pyr-d <sub>5</sub> ) of <b>163</b>	135
3.42	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of 1 <b>64</b>	139
3.43	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>165</b>	139
3.44	ESI MS spectrum of 166	138
3.45	<sup>1</sup> H NMR (600 MHz, a-CD <sub>3</sub> OD, b-Pyr-d <sub>5</sub> ) and key HMBC correlations	140
	of <b>166</b>	
3.46	HSQC-DEPT (600 MHz, Pyr-d <sub>5</sub> ) of <b>166</b>	140
3.47	HMBC (600 MHz, Pyr-d <sub>5</sub> ) of <b>166</b>	141
3.48	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and keys HMBC correlations of <b>167</b>	142
3.49	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>168</b>	143
3.50	HR ESI-MS spectrum of 170 and 171	144
3.51	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) of <b>170</b> and <b>171</b>	148
3.52	Key <sup>1</sup> H- <sup>1</sup> H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations <b>170</b> and <b>171</b>	149
3.53	Structures of isolated compounds from P. microcarpa	150
3.54	<sup>1</sup> H NMR (600 MHz, DMSO-d <sub>6</sub> ) and key HMBC correlations of <b>103</b>	151
3.55	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of $172$	153
3.56	HSQC-DEPT (600 MHz, CD <sub>3</sub> OD) of <b>172</b>	153

3.57	HMBC (600 MHz, CD <sub>3</sub> OD) of <b>172</b>	154
3.58	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of $173$	155
3.59	<sup>1</sup> H NMR (600 MHz, DMSO-d <sub>6</sub> ) and key HMBC correlations of <b>174</b>	156
3.60	Isolated compounds from the fruits of Z. leprieurii	159
3.61	ESI-MS spectrum of 175	160
3.62	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> ) of <b>175</b>	161
3.63	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>175</b>	161
3.64	HMBC (300 MHz, CDCl <sub>3</sub> ) of <b>175</b>	162
3.65	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> ) of <b>176</b>	163
3.66	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>176</b>	163
3.67	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> + drops MeOH) of <b>177</b>	166
3.68	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> + drops MeOH) and key HMBC	166
	correlations of <b>177</b>	
3.69	<sup>1</sup> H NMR (300 MHz, CDCl <sub>3</sub> ) of <b>178</b>	167
3.70	<sup>13</sup> C NMR (75 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>178</b>	167
3.71	<sup>1</sup> H NMR (300 MHz, CD <sub>3</sub> OD) of <b>179</b>	172
3.72	$^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>180</b>	172
3.73	HSQC-DEPT (600 MHz, CD <sub>3</sub> OD) of <b>180</b>	173
3.74	HMBC (300 MHz, CD <sub>3</sub> OD) of <b>180</b>	173
3.75	<sup>1</sup> H NMR (600 MHz, CD3OD) of <b>182</b>	174
3.76	$^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>182</b>	175
3.77	Structures of isolated compounds from the fruits of Z. zanthoxyloides	176
3.78	ESI-MS spectrum of <b>55</b>	177
3.79	<sup>1</sup> H NMR (300 MHz, CD <sub>3</sub> OD) of <b>55</b>	178
3.80	$^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>55</b>	178
3.81	<sup>1</sup> H NMR (300 MHz, CD <sub>3</sub> OD) of <b>183</b>	180
3.82	$^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>183</b>	180
3.83	<sup>1</sup> H NMR (600 MHz, acetone-d <sub>6</sub> ) of <b>184</b>	181
3.84	$^{13}$ C NMR (150 MHz, acetone-d <sub>6</sub> ) and key HMBC correlations of <b>184</b>	182
3.85	<sup>1</sup> H NMR (300 MHz, CD <sub>3</sub> OD) of <b>185</b>	183
3.86	$^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>185</b>	183
3.87	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) and key HMBC correlations of <b>128</b>	184
3.88	<sup>1</sup> H NMR (600 MHz, CDCl <sub>3</sub> ) and key HMBC correlations of <b>188</b>	185
3.89	ESI-MS spectrum of 188	189

3.90	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) of <b>188</b>	190
3.91	$^{13}\text{C}$ NMR (150 MHz, CD <sub>3</sub> OD), COSY (–) and Key HMBC (–)	190
	correlation of <b>188</b>	
3.92	HSQC-DEPT (600 MHz, CD <sub>3</sub> OD) of <b>188</b>	191
3.93	<sup>1</sup> H- <sup>1</sup> H COSY (600 MHz, CD <sub>3</sub> OD) of <b>188</b>	191
3.94	HMBC (600 MHz, CD <sub>3</sub> OD) of <b>188</b>	192
3.95	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) of <b>189</b>	194
3.96	$^{13}\text{C}$ NMR (150 MHz, CD <sub>3</sub> OD), COSY (–) and Key HMBC (–)	194
	correlation of <b>189</b>	
3.97	<sup>1</sup> H NMR (600 MHz, CD <sub>3</sub> OD) of <b>190</b>	195
3.98	$^{13}\text{C}$ NMR (150 MHz, CD <sub>3</sub> OD), COSY (–) and Key HMBC (–)	195
	correlation of <b>190</b>	
3.99	ESI-MS spectrum of <b>191</b>	198
	•	170
3.100	<sup>1</sup> H NMR (600 MHz, acetone- $d_6$ ) of <b>191</b>	198
<ul><li>3.100</li><li>3.101</li></ul>	<sup>1</sup> H NMR (600 MHz, acetone- $d_6$ ) of <b>191</b> <sup>13</sup> C NMR (150 MHz, acetone- $d_6$ ) of compound <b>191</b>	
		198
3.101	<sup>13</sup> C NMR (150 MHz, acetone- $d_6$ ) of compound <b>191</b>	198 199
3.101	<ul> <li><sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) of compound <b>191</b></li> <li>Possible biosynthesis pattern toward the quinolines isolated from <i>Z</i>.</li> </ul>	198 199
3.101 3.102	<ul> <li><sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) of compound <b>191</b></li> <li>Possible biosynthesis pattern toward the quinolines isolated from <i>Z</i>.</li> <li><i>zanthoxyloides</i></li> </ul>	198 199 201
3.101 3.102	<ul> <li><sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) of compound <b>191</b></li> <li>Possible biosynthesis pattern toward the quinolines isolated from <i>Z. zanthoxyloides</i></li> <li>Summarised biosynthetic pathway toward abietane, clerodane and</li> </ul>	198 199 201
<ul><li>3.101</li><li>3.102</li><li>3.103</li></ul>	<ul> <li><sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) of compound <b>191</b></li> <li>Possible biosynthesis pattern toward the quinolines isolated from <i>Z</i>.</li> <li><i>zanthoxyloides</i></li> <li>Summarised biosynthetic pathway toward abietane, clerodane and kaurane diterpenoids basic skeletons</li> </ul>	<ol> <li>198</li> <li>199</li> <li>201</li> <li>203</li> </ol>
<ul><li>3.101</li><li>3.102</li><li>3.103</li></ul>	<ul> <li><sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) of compound <b>191</b></li> <li>Possible biosynthesis pattern toward the quinolines isolated from <i>Z</i>.</li> <li><i>zanthoxyloides</i></li> <li>Summarised biosynthetic pathway toward abietane, clerodane and kaurane diterpenoids basic skeletons</li> <li>Summarised biosynthetic pathway toward basic flavonoid skeletons</li> </ul>	<ol> <li>198</li> <li>199</li> <li>201</li> <li>203</li> </ol>

## List of tables

Table	Title	Page
1.1	Examples of cytotoxic compounds isolated from plants	7
1.2	Secondary metabolites isolated from Z. leprieurii	43
1.3	Secondary metabolites isolated from Z. zanthoxyloides	52
2.1	List of medicinal plants studied	63
2.2	Cell lines used in bioassays	64
2.3	List of the different HPLC methods developed	72
2.4	Isolation of compounds from C. oligandrus	74
2.5	Isolation of compounds from J. hypocrateriformis	76
2.6	Isolation of compounds from P. microcarpa	78
2.7	Isolation of compounds from Z. leprieurii	80
2.8	Isolation of compounds from Z. zanthoxyloides	83
3.1	Yield of extraction of the different plants collected	94
3.2	Concentrations of extracts causing no more than 10% toxicity	95
3.3	Concentrations of fractions causing no more than 10% toxicity	98
3.4	Haem polymerisation inhibitory concentration (LC <sub>50</sub> in $\mu$ g/mL)	100
3.5	$^{1}$ H (600 MHz, CDCl <sub>3</sub> ) and $^{13}$ C NMR (150 MHz, CDCl <sub>3</sub> ) data of <b>7</b>	105
3.6	$^{1}$ H (600 MHz, CDCl <sub>3</sub> ) and $^{13}$ C NMR (150 MHz, CDCl <sub>3</sub> ) data of <b>149</b>	108
	and <b>150</b>	
3.7	<sup>1</sup> H NMR data of compounds <b>151-162</b>	128
3.8	<sup>13</sup> C NMR data of compounds <b>151-162</b>	130
3.9	<sup>1</sup> H NMR data of compounds <b>163-168</b>	145
3.10	<sup>13</sup> C NMR data of compounds <b>165</b> -1 <b>70</b>	147
3.11	<sup>1</sup> H and <sup>13</sup> C NMR data of <b>170</b> and <b>171</b>	148
3.12	$^{1}$ H (600 MHz, DMSO-d <sub>6</sub> ) and $^{13}$ C NMR (150 MHz, DMSO-d <sub>6</sub> ) data	151
	of <b>103</b>	
3.13	$^{1}$ H (600 MHz, CD <sub>3</sub> OD) and $^{13}$ C NMR (75 MHz, CD <sub>3</sub> OD) data of <b>172</b>	157
	and <b>173</b>	
3.14	$^{1}$ H (600 MHz, DMSO-d <sub>6</sub> ) and $^{13}$ C NMR (150 MHz, DMSO-d <sub>6</sub> ) data	158
	of <b>174</b>	
3.15	<sup>1</sup> H NMR (CDCl <sub>3</sub> , 300 MHz) and <sup>13</sup> C NMR data (CDCl <sub>3</sub> , 75 MHz) of	168
	175-179	
3.16	$^{1}\text{H}$ (600 MHz, CD <sub>3</sub> OD) and $^{13}\text{C}$ NMR (75 MHz, CD <sub>3</sub> OD) data of <b>180</b>	174

- 3.17 <sup>1</sup>H (600 MHz, CD<sub>3</sub>OD) and <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD) data of **181** 175
- 3.18 <sup>1</sup>H and <sup>13</sup>C NMR data of **55**, **128**, **183-184** and **186-187** 186
- 3.19 <sup>1</sup>H (600 MHz, CD<sub>3</sub>OD) and <sup>13</sup>C (150 MHz, CD<sub>3</sub>OD) NMR Data of 196 zanthoamide G-I (**188-190**)
- 3.20 <sup>1</sup>H (600 MHz, acetone-d<sub>6</sub>) and <sup>13</sup>C NMR data (150 MHz, acetone-d<sub>6</sub>) 199 of **191**
- 3.21 Cell viability effect (LC<sub>50</sub> in μM) of isolated compounds against 205 cancer and non-cancer (PNT2) cells
- 3.22 Least toxic concentration (no more than 10 % cells death) of isolated 208 compounds against AREc32 cells

# **Chapter 1 Introduction**

#### 1.1 Cancer

#### **1.1.1 General overview**

Cancer is one of the leading causes of mortality and morbidity worldwide, ranking second after heart disease (Ferlay *et al.*, 2013). The term cancer relates to 200 or more separate diseases characterized by uncontrolled growth and spread of abnormal cells in the body (Trichopoulos *et al.*, 1996). In 2015, approximately 8.8 million deaths worldwide were caused by cancer (WHO, 2018a), and in the UK, an estimated 359,960 new cases were registered in 2015, and about 163,444 deaths were due to cancer (CR-UK, 2018). Globally, one in six deaths are due to cancer and this number is projected to almost double by the year 2030, if no measures are taken to cure or prevent the disease (WHO, 2018a). Factors such as smoking, obesity, inadequate diet, lack of physical activity and pollution, increase the risk of getting cancer (Grundy *et al.*, 2017). Traditional cancer therapies include radiation, chemotherapy and surgery. However, these treatments have limited efficacy, huge side effects and the vital prognostic is unquestionable when the cancer has reached the stage of metastasis. As most cancer results from lifestyles and environmental factors, cancer prevention could be a better way to reduce the incidence of the disease. Besides, always 'prevention is better that cure'.

#### **1.1.2** Mechanism of cancer development

Cancer is a multiple step process and involves genetic and epigenetic changes. Such changes in just one cell can lead to the development of cancer (Fink, 1979). Carcinogenesis consists of three main stages: initiation, promotion and progression (Figure 1.1) (Murakami *et al.*, 1996). Initiation can be induced by exposure to a carcinogen giving rise to spontaneous changes such as DNA damage in one or more stable

cells, leading to mutations in the genetic code (Counts and Goodman, 1994). If the mutation occurs in a gene, which controls cell growth, that cell may become a precancerous cell (Cohen and Ellwein, 1995). If cell division occurs before repair of the damaged cell takes place, then the lesion will be present on the DNA of the daughter cell and will perpetuate into all the further progeny of that daughter cell. Further proliferations of the damaged cells due to exposure to promoting stimuli will give rise to a population of transformed cells and eventually will develop into a tumour (Cairns, 1981). Further stimulations of initiated cells in addition to mutations and chromosomal aberrations cause a fast increase in the tumour size and lead to increasing heterogeneity of the cell population (Devi, 2004). A tumour can be benign or malignant. Benign tumours are not usually life threatening and do not invade nearby cells or spread to others part of the body the way malignant tumours can. If untreated or unresponsive to treatment, malignant tumours can invade and destroy the surrounding tissues. As the tumour progresses under the effect of accumulation of multiple genetic alterations, some cells can be detached from the site of the primary growth, enter the circulating blood and lymph, and be transported to other organs/tissues. They can penetrate the vessels of the organ or tissues, continue to multiply in their new location and eventually develop into another clinically detectable tumour known as metastases (Yokota, 2000).

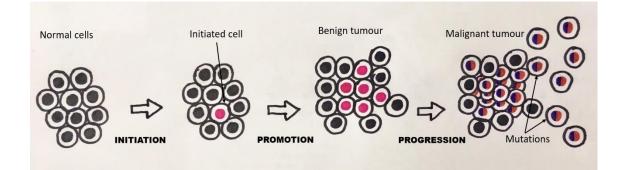


Figure 1.1 Different stages of cancer development

#### **1.1.3** Cancer occurrence and mortality rates

In 2017, the most common causes of cancer death in the world were, lung (1.69 million deaths), liver (788,000 deaths), colorectal (774,000 deaths), stomach (754,000 deaths) and breast (571,000 deaths) (WHO, 2018a). It has been predicted that 42% of males and 39% of females will suffer from cancer at some point in their life. Most common fatal cancers in females affect breast, colon, lung and cervix, while in males they affect lung, prostate, colon and stomach (WHO, 2014). Acute lymphoblastic leukaemia and brain tumours are the most common cancers causing death in children, except in Africa where non-Hodgkin lymphoma occurs more often (WHO, 2014). In the UK, breast, lung, colon and prostate are the most common cancer sites and accounted for more than half of all cancer deaths in 2014 (CR-UK, 2018). A total of 14 million cancer cases were registered in 2012, and this number is expected to rise by about 70% over the next two decades with almost half of all new cases being registered in developing countries (Ferlay et al., 2013). Currently, 70% of cancer deaths worldwide occur in low- and middle-income countries (WHO, 2018a). The chance of survival highly depends on the type of cancer and extent of disease at the beginning of treatment. The 5-year survival rates of many cancer patients are still low, 4% only for cancer affecting pancreas (Pezzuto et al., 2005). Making lifestyle changes such as stopping smoking, adopting a healthy diet and undertaking physical activity can reduce cancer incidence by 40% (Amin et al., 2009).

#### 1.1.4 Cancer chemotherapy

Cancer chemotherapy is the use of cytotoxic drugs to destroy cancer cells. It can be used alone or in combination with surgery and/or radiotherapy. The toxic drugs interfere with DNA and cell division to preferentially kill cancer cells (Cheung-Ong *et al.*, 2013). The

four main categories of cytotoxic drugs include alkylating agents, antimetabolites, hormonal agents and mitotic inhibitors. Alkylating agents derived from mustard gas, their anticancer properties are because of their ability to bind covalently with DNA via their alkyl groups (Lind, 2011). Therefore, DNA molecules are bound tightly together and cannot split during mitosis, thus preventing the proliferation of the cancer cell. Examples of these types of drugs include cisplatin and cyclophosphamide. Antimetabolites have similar structures with the metabolites with whom they interfere with in the organism. They act by blocking DNA dereplication thus preventing cell division. Some examples of antimetabolites are capecitable and fludarable. Hormonal agents are used to stimulate or inhibit the production and activity of some hormones involved in the development of certain cancers. Examples of such are leuprorelin and tamoxifen used in the treatment of prostate and breast cancers, respectively. Mitotic inhibitors are derived from plant natural products. They act by blocking cell division by preventing microtubule function (Rowinsky and Donehower, 1991). Examples of these drugs include etoposide and taxol (Figure 1.2).

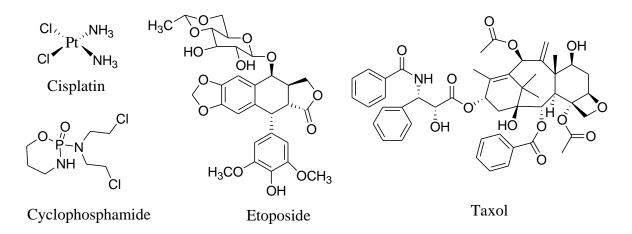


Figure 1.2 Some anticancer dugs

#### **1.1.5** Treatment of cancer by traditional medicine

Traditional medicine is commonly used as both adjuvant and alternative to the use of synthetically manufactured chemotherapeutic drugs. Many surveys have identified plants used in different ethnic groups across the world for the treatment of cancer. An example of such survey could be the one carried out by Hartwell (1971), who combined data of 3000 plants used traditionally for the treatment of cancer. Phytochemical studies of these plants led to the isolation of almost 1000 compounds with potent anticancer properties and the discovery of lead compounds for the development of new cancer drugs (Cragg and Newman, 2016). Table 1.1 lists some examples of isolated anticancer agents from plants.

#### 1.1.6 Cancer chemoprevention

Cancer chemoprevention is defined as the use of vaccines or pharmaceutical agents to inhibit, retard, or reverse the carcinogenesis process (Mettlin, 1997). Since carcinogenesis is a multiple stage process and takes many years or decades to develop, there are several chances to intervene in this process (Tamimi *et al.*, 2002). Moreover, exogenous causative factors including smoking, diet, physical exercise, environmental pollution, exposition to potential carcinogens and radiations, can be controllable and preventable (Grundy *et al.*, 2017). Inhibition of carcinogens, treatment of precancerous lesions, protection of persons at genetic risk and reduction of the incidence of cancer due to dietary epidemiology represent the four main lines address by cancer chemoprevention strategies (Surh, 1999). The ultimate goals of cancer chemoprevention research include identification of the most effective agents and/or to develop efficient approaches for clinical trial and ultimately, application to human population. Plant derived natural products offer a diverse source of potential chemopreventive agents (Manna *et al.*, 2000). Indeed, several compounds isolated from plants i.e. avicins (*Acacia victoriae*), ursolic acid (Basil), resveratrol (red grapes), curcumin (*Curcuma longa*), all from dietary vegetables and fruits, have shown the ability to inhibit human cancer cells growth and therefore, may serve as chemopreventive agents (Shishodia *et al.*, 2003; Aggarwal *et al.*, 2004; Howes, 2015).

Plant sources	Isolated compounds	Activity	References
Acronychia bauer L.	Acronycine	Antitumour	(Duke, 1992)
Brucea species Miller	Bruceantin	Antitumour/	(Shah et al.,
		Antileukaemia	2014)
Bryophyllum pinnatum	Bryophyllin-A	Antitumour	(Duke, 1992)
Salisb.			
Camptotheca acuminata	Camptothecin	Antileukaemia	(Pezzuto,
Decne			1997)
Cassia obtusifolia L.	Emodin	Against breast	(Huang, et al.,
		cancer	2008)
Catharanthus roseus (L.)	Vincristine and	Antileukaemia	(Pezzuto,
G. Don	vinblastine		1997)
Cephalotaxus harringtonii	Homoharringtonine	Antileukaemia	(Shah et al,
(Forbes) K. Koch			2014)
Rabdosia rubescens L.	Rubescensine B	Antihepatoma	(Chang, 1998)
Podophyllum peltatum L.	Etoposide and	Antitumour	(Pezzuto,
	teniposide		1997)
Stephania tetrandra Lour	Tetrandrine	Against lung	(Duke, 1992)
		cancer	
Zanthoxylum nitidum L.	Nitidine chloride	Against lung	(Chang, 1998)
		cancer	

Table 1.1 Examples of cytotoxic compounds isolated from plants

Chemopreventive agents (Figure 1.3) can be natural or synthetic substances. Examples include the synthetic drugs raloxifene and tamoxifen used to prevent breast cancer in women (Chlebowski, 2015) and phenethyl isothiocyanate, a naturally occurring compound in some cruciferous vegetables with potential chemopreventive effects on prostate cancer (Wang, 2010). Chemopreventive agents are classified into two groups, blocking agents and suppressing agents (Wattenberg, 1985). Suppressing agents act in either the promotion or the progression stages during carcinogenesis by inhibiting the proliferation of initiated cells. Blocking agents act on the initiation stage of the carcinogenesis either by blocking the formation of the carcinogen, or by inhibiting interaction of the ultimate carcinogen with crucial cellular macromolecules such as DNA, RNA and protein (Figure 1.4) (Surh, 1999; Kuo et al., 2012).

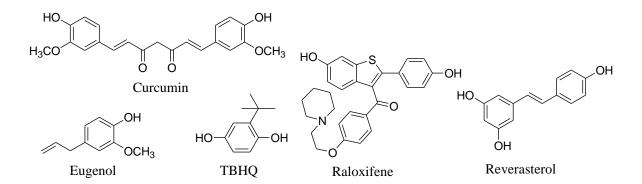


Figure 1.3 Some chemopreventive agents

#### **1.1.7** Chemoprevention by dietary phytochemicals

Dietary phytochemicals are natural products commonly found in edible plants. They became an attraction for cancer chemoprevention research when researchers noticed that people adopting the Mediterranean diet (mean a diet of fish, vegetables, legumes, whole grains, potatoes, fruit and extra virgin olive oil) have the lower risk of developing cancer (Gotsis, 2015). Evidences later confirmed their anticancer and chemopreventive potential

(Shishodia *et al.*, 2003; Aggarwal *et al.*, 2004). Dietary phytochemicals mainly act as blocking agents by intervening in a range of cellular processes resulting in the inhibition of the early stages of the carcinogenesis (Sapienza and Issa, 2016). They prevent carcinogens from reaching targeted sites, support detoxification of highly reactive oxygen species, enhance innate immune surveillance and improve the elimination of initiated cells (Kotecha *et al.*, 2016). The above-mentioned properties of dietary phytochemicals involve the activation of the nuclear factor erythroid-2 (NF-E2)-related factor 2 (Nrf2)-Kelch-like ECH associated protein 1 (Keap1).This complex activation consequently results in the induction of cellular defence mechanisms, including phase II detoxifying enzymes, phase III transporters, antioxidative stress proteins, and other stress-defence molecules that protect normal cells from reactive oxygen species (ROS) and/or reactive nitrogen species (RNS) and reactive metabolites of carcinogenic species known to be the causative agents of more than 600-types of cancer (Lee *et al.*, 2013; Landis-Piwowar *et al.*, 2014).

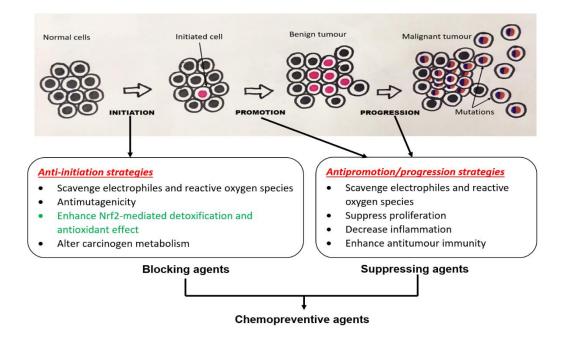


Figure 1.4 Multistage carcinogenesis and strategies for cancer chemoprevention (Adapted from Kuo *et al.*, 2012)

#### 1.1.8 Phase II detoxification enzymes

One significant mechanism by which dietary phytochemicals protect cells from cancer is by the induction of the activity of drug metabolizing enzymes particularly of phase II detoxifying enzymes. Phase II detoxification enzymes provide the defence against foreign chemicals, to which humans are constantly exposed. Many xenobiotics are toxic and if they accumulate in the body, they may cause cell damages which will eventually kill the cell or develop into a cancer. Therefore, many enzymes with various specificities are required to reduce the toxicity of reactive intermediates, protect cells against oxidative or electrophilic challenges, and maintain chemical homeostasis in cells (Kwak *et al.*, 2001). Examples of reactions and Phase II enzymes involved in detoxification include: reduction reaction catalysed by NAD(P)H dehydrogenase, quinone 1 (NQO1); conjugation reactions catalysed by UDP-glucuronosyltransferase (UGT) and glutathione transferase (GST); oxygenation reaction catalysed by heme oxygenase-1 (HO-1) (Zimniak, 2008). The genes encoding the detoxifying enzymes are typically regulated through a consensus cis-enhancer sequences known as the antioxidant responsive element (ARE) or electrophile response element (EpRE), which are in their promoter region. ARE-mediated gene expression plays a central role in the cellular defence against oxidants and electrophiles (Chen and Kong, 2004). It is activated through the binding of the transcription factor, Nrf2, in response to cellular attack by ROS or any other compound with the capacity to either undergo redox cycling or be metabolically transformed to a reactive or electrophilic species (Nguyen et al., 2009). NQO1, for example, is a key enzyme that protects against quinone-derived reactive intermediates and maintains cellular pool of antioxidants such as tocopherol (Nioi and Hayes, 2004). Generally, an increase in the activity of NQO1 enhances the ability of an organism to detoxify numerous potentially harmful xenobiotics (Dinkova-Kostova and Kostov, 2012; Landis-Piwowar et al., 2014). Therefore, substances that increases the activity of NOO1 can be potential chemopreventive agents with the ability to inhibit chemically induced cancer formation (Perez *et al.*, 2010).

#### 1.1.9 Nrf2 as a target for cancer chemoprevention

Nrf2 is a protein that regulates the expression of many phase II detoxifying/antioxidant enzymes. Under normal condition, Nrf2 is sequestered in the cytoplasm by an actinbinding protein, Kelch-like ECH associating protein 1 (Keap-1) and maintains at very low levels regulated by proteasomal degradation (Jaramillo and Zhang, 2013). Upon exposure of cells to inducers such as chemopreventive agents, the sensor cysteines of the Keap-1 that control the Keap-1-dependent Nrf2 degradation is inactivated by the inducers and this therefore, allows *de novo* synthesized Nrf2 to accumulate, to dissociate from the Keap-1 and to translocate to the nucleus where it binds to ARE (Carvalho et al., 2010), and transactivates phase 2 detoxifying and antioxidant genes and transporters, that defend cells from subsequent carcinogenic insults by reducing reactive compounds such as electrophiles and free radicals to less toxic intermediates whilst increasing the ability of the cell to repair any damage ensued (Jeong et al., 2006; Lau et al., 2008; Bryan et al., 2013). Several studies have demonstrated that the major mechanism of protection against carcinogenesis is through Nrf2-mediated induction of drug metabolizing enzyme, particularly phase II detoxification and antioxidant enzymes (Lau et al., 2008). Various compounds, such as naphthoflavone and *tert*-butyl hydroquinone (tBHQ), are potent chemopreventive agents because of their ability to induce phase II enzymes in mammalian cells (Jeong et al., 2008). As blocking agents, dietary phytochemicals protect DNA from mutation by activating the Nuclear factor (erythroid-derived 2)-like 2 (Nrf2) mediation detoxification and antioxidant effect thus making the Nrf2 to be a good drug

target for cancer prevention by phytochemicals (Jaramillo and Zhang, 2013; Reuland *et al.*, 2013; Landis-Piwowar *et al.*, 2014).

#### 1.2 Malaria

#### 1.2.1 General overview

Malaria is a disease endemic to tropical and subtropical regions of the world (Bloland, 2001). The disease is caused by protozoan parasites *Plasmodium*, transmitted to humans through the bites of infected female Anopheles mosquitoes. The most life-threatening and drug-resistant cases are due to infection by the species P. falciparum (Daily, 2006). The development of resistance to drugs poses one of the greatest threats to malaria control and has been linked to recent increases in malaria morbidity and mortality. The World Health Organization's 2016 'World Malaria Report' estimates that in 2016, 216 million cases of malaria were reported causing 445,000 deaths compared to 212 million cases reported in 2015 causing 446,000 deaths (WHO, 2017). Resistance to antimalarial drugs has threatened global malaria control since the emergence of resistance to chloroquine in the 1970s. WHO advocates Artemisinin Combination Therapy (ACT) as first line treatment for acute uncomplicated malaria, where the potentially life threatening parasite P. falciparum is the predominant infecting species with the aim to improve efficacy and to retard the development of resistance (Reyburn, 2010). However, this treatment is also vulnerable and resistance to artemisinin has already been registered in Western Cambodia (Noedl et al., 2008). Because of the potential consequences if resistance to artemisinin were to become widespread, one of the great scientific and medical concerns today is to develop new drugs or vaccines to fight against malaria.

#### 1.2.2 Malaria occurrence and mortality rate

Malaria is one of the oldest diseases that affects human. The evidence of the occurrence of this disease, 5,000 years ago, was found in Chinese manuscript (Cox, 2010). The word malaria derived from the Italian word 'mal aria', which signifies bad air. The disease was first believed to have been caused by vapours given off by swamps by inhalation and was also called swamp fever. It was not until 1880 that the parasite responsible for the disease, known today as *Plasmodium*, was described by Alphonse Laveran (Cox, 2010). The vector responsible for the transmission of the disease, the mosquito *Anopheles*, was later identified by Ronald Ross in 1899 (Cox, 2010). Malaria is widespread in tropical and subtropical regions in sub-Saharan Africa, Central and South America, the Caribbean island of Hispaniola, the Middle East, the Indian subcontinent, South-East Asia, and Oceania (Figure 1.5). Despite its discovery several decades ago, and all the research implemented to prevent and eradicate the disease, malaria is still one of the leading causes of human mortality. Malaria is prevalent in 91 countries and occurs mainly in sub-Saharan Africa.

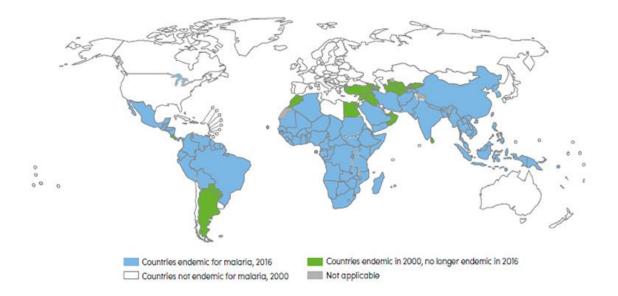


Figure 1.5 Geographic distribution of malaria (Source: WHO. World Malaria Report. Geneva, 2016. Licence: CC BY-NC-SA 3.0 IGO.)

According to the WHO malaria report 2017, an estimated of 216 million cases of malaria were reported in 2016, an increase of 4 million cases compared to 2015. The disease made 445,000 casualties in 2016, a similar number of deaths were observed the previous year (446,000). Sub-Saharan Africa is the most affected area with 90% of malaria cases and 91% of malaria deaths. South-East Asia is the second affected region with 7% and 6% of reported malaria cases and deaths, respectively. Children under 5 years old are the most affected by the disease and 70% of all malaria deaths occur in this age group. Malaria transmission and infection highly depend on the climate conditions and geographical position (Bloland and Organization, 2001). Warm temperatures promote the breeding of the mosquito vector, and highland (altitude >2,000 m) and arid areas (<1000 mm rainfall/year) are less prone to the disease in areas where malaria does occur (Ermert *et al.*, 2011).

#### 1.2.3 Life cycle of malaria *Plasmodium*

The *Plasmodium* is a genus of parasites comprising about 200 species (Chavatte *et al.*, 2007). Five species including *P. falciparum*, *P. vivax*, *P. malariae*, *P. ovale* and *P. knowlesi* are responsible for malaria in humans. Malaria cases associated with the species *P. malariae* and *P. ovale* are less common. *P. knowlesi* mostly affects primates and do not seem to be a major threat to humans (Singh *et al.*, 2004). Infections due to *P. falciparum* and *P. vivax* are the most life threatening. *P. vivax* can persist for years in the dormant stage in the liver and can cause clinical relapses at regular intervals (Carvalho *et al.*, 2010). Almost 99% of malaria cases in sub-Saharan Africa are due to *P. falciparum*, which is the most prevalent in that region. *P. vivax* is predominant in the others area affected by the disease outside Africa and were responsible for 4% of malaria deaths globally in 2016 (WHO, 2017). All the *Plasmodium* species causing malaria in humans

are transmitted by mosquito species of the genus *Anopheles* (Igweh and Okwa, 2012). The life cycle of the malaria parasite, discovered by Ronald Ross in 1899, is complex and takes place in both human and mosquito tissues (Cox, 2010). A simplified schematic representation of the malaria cycle is depicted in Figure 1.6. The cycle starts with the injection of sporozoites by the vector, the mosquito *Anopheles*, into the subcutaneous tissue or directly into blood vessels (Miller *et al.*, 2002). The sporozoites are then transported to the liver via the bloodstream where they infect the liver cells (hepatocytes) (Kappe *et al.*, 2010). Within the liver, sporozoites can invade any cells and then, undergo asexual replication to develop into schizonts which are formed by thousands of merozoites, which are later released into the bloodstream upon rupture of the schizont membrane and infect the red blood cells (RBC) (Cowman and Crabb, 2006; Kappe *et al.*, 2010). The liver stage normally lasts about 9-16 days. For infections due to *P. vivax* and *P. ovale*, sporozoites do not develop into schizonts immediately. They remain dormant in the liver as hypnozoite and that makes it difficult to diagnose and treat (Igweh and Okwa, 2012).

Within the RBC, a fraction of the merozoites initiate the intra-erythrocytic cycle that lasts about 36-48 h. They first develop into a ring-shaped form, then to a trophozoite form and later into schizont. Upon destruction of the RBC, the schizont released new merozoites into the bloodstream, which can infect new RBC and reinitiate the cycle. After several repeated cycles, when 10% of the bloodstream RBC are infected, clinical features associated with the disease can be observed in the patient (Wirth, 2002). The remaining merozoites differentiate into male and female gametocytes. The Mosquito *Anopheles* ingests those gametocytes during its blood meal. Within the mosquito, the gametocytes emerge from the red blood cells as gametes. The fusion of male and female gametes forms a zygote, which elongates into ookinete. The ookinete then later develops into an oocyst whose membrane's rupture will release thousands of sporozoites that will travel to the insect salivary glands. After this stage, the cycle is complete and the mosquito is ready to initiate a new infection on a healthy individual during another blood meal.

Malaria symptoms appear approximately one week after infection. Common symptoms include periodic fever, shivering, cough, respiratory distress, pain in the joints, headache, watery diarrhoea, vomiting and convulsions (Miller *et al.*, 2002). If malaria remains untreated, erythrocytes filled with mature stages of the parasite can adhere to the walls of capillary veins, causing vascular occlusion, severe anaemia, impaired consciousness, renal failure and cerebral death (cerebral malaria) (Miller *et al.*, 2002; Tuteja, 2007).

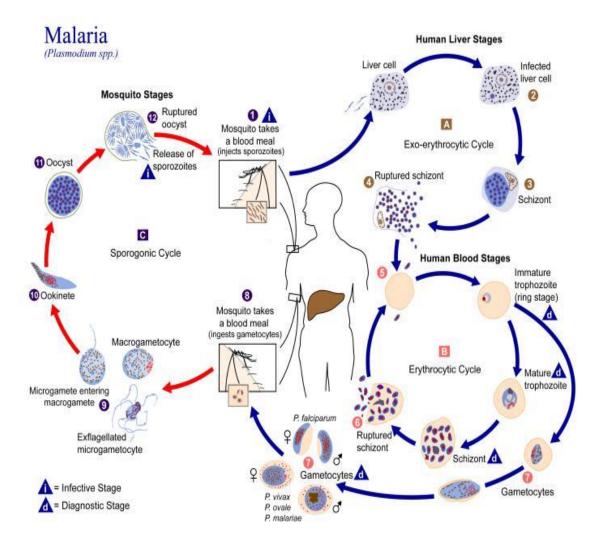


Figure 1.6 Life cycle of malaria cycle infection

(https://hu.wikipedia.org/wiki/Fájl:Plasmodium\_lifecycle\_PHIL\_3405\_lores.jpg )

#### **1.2.4** Current antimalarial drugs and prevention

Malaria is a preventable and treatable disease. Several drugs have been developed to prevent and treat the disease, and several vaccines are currently under development or in clinical trials. The WHO recommends the use of the artemisinin based combined therapy as first line treatment for uncomplicated malaria due to infection by *P. falcifarum* (Prasad *et al.*, 2015). Quinine along with its derivative has been used for the treatment of severe cases of malaria. These therapies have considerably lowered the mortality and morbidity rates due to malaria during the last three decades (Bloland and Organization, 2001).

#### 1.2.4.1 Quinine and its derivatives

Quinine was the first drug to be used to treat malaria. It was serendipitously discovered in the 1600s, when a native from south America suffering from high fever was miraculously cured after drinking from a pool of water contaminated with the *Cinchona* tree (Cragg and Newman, 2005a). Extract of the bark of *Cinchona* with wine or water was used to treat fever associated to malaria until the isolation of quinine as its crystalline sulphate in 1820 by Pierre J. Pelletier and Joseph Caventou. The molecular formula of quinine was proposed by Paul Rabe in 1907, and later confirmed following its total synthesis by Robert B. Woodward and William Doering in 1944 (Achan *et al.*, 2011). Quinine was the main antimalarial drug until the 1920s after which analogues notably chloroquine became more commonly used, due to them being more effective and having less side effects (Plowe, 2010). Quinine is still used today and remains the drug of last resort for the treatment of multidrug-resistant malaria (WHO, 2015). Other quinine derivatives in use today include chloroquine, primaquine, mefloquine and amodiaquine (Figure 1.7) (Bloland and Organization, 2001).

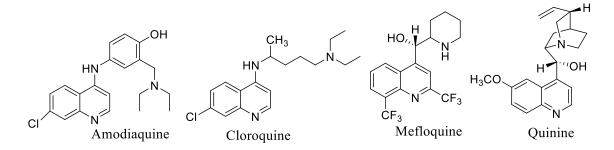


Figure 1.7 Some quinoline antimalarials

#### 1.2.4.2 Antifolates

Folate compounds are derived from folic acid, which is formed from a pteridine ring, *p*-aminobenzoic acid and glutamate. They act on the folate biosynthetic pathway of the *Plasmodium*, a pathway essential to malaria parasite survival, either as inhibitors of dihydropteroate synthase (antifolates class I) or as inhibitors of dihydrofolate reductase (antifolates class II), hence the name antifolates (Nzila, 2006). Class I antifolates in use include proguanil, chlorproguanil, pyrimethamine and trimethoprim. Class II antifolates also called sulpha-drugs in useuse include dapsone, sulfamethoxazole and sulfadoxine (Figure 1.8) (Bloland, 2001). Each class when used alone are effective antimalarials, however, parasitological resistance can develop rapidly in that situation (Bloland and Organization, 2001). When used in combination, they produce a synergistic effect on the parasite and can be effective even in the presence of resistance to the individual components (Nzila, 2006). Antifolates are mainly used in sub-Saharan Africa regions where chloroquine resistant *P. falcifarum* is widespread. They have less or no action of *P. vivax* and *P. ovale* (Gregson, 2005).

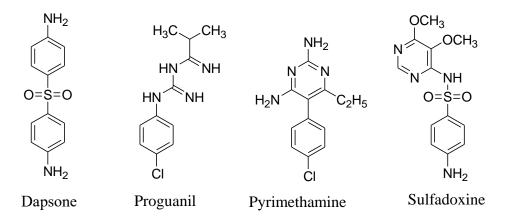


Figure 1.8 Some antifolates used in malaria therapy

#### 1.2.4.3 Artemisinin and related drugs

Artemisinin is a sesquiterpene lactone with antimalarial properties that was first isolated from the leaves of *Artemisia annua* (Tu, 2011). *A. annua* has been used for more than 2000 years in Chinese traditional medicine for the treatment of fever and chills (Tu, 2011). Artemisinin was discovered by Youyou Tu in 1972 as part of an antimalarial research project by the Chinese government. This discovery led her receive the Nobel Prize in Physiology or Medicine in 2015 (Xie, 2016). Artemisinin has later been used as a lead compound for the discovery of related antimalarials including artemether, dihydroartemisinin, artesunate and arteether (Figure 1.9) (Tu, 2017). WHO recommends the use of Artemisinin Combined Therapy (ACT) as first line treatment of uncomplicated malaria due to *P. falciparum* infection (WHO, 2018b). ACT is a combination of artemisinin or an artemisinin-derivative and a second drug with a different mechanism of action to enhance efficacy (Reyburn, 2010). Artemisinin and its derivatives must not be used as oral monotherapy, as this promotes the development of artemisinin resistance (WHO, 2018b).

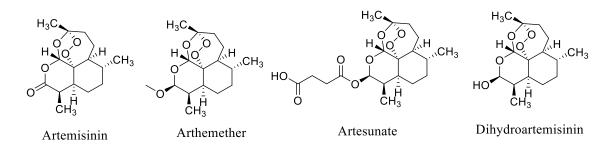


Figure 1.9 Artemisinin and derivatives

## 1.2.4.4 Other used antimalarial drugs

Other antimalarials currently in use include antibiotics, such as tetracycline, and its derivatives, clindamycin and thiostrepton. They are used either as a treatment or for prophylaxis in combination with quinine (Bloland and Organization, 2001). Lumefantrine, atovaquone and halofantrine are also antimalarials (Figure 1.10). They are used in area with *P. falciparum* multidrug resistance.

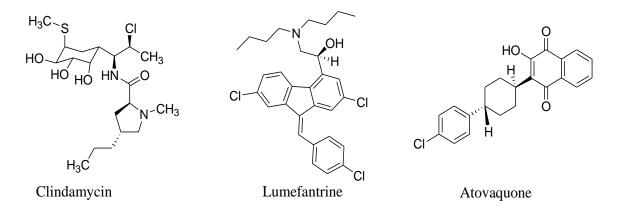


Figure 1.10 Other antimalarials: clindamycin, lumefantrine and atovaquone

#### **1.2.5** Malaria prevention

The WHO has developed several policies to protect persons at risk of getting malaria in endemic regions and those travelling from non-malaria regions to malaria regions. These include the control of the vector and chemoprophylaxis. In endemic regions, persons are protected from the mosquito Anopheles by the usage of long-lasting insecticidal nets (LLIN) and indoor residual spraying insecticides. LLIN are provided free of charge and all people at risk of malaria mainly children under 5 years old and pregnant women are encouraged to sleep under a well maintained LLIN every night. Preventive treatment with sulfadoxine-pyrimethamine are administrated on a trimestral basis to pregnant women and in addition to sulfadoxine-pyrimethamine, amodiaquine is administrated to children under 5 years old monthly. Similarly, for travellers from non-malaria to malaria regions, chemoprophylaxis using drugs that act on the blood stage of the Plasmodium and mosquito repellent lotions are used to prevent malaria. The WHO with partner organisations have also developed some vaccines against malaria (WHO, 2018b). The first of such, RTS, S/AS01 (RTS, S) developed with the partnership of GlaxoSmithKline will be available for use in 2018. RTS, S will be used to prevent children under 3 years old from getting malaria due to infection by P. falciparum. Another vaccine currently under development is Sanaria® PfSPZ Vaccine (Plasmodium falciparum sporozoite Vaccine), a clinical trial of which is currently being conducted in Equatorial Guinea (Olotu *et al.*, 2018).

# 1.2.6 Malaria drug resistance

Even though the number of global malaria deaths has fallen by 48% since 2000, malaria is still a life threatening disease due to the development of resistance to existing drugs in most endemic areas affected and the difficulty to control the most effective malaria vector *Anopheles gambiae* (Prasad *et al.*, 2015). The WHO defines antimalarial drug resistance as the ability of a parasite strain to survive and/or multiply despite the administration and absorption of a drug given in doses equal to or higher than the recommended but within

tolerance of the subject. The parasite must be sensitive to the used antimalarial and this later must be able to gain access to the parasite or the infected RBCs for the duration of the time necessary for its normal action (Bruce-Chwatt, 1986). Despite the long list of available antimalarials, parasites have developed resistance over time to almost all antimalarial drugs. Resistance to quinine was first reported in 1910 after its introduction in pure form in the 1840s, while resistance to artemisinin, introduced in malaria therapy in the 2000s, was recently registered in Cambodia and has since been spread in other areas of the world including south Asia, Serra Leone, Nigeria and Madagascar (Hien *et al.*, 2012). Antimalarial drug resistance has been the main problem in the management of the disease.

#### 1.2.6.1 Resistance to quinoline antimalarials

Resistance of *Plasmodium* to quinine was first reported in 1910 and later confirmed in 1938 (Clyde *et al.*, 1972). Quinine resistant *Plasmodium* strains are rare in Africa but widespread in Southeast Asia and South America (Legrand *et al.*, 2007). Resistance to quinine is usually low grade, with the drug retaining some activity but having its action delayed or diminished (Achan *et al.*, 2011). Chloroquine became the cornerstone of malaria treatment in 1950s. However, just ten years after its introduction, high levels of resistance were recorded in Thailand and South America and have since been found in other countries including Malaysia, Vietnam, Cambodia and Burma and more recently in Sub-Saharan Africa.

#### 1.2.6.2 Resistance to artemisinin and its derivatives

Artemisinin was first used in the 1980s in China. Interest to artemisinin began to grow with the advent of resistance of *Plasmodium* to existing antimalarials. In the 1990s, artemisinin was introduced in malaria therapy in Thailand and Cambodia, and then in other malaria regions (Meshnick, 2002). The first resistance of P. falciparum parasites to artemisinins was detected in Thailand-Cambodia border in the early 2000 (Denis et al., 2006). The resistance was confirmed in 2009 and has since been detected in Southeast Asia (Cambodia, Vietnam, Myanmar and Thailand) and in Africa (Nigeria, Madagascar and Sierra Leone) (Hien *et al.*, 2012). Artemisinin and its derivatives are still effective; however, measures have been taken to reduce the chances of developing resistance in other malaria regions and to prevent the spread of resistance (Reyburn, 2010); the WHO bans monotherapy and recommends the use of artemisinin combined therapies (ACT) for the treatment of uncomplicated malaria. There are five available ACTs including artemether/lumefantrine, artesunate/amodiaquine, artesunate/mefloquine, dihydroartemisinin/piperaquine and artesunate/sulfadoxine-pyrimethamine (Prasad et al., 2015).

#### 1.2.6.3 Resistance to antifolates

Resistance to pyrimethamine and proguanil, the firsts DHFR inhibitors introduce in the malaria therapy, developed shortly after their introduction in the late 1940s and early 1950s respectively in Southeast Asia (Gregson, 2005). Sulpha-drugs such as sulfadoxine and dapsone were introduced in the malaria therapy in the 1960s due to their long half-life and least toxicity (Triglia and Cowman, 1999). However, their usage was almost immediately limited due to the parasite to develop rapid resistance. Antifolates were

reintroduced in form of a combination formulation of DHFR and sulfa-drug (e.g., sulfadoxine-pyrimethamine), when subsequent studies revealed their strong synergistic action and ability to block the plasmodium DNA replication (Abdul-Ghani *et al.*, 2013), however resistance to sulfadoxine-pyrimethamine developed and spread rapidly in South East Asia from the mid-1960s (Nzila, 2006). Because of the affordability of this combination, the WHO still recommends it for the treatment of chloroquine-resistant *P. falciparum* in Africa where the resistance to the combination is still limited to Tanzania (White, 2004; Nzila, 2006).

# **1.2.7** Mechanism of action of chloroquine: hemozoin as target for new antimalarials development

Among existing antimalarial drugs, resistance to quinine has been shown to develop slowly compared to others antimalarials. Quinine, 200 years after its discovery remains one of the most used effective antimalarials. Quinine in combination with an antibiotic e.g., doxycycline, tetracycline or clindamycin is recommended as second line treatment for uncomplicated malaria (when the first-line drug fails or is not available) and remains the first line treatment in combination with clindamycin of malaria in the first trimester of pregnancy (WHO, 2015). Several studies have been carried out to investigate the mechanism of action of quinine and other quinoline antimalarials. Quinoline antimalarials such as quinine and chloroquine, act by interfering with the digestion of haemoglobin in the blood stage of the malaria life cycle (Foley and Tilley, 1997). During the intra-erythrocytic stage, haemoglobin is digested by the parasite *Plasmodium* in its acidic vacuole. Toxic haem, which is dangerous for the parasite, is released and spontaneously converted to a less reactive dimer, hemozoin (Figure 1.11) (Slater, 1993; Egan *et al.*, 1994). Quinoline, inhibits the haem's polymerisation to hemozoin and the consequence

is that the haem released during haemoglobin digestion is accumulated in the parisite's vacuole killing the parasite with its own toxic waste (Slater and Cerami, 1992; Egan *et al.*,1994; Sullivan, 2002). Resistance to quinoline antimalarials is manifested by the capacity of the parasite to expel chloroquine at a rate that does not allow chloroquine to reach levels required for inhibition of haem polymerization (Egan *et al.*, 2001). Hemozoin synthesis is an indispensable pathway for the survival of the parasite and therefore, a good target for the development of new antimalarial drugs. Several screening techniques based on the quinolines antimalarials drugs mechanism have been developed during the recent years to assess the formation of hemozoin *in vitro* and screen compounds as inhibitors of the formation of hemozoin (Tripathi *et al*, 2001; Ncokazi and Egan, 2005).

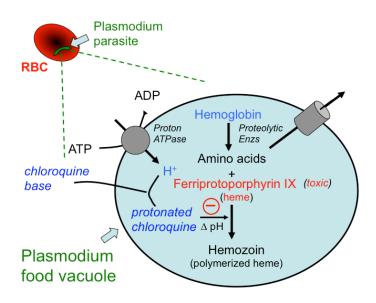


Figure 1.11 Quinoline antimalarials mode of action (from Ivers and Ryan, 2012)

# 1.2.8 Malaria and natural products

The two essential drugs used for the treatment of malaria, quinine and artesunate, are derived from the medicinal plants *Cinchona sp.* and *Artemisia annua* respectively. Plants

have been the source of medicines for humans and animals for centuries. Over a thousandplant species are traditionally used for the treatment and prevention of malaria symptoms. Phytochemical studies of those plants led to the isolation of a range of compounds with potent antimalarial properties. Beaufay *et al.* (2018), Bero *et al.* (2010), and Kaur *et al.* (2009) have published comprehensive reviews (covering up to 2016) on natural occurring compounds from plants with promising antimalarial activities. More recently, the new tirucallane triterpene, *seco*-tiaminic acid B isolated from *Entandrophragma congoënse*, and the macrolide bastimolide B (Figure 1.12) isolated from the tropical marine cyanobacterium *Okeania hirsute* were found to possess antiplasmodial activity with IC<sub>50</sub> values of 2.3 µg/mL and 5.7 µM, respectively (Happi *et al.*, 2018; Shao *et al.*, 2018).

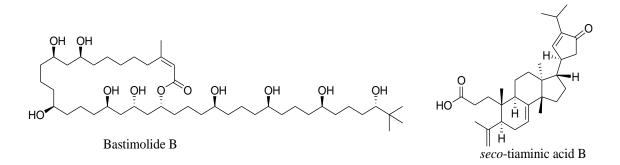


Figure 1.12 Recent antiplasmodial compounds from natural source

# 1.3 Natural products drug discovery

The use of medicinal plants in the treatment of human ailments can be traced for millennia (Solecki, 1975). Plants were used about 2600 B.C. in various forms including infusion, pills, ointments and oils gargles for the treatment of diverse ailments and cosmetics (Cragg and Newman, 2005b). Even today, plants are the almost exclusive source of healthcare for 80% of the world's population and particularly in developing countries (Ekor, 2014). On the international market, global trade of medicinal plants and related

products is estimated at about 40 billion GBP annually (Te, 2013). The knowledge associated with traditional medicine has promoted further investigations of medicinal plants as potential medicines (Dias et al., 2012). Phytochemical screenings show that antibacterial substances like alkaloids, flavonoids, glycosides, saponins and terpenoids are distributed in plant materials (Edeoga et al., 2005). Plant-derived compounds (phytochemicals) represent an interesting source of new bioactive agents (Ghosh et al., 2007). Almost 21% of drugs in use today are derived from natural products and 51% of the 1562 new chemical entities introduced between 1981 and 2014 can be traced to a natural product origin (Newman and Cragg, 2016). The ethnopharmacological properties of plants have been used as a primary foundation for early drug discovery (McRae et al., 2007). This has led to the discovery of the cancer drug Taxol isolated from the bark of Taxus brevifolia and the antimalarial drug quinine isolated from the bark of the Cinchona tree (Wani et al., 1971; Kremsner et al., 1994). Different approaches have so far been developed for drug discovery from natural products (the following will discuss the case of plants only) including random and non-random screenings such as taxonomic, chemotaxonomic, ethnobiological, metabolomic and information managed screening (Cordell et al., 1993; Schwikkard and Mulholland, 2014).

#### 1.3.1 Random screening

Random screening is the arbitrary collection of a huge number of taxa for a large-scale screening (Balunas and Kinghorn, 2005). The plant kingdom comprises about 250,000 plant species on which only around 10% have been screened for their potential bioactivity (Iqbal *et al.*, 2017). This approach is the method of choice in the absence of known taxa with the activity of interest. The collection process involves no intellectualization. All taxa are collected and screened irrespective of their usage. Bioassay-guided isolation is

used to identify the compounds of interest using various chromatographic and analytical techniques. Paclitaxel, sold under the brand name Taxol and used for the treatment of various types of cancer, was first isolated from the bark of *Taxus brevifolia* tree, collected in Washington State as part of a random screening approach carried out by the USA National Cancer Institute in their search for new anticancer drug (Wani *et al.*, 1971; Cragg *et al.*, 1993). Discovery of lead compounds by this approach is completely serendipitous and its success depends on the specificity of the bioassay and the volume of extracts available. In addition, random screening is time consuming, cost effective and only 1-5% of the collected samples screened usually show preliminary bioactive activity (DiMasi *et al.*, 1991). However, with the advent of robotic and high throughput screening techniques the efficiency of random screening has increased, and it is now possible to carry out rapid and sensitive screening of high numbers of samples at low cost (Pereira and Williams, 2007).

# 1.3.2 Ethnobotanical screening

The ethnobotanical or ethnopharmacological screening is essentially based on the empirical usage of plants for therapeutic purposes. The decision to investigate a plant species is determined by the fact that the plant is already being used as medicine or as an insect repellent or for some cultural purpose (Gullo, 1994). The screening of the plant extracts and any isolated compounds is usually guided by what the plant is traditionally used for and any positive results would serve to validate the use of the plant traditionally as well as provide useful leads for further drug development. Several manuscripts describing the usage of plants for medicinal purposes, their dosage and mode of administration in various countries, notably in China, India, Mediterranean and South American countries are available. Ethnobotanical screening has been useful for the

discovery of lead compounds of valuable interest that have undergone chemical modifications to enhance their activity or reduce their toxicity. Fabricant and Farnsworth (2001) have demonstrated that 80% of the 94 medicinal plants from which 122 compounds have been isolated and used as drugs, have had an ethnomedical use identical or related to the current use of the active principle of the plant. In 1785, Withering, an English scientist, published a book on the use of the foxglove, *Digitalis purpurea*, for the treatment of heart disorders; the study of foxglove later led to the isolation of the cardiotonic agent, digitoxin (Fabricant and Farnsworth, 2001). There are many other significant drugs developed from traditional medicinal plants, e.g., quinine and artemisisin.

The main advantage of the ethnobotanical screening is that the active compounds isolated from a plant are likely to show the activity of interest than those derived from plant species with no history of human use. This approach is also valuable for the discovery of complex skeletons unconceivable during synthesis that can then been used as lead compounds for drug discovery. The main inconvenient with this approach is the inadequate design of ethnobotanical data collection, and the misinterpretation of the role medicinal plants play in the medical systems of local and indigenous communities (Etkin, 1993; De Albuquerque and Hanazaki, 2009). In addition, the plant identified may not contain any pharmacologically active compounds with the therapeutic effect observed being due the results of different factors (Moerman, 2007) including interactions with other components of the formulation as it is common for a secondary species of plant or additives such as wine to be added to complete the traditional formulation. Another limitation is the occurrence in the plant of compounds of interest at low concentration or as mixture of analogues making their separation difficult (Gullo, 1994).

#### **1.3.3** Taxonomic and chemotaxonomic screening

Chemotaxonomic screening resides on the ability of plants from the same genus to biologically synthesize the same class of metabolites (Gullo, 1994). This approach then involves the study of species within a plant family or genera which are known to produce biologically active compounds such as alkaloids or flavonoids (Cordell *et al.*, 1993). An example of drug, discovered using such approach, is digitoxin isolated from *Digitalis lanata*, a species from the same genus as *Digitalis purpurea* known to produce the cardiac glycoside digoxin (Farnsworth *et al.*, 1985). This approach is useful for taxonomic study to characterize and classify plants in a family by generating a profile of their secondary metabolites. Compounds obtained are similar in structure and can be used for structure activity relationship studies (Gullo, 1994). The limitation of this approach is the variation of plant metabolites within the period of the year, site of collection and the absence of metabolites of interest due to mutations or genetic factors affecting the gene clusters responsible for their accumulation in the plant (Makins *et al.*, 1983).

# **1.3.4** Virtual screening

Virtual Screening, which is also known as the 'information-managed approach' uses all available biological information and chemical data available in journals and databases to direct the search for new drugs (Cordell *et al*, 1993; Sarker and Nahar, 2018). This approach is useful in identifying structures that are more likely to bind to the target of interest (Rollinger *et al.*, 2008). In addition to the panoply of articles published daily, there are several databases containing extensive libraries of compound properties as well as software programs available to simulate or predict the binding capability of a compound to the receptor or target of interest. NAPRALERT (Natural Product ALERT)

is the first of such databases, created and developed by Farnsworth and coworkers (Loub *et al.*, 1985) which contains over 140,000 compounds isolated from natural sources. Other examples are the Combined Dictionary of Natural Products, Chemlab, ChEMBL and the Collaborative Drug Discovery (CDD) database. Such virtual screening approach is useful to identify active pharmacophores which may be optimised by chemical modification. The main benefit of this approach is to prevent the re-study of known compounds that have already been investigated. By dereplication strategies, databases are linked to hyphenated chromatographic methods e.g. CE-MS, GC-MS, HPLC-DAD-ESI-MS/MS, which allows the rapid identification of new chemical entities from complex matrices such as plant extracts. (Dinan, 2006; Sarker and Nahar, 2012). This virtual screening approach is very productive as evidenced by its use in the pharmaceutical industry (Cheng *et al.*, 2012). However, it is expensive, time consuming and such databases must be highly collaborative and updated on a regular basis.

#### **1.4** Cameroonian medicinal plants and their potential

Cameroon's population rely on medicinal plants for their primary healthcare. Indeed, due to the high cost of modern drugs, only 3 out of 20 patients can afford prescription drugs (Nkongmeneck *et al.*, 2007; Kuete and Efferth, 2010). Cameroonian medicinal plants have exhibited a range of valuable activities including antimicrobial, antimalarial, antiproliferative, anti-inflammatory, antidiabetic, analgesic and antituberculotic (Kuete and Efferth, 2010). They have also shown to be rich in distinctive bioactive compounds (Kuete and Efferth, 2010; Ntie-Kang *et al.*, 2013) such as michellamine B, an alkaloid isolated from the leaves of a Cameroonian native plant *Ancistrocladus korupensis* that has been found to be capable of the complete inhibition of cytopathic effects of HIV-1 and HIV-2 on human lymphoblastoid target cell *in vitro* (Boyd *et al.*, 1994). Cameroonian

flora is rich with an estimated 7,850 known species of which only 37% have been studied (Onana, 2015). Additionally, current studies of Cameroonian medicinal plants continue to show that they are a potential rich source of biologically active compounds for the development of new medicines (Noté *et al.*, 2016; Fankam *et al.*, 2017; Happi *et al.*, 2018).

# 1.5 Selected Cameroonian medicinal plants for this study

Plant materials were selected for this study based on ethnopharmacological information and an extensive literature survey. Plants used traditionally as remedies by local communities in Cameroon in their primary healthcare system for the treatment of malaria or anaemia related diseases were identified. Plants with minimal or no scientific data to validate their use were then selected and collected for further investigation. The following five plants belonging to four families were harvested: *Croton oligandrus* Pierre ex Hutch, *Pseudospondias microcarpa* (A. Rich.) Engl., *Ruspolia hypocrateriformis* (Vahl) Milne-Redh, *Zanthoxylum lepreurii* Guill. and Perr. and *Zanthoxylum zanthoxyloides* (Lam.) Zepern. and Timler.

#### **1.5.1** Croton oligandrus Pierre ex Hutch

Scientific name	Croton oligandrus Pierre ex Hutch
Synonyms	Croton oligandrum
Family	Euphorbiaceae
Subfamily	Crotonoideae
English name	Croton Tribe Crotoneae
Vernacular name	Ebin (Cameroon)

#### **Botanical description**

*Croton oligandrus* Pierre ex Hutch (Figure 1.13) belongs to the Euphorbiaceae family. The Euphorbiaceae, also known as the spurge family, is the largest family of flowering plants of about 322 genera and 8910 (Webster, 1993). *Croton* is a genus with nearly 1300 species distributed in tropical and subtropical regions of the World (Berry *et al.*, 2005). *C. oligandrus*, formally known as *Croton oligandrum*, is a tree of about 9-15 m high commonly found in western and central African secondary forests, especially in Cameroon and Gabon. Its leaves are sub-silvered underneath, oblong-elliptic or elliptic-lanceolate, sub-caudate and rounded at the base. The seeds are oblong, convex on the back, 12 mm long, 7 mm broad, slightly shining and a little wrinkled. The wood is white, and the bark has an ash-grey colour with a remarkable aromatic odour.

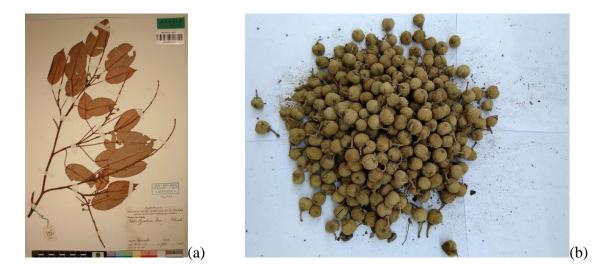


Figure 1.13 a) Ptericarp (scientific tropical archive, http://actd.iict.pt), and b) seeds of *C*. *oligandrus* 

#### Ethnobotanical uses

*Croton oligandrus* is usually harvested in the wild for use as traditional medicine or as timber. In Cameroon, the bark decoction is taken orally to treat pneumonia and splenomegaly as well as for ritual practice (Jiofack *et al.*, 2009; Mpondo *et al.*, 2017). The same bark decoction is used in Gabon for the treatment of anaemia and colic (Betti,

2013). The powder is sniffed to treat nasal tumours and it is externally applied to treat scabies (Schmelzer, 2008).

#### Previous phytochemical investigations

Linalool (1) has been identified as the main constituent of *Croton* oil (Agnaniet *et al.*, 2005). Abega *et al.* (2014) isolated the diterpenes 7-acetoxytrachiloban-18-oic acid (2), crotonadiol (3), crotonoligaketone (4), crotonzambefuran (5) and imbricatadiol (6), and the triterpenes 3-*O*-acetylaleuritolic acid (7), lupeol (8),  $\beta$ -sitosterol (9), and stigmasterol (10) from the stem bark of *C. oligandrus* (Figure 1.14).

#### Previous pharmacological investigations

There is only one report which investigated the pharmacology of *C. oligandrus*. The stem bark's essential oil was screened for its antiradical and antioxidant activities, with no activity being found in both cases (Agnaniet *et al.*, 2005).

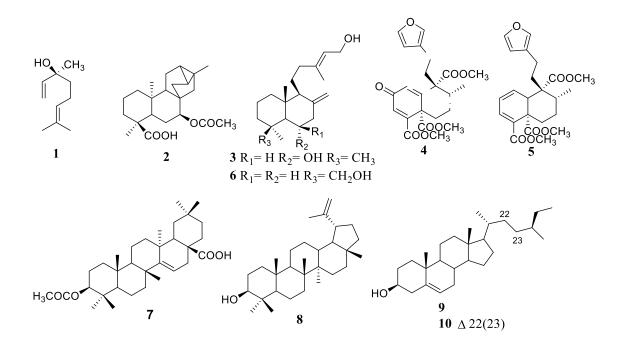


Figure 1.14 Isolated compounds from the bark of C. oligandrus

1.5.2	Ruspolia	hypocraterifor	mis (Vahl)	Milne-Redh
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Scientific name	Ruspolia hypocrateriformis (Vahl) Milne-Redh	
Synonyms	Eranthemum affine Spreng.	
	Eranthemum hypocrateriforme (Vahl) R.Br. ex Roem.and	
	Schult.	
	Justicia hypocrateriformis Vahl	
	Pseuderanthemum hypocrateriforme (Vahl) Radlk.	
	Siphoneranthemum hypocrateriforme (Vahl) Kuntze	
Family	Acanthaceae	
Subfamily	Acanthoideae Tribe Justicieae	
English name	Ruddy Rose, Pricklybush, Cameroun tea, blood tonic plant	
Vernacular name	kifu ke menseh (Cameroon)	

#### Botanical description

The Acanthaceae, acanthus family, is made up of 221 genera and 4000 species (Scotland and Vollesen, 2000). The family, distributed in tropical and subtropical regions of the World with a few species found in temperate regions, includes herbs, shrubs or rarely trees (Burch and Demmy, 1986). The *Ruspolia* comprises five species native to Africa with one species endemic to Madagascar (Milne-Redhead, 1936). *Ruspolia hypocrateriformis* (Vahl) Milne-Redh (synonym: *Justicia hypocrateriformis* Vahl) (Figure 1.15), is a shrub with scattered growth measuring above 1 m in height. The leaves are ovate or elliptic and glabrous when mature. The flowers are borne in showy terminal inflorescences with coral-red tubular flowers 3.75 cm long. (Burch and Demmy, 1986). Its natural habitat includes savanna, and secondary and deciduous forest areas from Senegal to west Cameroon, and it is dispersed to Uganda, Kenya and southern Africa (Milne-Redhead, 1936). In Cameroon, it is seen growing around homes and gardens.



Figure 1.15 Young flourishing plant of R. hypocrateriformis (Orji et al., 2016)

#### Ethnobotanical uses

*Ruspolia hypocrateriformis* is a shrub used as ornamental and medicinal plant in Cameroon. The leaves are soaked or boiled in water, and the obtained juice is drunk to treat diarrhoea and anaemia (Adjanohoun *et al.*, 1996; Noumi, 2015).

# Previous phytochemical investigations

Phytochemical screening has revealed the presence of alkaloids, anthraquinones, flavonoids and saponins from the leaves of the species harvested in Cameroon (Agbor *et al.*, 2014). Alkaloids and anthraquinones were absent from the Nigerian species (Orji *et al.*, 2017). Pyrrolidine alkaloids including hypercratine (**11**), norruspoline (**12**), norruspolinone (**13**) and ruspolinone (**14**) have been isolated from the roots (Figure 1.16) (Roessler *et al.*, 1978; Neukomm *et al.*, 1983).

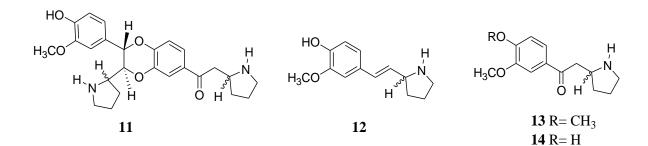


Figure 1.16 Isolated compounds from the roots of R. hypocrateriformis

## Previous pharmacological investigations

Previous pharmacological investigations revealed that *R. hypocrateriformis* aqueous extract possesses antidiarrhoeal activity supported by its antioxidant potential (Agbor *et al.*, 2014) while a supplement of the hydro-ethanol extract daily may help in the management of lead poisoning (Orji *et al.*, 2016).

# 1.5.3 Pseudospondias microcarpa (A. Rich.) Engl.

Scientific name	Pseudospondias microcarpa (A. Rich.) Engl.
Synonyms	Sorindeia obliquifoliolata Engl.
	Spondias angolensis O. Hoffm.
	Spondias microcarpa A. Rich.
Family	Anarcadiaceae
Subfamily	Spondiadoideae Tribe Spondiadeae
English name	African grape
Vernacular name	Gueme, Atom koe mpom (Cameroon)

# Botanical description

The genus *Pseudospondias* belongs to the family Anacardiaceae. There are only two recognised species including *P. microcarpa* Engl. and *P. longifolia* Engl. distributed throughout sub-Sahara tropical Africa (Aubreville, 1950). *P. microcarpa* (A. Rich.) Engl. basionym *Spondias microcarpa* A. Rich. (Figure 1.17), is a spreading tree of up to 20 m high. The trunk is often irregular, twisted, the branches growing near the base and often covered with other plants. The leaves are oblong-ovate to elliptic, odd pinnate on stalks to 30 cm, with 2-8 pairs leaflets plus 1, each leaflet stalked, rather stiff, oval 5-20 cm, base very unequal, tip long pointed, darker above than below, the basal ones smaller. The

fruit are soft drupes measuring about 2.5 cm, red or blue-black when ripe are resinous, the stone inside is 4-sided and contains the seeds (Burkill, 1985).



Figure 1.17 Fruits and ptericarps of *P. microcarpa* (img. By J. Stevens, http://www.zambiaflora.com/speciesdata)

#### Ethnobotanical uses

*Pseudospondias microcarpa* produces edible fruits. The wood is soft and used as firewood, to produce charcoal and canoes (Ruffo *et al.*, 2002). Many medicinal properties have been attributed to *P. microcarpa* across Africa. In Congo, the bark powder is used to treat scabies (Nglobua *et al.*, 2013). The plant is also used for the treatment of malaria, dyspepsia, diarrhoea and opportunistic infections (Mbatchi *et al.*, 2006; Bruno, 2013). The leaves decoction is drunk in Tanzania to relieve chronic cough and malaria (Kisangau *et al.*, 2007). The wood is used as chewing stick in Benin while the bark decoction alleviates teeth problems (Adjanohoun *et al.*, 1996; Akpona *et al.*, 2009). In Sierra Leone, the bark and young leaves decoction is drunk to stimulate appetite (Macfoy, 2013). The plant is used in Ghana as a sedative and for treatment of diseases affecting the central nervous system (Burkill, 1985). The reddish resin contained in the bark is used in Liberia to treat jaundice and other diseases affecting the eyes (Burkill, 1985). The stem bark macerate is mixed with *Coster afer* and palm wine in Cameroon for the treatment of

helminthiasis and constipation (Noumi and Yomi, 2001). Other medicinal uses of the bark and leaves include the treatment of malaria, diabetes, anaemia, rheumatism, gonococci and elephantiasis (Burkill, 1985).

#### Previous phytochemical investigations

The presence of alkaloids, anthraquinones, cardiac glycosides, coumarins, flavonoids, polyphenols, quinones, saponins, tannins, and triterpenes were reported from the stem bark aqueous and MeOH-DCM extracts (Yondo *et al.*, 2009). Alkaloids, tannins, terpenoids, steroids and heteroside cardiotonics have been reported from the leaves (Akpona *et al.*, 2009).

## Previous pharmacological investigations

The ethanolic extracts of the roots and stem bark of the plant harvested in Tanzania exhibited good antiplasmodial activity, with IC<sub>50</sub> of 1.13 µg/mL and 4.33 µg/mL, respectively (Malebo *et al.*, 2010). From the species harvested in Cameroon, the leaves and stem bark hydro-ethanolic extracts and the roots ethyl acetate extract showed moderate antimicrobial and antioxidant properties (Yondo *et al.*, 2009). A moderate antiplasmodial activity against *Plasmodium* K1 strain and cytotoxic activity against HepG2 cell lines were also observed (Sidjui *et al.*, 2016). *P. microcarpa* hydro-ethanolic extract possesses antidepressant-like effect, analgesic and anticonvulsant activities (Adongo *et al.*, 2015). The extract has also showed anxiolytic-like activity similar to that of diazepam (Adongo *et al.*, 2016).

# 1.5.4 Zanthoxylum lepreurii Guill. and Perr.

Scientific name	Zanthoxylum lepreurii Guill. and Perr.	
Synonyms	Fagara angolensis Engl., Fagara attiensis Hutch. and Dalziel	
	Fagara beniensis Engl., Fagara membranifolia Mildbr.	
	Fagara kibomboensis De Wild., Fagara nitens (Hiern) Engl.	
	Fagara leprieurii (Guill. and Perr.) Engl., Fagara olung Engl.	
	Fagara polyacantha Engl., Fagara stuhlmannii (Engl.) Engl.	
	Zanthoxylum crenatum A. Cheval., Zanthoxylum nitens Hiern	
	Zanthoxylum stuhlmannii Engl., Fagara kelekete De Wild.	
Family	Rutaceae	
Subfamily	Toddalioideae <b>Tribe</b> Zanthoxyleae	
English name	Satinwood, sand knobwood	
Vernacular name	Melan, bongo (Cameroon)	



Figure 1.18 Dried Fruits of Z. leprieurii

# Botanical description

The genus *Zanthoxylum* L. belongs to the Rutaceae family and comprises about 200 species of pantropical trees. In Africa, the genus is represented by 35 species of economic importance for their usage as spice, edible fruit, medicinal plants and wood for

construction (Matu, 2011). The major characteristic of *Zanthoxylum* trees is that their trunks, branches, branchlets, leaf stalks and inflorescence axes are covered by prickles or spines (Adesina, 2005). *Zanthoxylum leprieurii* Guill. and Perr. is a medium tree of up to 15-25 m high found in rain forests and savanna woodland with brown branches armed with straight or sometimes curved spines (Tabuti, 2016). The leaves are alternate and impair-pinnate with 8-16 leaflets; leaflets are opposite or almost opposite, elliptical to oblong-elliptical, apex acuminate or caudate with oblique base. The flowers are unisexual, small and nearly sessile. Fruits (Figure 1.18) and seeds are almost globose measuring 4-5 mm and 3-3.5 mm in diameter, respectively. While the fruit are reddish and glandular pitted, the seeds are black and shiny. *Z. leprieurii* has a wide distribution and occurs from Senegal east to Ethiopia and south to Mozambique and eastern South Africa (Tabuti, 2016).

## Ethnobotanical uses

The wood of *Z. leprieurii* is used in house and boat-building, carpentry, decorative and musical instrument construction and in the paper and pulp industry (Adesina, 2005). Different parts of the plant are traditionally used in Africa for medicinal purposes. The leaves are used for the treatment of stomatitis, gingivitis and bilharzia, while the roots are used as anti-ulcerative, antiseptic, urinary antiseptic, antisickling and antibacterial (Ngane *et al.*, 2000). The stem bark is used as antimicrobial, digestive aid, antidiarrhoeic, anticancerous, anti-odontalgic and parasiticide (Ngoumfo *et al.*, 2010). In Ghana, a decoction of the stem bark and root bark is used as a diuretic. The stem bark and leaves are used topically to treat wounds, syphilitic sores and leprous ulcers. The bark, when boiled in hot water produces vapour, which is inhaled to treat toothache and rheumatic pain. Decoction and poultice of stem bark is used for the treatment of skin and urinary tract infections, dysentery and intestinal worm infestation (Adesanya and Sofowora,

1983; Agyare *et al.*, 2009). In Senegal, the powdered bark together with the latex of *Baissea axillaris* (Benth.) Hua is applied to tumours (Tabuti, 2016). In Cameroon, fruit are sold on local markets for their usage as spice (Ngoumfo *et al.*, 2010). The leaves, stems and roots are used in the treatment of gonococci, urinary infections, dysentery, kidney pain and sterility (Noumi and Yomi, 2001). The fruit infusion is used for the treatment of sickle cell anaemia (Tabuti, 2016).

## Previous phytochemical investigations

The phytochemistry of the *Zanthoxylum* genus has been thoroughly investigated in relation to its use for the treatment of sickle cell anaemia. Preliminary phytochemical screening of the stem bark aqueous-methanol extract of *Z. leprieurii* revealed the presence of alkaloids, flavonoids, carbohydrates and saponins (Tatsadjieu *et al.*, 2003). Alkaloids such as aporphine, acridone, benzophenanthridine and quinoline derivatives were found to be the main constituents of the stem bark, pericarps, roots and fruit. Flavonoids, lignans and isobutylamides have also been reported. Table 1.2 summarises the isolated compounds (Figure 1.19) from *Z. leprieurii*.  $\alpha$ -pinene (**62**), (E)- $\beta$ -ocimene (**63**), limonene (**64**) and myrcene (**65**) (Figure 1.19) were found to be the major constituents of the fruit essential oil of *Z. leprieurii* (Fogang *et al.*, 2012).

#### Previous pharmacological investigations

*Zanthoxylum leprieurii* fruit essential oil showed antimicrobial and antibacterial activity, inhibiting the growth of several microorganisms and bacteria strains (Tatsadjieu *et al.*, 2003). The oil also possesses antioxidant, anti-inflammatory, antidermatophitic and moderate cytotoxic properties (Misra *et al.*, 2013; Tchabong *et al.*, 2017). The aqueous-methanol stem bark extract was found to have wound healing properties as well as antimicrobial, cytotoxic and antioxidant activities (Kuete *et al.*, 2011; Agyare *et al.*, 2014). The aqueous

ethanolic extracts of the leaves, roots and stem bark have demonstrated moderate antifungal activity while the chloroform extract of the fruit showed moderate toxicity with the brine-shrimp assay (Ngane et al, 2000; Ngoumfo et al, 2010). Acridone alkaloids, 1hydroxy-3-methoxy-N-methylacridone (16), arborinine (13), 1,3-dihydroxy-2-methoxy-N-methylacridone (19) and tegerrardin A (26) isolated from the fruit of Z. leprieurii were found to be moderately active against lung carcinoma cells (A549), colorectal adenocarcinoma cells (DLD-1) and normal cells (WS1) with IC<sub>50</sub> values ranging from 27 to 77 µM (Ngoumfo et al., 2010; Kuete et al., 2011). Nitidine (45) demonstrated strong antileukemic activities against L1210 and P388 and showed growth inhibition of Lewis lung carcinoma (Wall et al., 1987). 2-Hydroxy-1,3-dimethoxy-N-methyl-9-acridone (25) and 3-hydroxy-1,5,6-trimethoxyacridone (22) showed good activity against pan-sensitive and resistant strains of *Mycobacterium tuberculosis* (Bunalema et al., 2017). Arborinine (23) and xanthoxoline (27) demonstrated good in vitro antiplasmodial activity against *Plasmodium falciparum* 3D7 strains with IC<sub>50</sub> values of  $4.5\pm1.0$  and  $4.6\pm0.6$  µg/mL, respectively (Tchinda et al., 2009). The same study reported arborinine (23) and tegerrardin A (26) as good chelating agents with 90% and 61% radical scavenging activity, respectively (Tchinda et al., 2009).

Isolated compounds	Plant parts	References
Acridone alkaloids		
1-Hydroxy-3,4-dimethoxy-N-	Fruits, roots	(Fish and Waterman, 1972a;
methylacridone (15)		Tchinda <i>et al.</i> , 2009)
1-Hydroxy-3-methoxy-N-	Fruits, roots	(Adesina, 2005; Ngoumfo et
methylacridone (16)	and stem	al., 2010; Bunalema et al.,
	bark	2017)

Table 1.2 Secondary metabolites isolated from Z. leprieurii

# Table 1.2 continued

1-Hydroxy- <i>N</i> -methylacridone ( <b>17</b> )	Fruits, roots	(Adesina, 2005)
	and stem	(1.000,000)
	bark	
1,2-Dihydroxy-	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
3-methoxy- <i>N</i> -methylacridone (18)	roots	
1,3-Dihydroxy-2-methoxy- <i>N</i> -	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
methylacridone (19)	roots	
1,3-Dihydroxy-4-methoxy- <i>N</i> -	Roots	(Fish and Waterman, 1972b)
methylacridone (20)		
2-Hydroxy-1,3-dimethoxy-N-methyl-	Stem bark	(Bunalema <i>et al.</i> , 2017)
acridone (21)		
3-Hydroxy-1,5,6-trimethoxy-acridone	Fruits and	(Wouatsa <i>et al.</i> , 2013a;
(22)	stem bark	Bunalema et al., 2017)
Arborinine (23)	Fruits and	(Tchinda et al., 2009;
	roots	Ngoumfo et al., 2010)
Helebelicine A (24)	Fruits	(Ngoumfo <i>et al.</i> , 2010)
Helebelicine B (25)	Fruits	(Ngoumfo et al., 2010)
Tegerrardin A (26)	Fruits and	(Tchinda et al., 2009;
	roots	Ngoumfo <i>et al.</i> , 2010)
Xanthoxoline (27)	Fruits, roots	(Adesina, 2005; Tchinda et
	and stem	al., 2009; Ngoumfo et al.,
	bark	2010)
Alkamides		
(2E,4E)-N-Isobutyleicosa-deca-2,4-	Roots	(Adesina, 2005)
dienamide (28)		
(2E,4E,8E,10E,12E)-N-Isobutyl-	Roots	(Adesina, 2005)
2,4,8,10,12-tetradecapentaenamide ( <b>29</b> )		
<i>N</i> -Isopentyl-cinnamamide ( <b>30</b> )	Roots	(Adesina, 2005)
<i>N</i> -Docosanoyltyramine ( <b>31</b> )	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
	roots	
Violyedoenamide (32)	Fruits and	(Ngoumfo et al., 2010)
	roots	

# Table 1.2 continued

Aporphine alkaloids		
Magnoflorine (33)	Roots and	Adesina, 2005
	stem bark	
<i>N</i> -Methylcorydine ( <b>34</b> )	Roots	Adesina, 2005
Tembetarine (35)	Roots and	(Fish and Waterman, 1972b)
	stem bark	
Benzophenanthridine alkaloids		
10-O-Demethyl-12-O-	Roots	(Ngoumfo et al., 2010)
methylarnottianamide (36)		
10-O-Demethyl-12-O-	Roots	(Ngoumfo <i>et al.</i> , 2010)
methylisoarnottianamide (37)		
Angoline (38)	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
	roots	
Arnottianamide (39)	Roots	(Adesina, 2005)
Chelerythrine (40)	Roots and	(Adesina, 2005)
	stem bark	
Decarine (41)	Fruits and	(Ngoumfo et al., 2010)
	roots	
Dihydroavicine (42)	Roots	(Adesina, 2005)
Dihydronitidine (43)	Fruits and	(Ngoumfo et al., 2010)
	roots	
Fagaronine (44)	Roots	(Adesina, 2005)
Nitidine (45)	Roots and	(Adesina, 2005; Ngoumfo et
	stem bark	al., 2010)
Norchelerythrine (46)	Roots	(Adesina, 2005)
Coumarins		
6,7-Dimethylcoumarin (47)	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
	roots	
6,7,8-Trimethoxycoumarin ( <b>48</b> )	Bark	(Eshiet and Taylor, 1968)
7, 8-Dimethoxycoumarin (49)	Fruits	(Misra <i>et al.</i> , 2013)

# Table 1.2 continued

Scoparone (50)	Roots and	(Adesina, 2005; Tchinda et
	stem bark	al., 2009)
Xanthotoxin (51)	Roots and	Adesina, 2005
	stem bark	
Lignans		
	Roots	(Adaging 2005)
Asarinin (52)		(Adesina, 2005)
Lirioresinol-B-dimethyl ether (53)	Roots	(Adesina, 2005)
Sesamin (54)	Fruits, roots	(Adesina, 2005; Ngoumfo et
	and stem	al., 2010)
	bark	
Quinoline alkaloids		
Skimmianine (55)	Roots	Adesina, 2005
Triterpenes		
β-Sitosterol (9)	Fruits	(Ngoumfo et al., 2010)
β-Sitosterol palmitate (56)	Bark	(Eshiet and Taylor, 1968)
Lupeol (8)	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
	roots	
Other compounds		
Candicine (57)	Stem bark	(Fish and Waterman, 1972a)
	and roots	
Glucose (58)	Fruits and	(Ngoumfo <i>et al.</i> , 2010)
	roots	
Hesperidin (59)	Fruits and	(Ngoumfo et al., 2010)
	roots	
Sacchore (60)	Fruits and	(Ngoumfo et al., 2010)
	roots	
Sinapic acid (61)	Fruits	(Misra <i>et al.</i> , 2013)

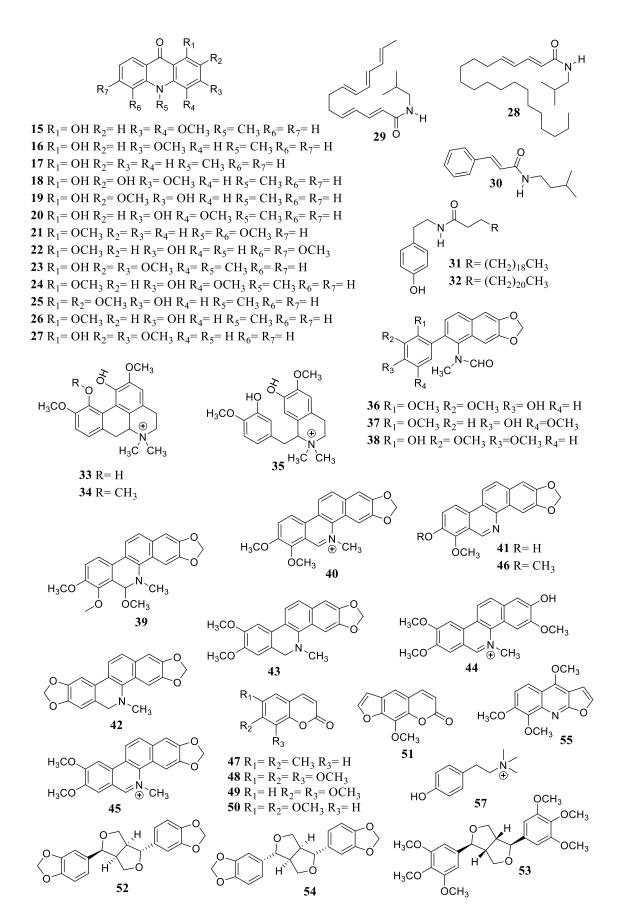


Figure 1.19 Isolated compounds from Z. leprieurii

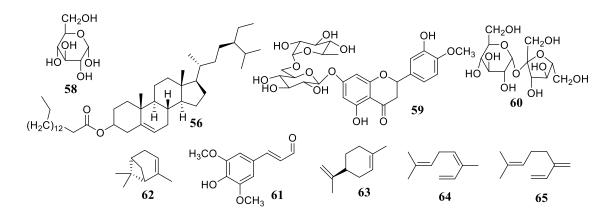


Figure 1.19 continued

# 1.5.5 Zanthoxylum zanthoxyloides (Lam.) Zepern. and Timler

Scientific name	Zanthoxylum zanthoxyloides (Lam.) Zepern. and Timler
Synonyms	Fagara senegalensis (DC.) A. Chev.
	Fagara zanthoxyloides Lam.
	Zanthoxylum senegalense DC
	Zanthoxylum polyganum Schum.
Family	Rutaceae
Subfamily	Toddalioideae Tribe Zanthoxyleae
English name	Candlewood
Vernacular name	Nwah che (Cameroon)

# Botanical description

*Zanthoxylum zanthoxyloides* is a small tree of up to 6-12 m tall widespread in Africa savanna and dry forest vegetations (Matu, 2011). The trunk is spiny with straight, often short bole and rounded and a quite dense crown. The bark is grey to beige, rough, with fine vertical fissures, often with woody prickle-bearing protuberances; the slashes are yellow, odorous, orange-mottled beneath while the stems are glabrous, grey, with solitary prickles. The leaves are alternate, glabrous, impari-pinnate with 5-7(-11) opposite or

alternate leaflets, up to 12(-20) cm long; leaflets are obovate to elliptical, 5-10 cm x 2-4 cm, with cuneate to rounded base, obtuse or rounded apex, sometimes apiculate or notched, with many glandular dots, smelling of pepper and lemon when crushed, rigidly papery, pinnately veined with 10-14 pairs of lateral veins, barely prominent, fusing near the margin. The flowers are unisexual, regular, white or greenish, sessile; corolla barely open; Male flowers pentamerous. Petals imbricate. Rudiment of pistil subulate, upon a thickened central disk. The fruits (Figure 1.20) are rather crowded, in paniculate clusters shorter than the leaves, 1-carpellary, 1-seeded. Seeds are shining, blue-black; fruits contain an ovoid follicle, 5-6 mm in diameter, brown, with glandular dots, dehiscent, 1-seeded (Burkill, 1985).



Figure 1.20 Dried Fruits of Z. zanthoxyloides

#### Ethnobotanical uses

Zanthoxylum zanthoxyloides as the other African Zanthoxylum species has a broad spectrum of ethnopharmacological usage across Africa. In Ouganda, the root-bark extract is used in treating elephantiasis, toothache, impotence, gonorrhoea, malaria, dysmenorrhoea and abdominal pain (Adjanohoun *et al.*, 1993). In Senegal, the plant is used in the treatment of enteritis, dysentery, diarrhoea, guinea worm, urethritis and as an

anti-odontalgic. In Cameroon, the leaves, stems and roots of these plants are used in the treatment of gonococci, urinary infections and dysentery (Noumi and Yomi, 2001). The seeds are used as spices. Leaves and bark are used for the treatment of cough, fever, colds, toothache and snake bites. The leaves are also used as scaring, antiseptic, astringent and laxative, while the stem bark is used as antirheumatic, anti-odontalgic, diuretic, urinary antiseptic, digestive aid and parasticide. The roots are used as antiseptic, antisickler, digestive aid and parasticide (Ngane *et al.*, 2000; Kassim *et al.*, 2004). In Ivory Coast, sap from the pulped bark is applied as eye drops to treat eye infections. In Ghana, roots and stem bark powder is taken to treat whooping cough (Arbonnier, 2004). In Nigeria, stem bark and roots decoction are drunk to treat cancer. Roots and stem bark of *Z. zanthoxyloides* are also used as chewing sticks for teeth cleaning (Arbonnier, 2004).

#### Previous phytochemical investigations

Zanthoxylum zanthoxyloides is a rich source of bioactive metabolites including alkaloids, aliphatic and aromatic amides, lignans, coumarins, sterols and carbohydrate residues (Figure 1.21). A list of isolated compounds from different parts of the plant as well as their activities is presented in Table 1.3. The main constituents of the fruit essential oils were found to be  $\alpha$ -pinene (62), (*E*)- $\beta$ -ocimene (63), citronellal (143), citronellol (144), citronellyl acetate (145), geraniol (146), limonene (64) and myrcene (65) (Figure 1.21) (Tatsadjieu *et al.*, 2003; Fogang *et al.*, 2012).

#### Previous pharmacological investigations

*Zanthoxylum zanthoxyloides* has demonstrated a panel of biological activities. The plant powder and extract have good insecticidal and antifeedant activities against a range of crop pests (Ojo *et al.*, 2016). The fruit essential oil possesses antimicrobial, antibacterial, insecticidal and antioxidant properties (Tatsadjieu *et al.*, 2003; Fogang *et al.*, 2012).

Roots and stem bark methanolic extracts were found to be least toxic with LD<sub>50</sub> 5.0 g/Kg and 1.5 mg/kg body weight, respectively in Wistar rats and the ethanolic extracts have moderate antifungal activity (Ngane *et al.*, 2000; Misra *et al.*, 2013). The crude alkaloid extracts obtained from the trunk bark demonstrated good antimalarial activity with IC<sub>50</sub> ranging from 1.91 to 4.32 µg/mL (Gansane *et al.*, 2010). The DCM extract of the stem and roots bark and the methanol extract of the stem bark have also been found to possess good antiplasmodial activity with IC<sub>50</sub> ranging from 1 and 10 µg/mL. Extract of the root bark has also demonstrated significant anti-inflammatory, anti-sickling and analgesic activities and moderate antibacterial, antiprotozoal and antiviral activities against a range of pathogenic bacteria, *Herpes simplex* virus and *Leishmania major* (Kassim *et al.*, 2005; Ahua *et al.*, 2007; Barnabas *et al.*, 2011). Extract of the root bark is also reported to have gastroprotective activity (Boye *et al.*, 2012). Ethanolic extract of the leaves has shown antifungal, anthelmintic, antidiabetic, hypolipidaemic and antioxidant activities (Aloke *et al.*, 2012; Adekunle *et al.*, 2014). A moderate cytotoxic activity of the fruits ethanolic extract has also been reported (Choumessi *et al.*, 2012).

The acridones 3,4,5,7-tetrahydroxy-1-methoxy-*N*-methylacridone (**67**) has been found to possess a moderate antibacterial activity against *Micrococcus luteus* and *Pseudomonas aeruginosa* and 3-hydroxy-1,5,6-trimethoxy-acridone (**22**) showed moderate cytotoxic effect against liver cancer cell line WRL-68 (Wouatsa *et al.*, 2013a). Furoquinoline alkaloid atanine (**128**) possessed antiprostaglandin synthetase activity (Prempeh and Mensah-Attipoe, 2009). Fagaronine (**44**), 2-hydroxymethylbenzoic acid (**109**), burkinabins A, B and C (**111-113**) have demonstrated strong antileukaemic acid justifying of the usage of *Z. zanthoxyloides* plant in the traditional treatment of sickle cell anaemia (Adesina, 2005; Ouattara *et al.*, 2009). Chelerythrine (**40**), berberine (**90**) and 6-canthinone (**136**) possessed antimicrobial activity (Adesina, 2005).

# Table 1.3 Secondary metabolites isolated from Z. zanthoxyloides

Isolated/identified compounds	Plant parts	References		
Acridone alkaloids				
1-Hydroxy-3-methoxy-N-methylacridone	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
(16)				
1,6-Dihydroxy-3-methoxy-acridone (66)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
3-Hydroxy-1,5,6-trimethoxy-acridone (22)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
3,4,5,7-Tetrahydroxy-1-methoxy-10-	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
methyl-9-acridone (67)				
4-Hydroxyzanthacridone (68)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
4-Hydroxyzanthacridon-(2,4')-oxide (69)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
4-Methoxyzanthacridone (70)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
Helebelicine A (24)	Fruits	(Wouatsa <i>et al.</i> , 2013a)		
Alkamides				
(2E,4E)-N-Isobutyldeca-2,4-dienamide	Roots	(Chaaib <i>et al.</i> , 2003)		
(71)				
(2E,4E)-N-Isobutylhexadeca-2,4-	Fruits, roots	(Adesina, 2005)		
dienamide (72)	and bark			
(2 <i>E</i> ,4 <i>E</i> )- <i>N</i> -isobutylocta-2,4-dienamide ( <b>73</b> )	Roots	(Chaaib <i>et al.</i> , 2003)		
(2E,4E)-N-Isobutyltetradeca-2,4-	Fruits	(Adesina, 2005)		
dienamide (74)				
(2E,4E,8E,10E,12E)-N-Isobutyl-	Roots	(Adesina, 2005)		
2,4,8,10,12-tetradecapentaenamide (29)				
Cis-fagaramide (75)	Roots and	(Chaaib et al., 2003;		
	stem bark	Adesina, 2005)		
<i>N</i> -Isopentylcinnamamide ( <b>30</b> )	Roots	(Adesina, 2005		
Pellitorine (76)	Roots	(Adesina, 2005)		
Piperlonguminine (77)	Roots	(Adesina, 2005)		
Rubebmamin (78)	Roots and	(Adesina, 2005;		
	stem bark	Oriowo, 1982)		
Trans-fagaramide (79)	Roots	(Chaaib et al., 2003;		
		Queiroz et al., 2006)		

# Table 1.3 continued

Aporphine alkaloids			
Berberine (80)	Roots	(Adesina, 2005)	
Magnoflorine (33)	Roots	(Adesina, 2005)	
<i>N</i> -Methylcorydine ( <b>34</b> )	Roots	(Adesina, 2005)	
<i>N</i> , <i>N</i> -Dimetylindicarpine ( <b>81</b> )	Roots	(Queiroz <i>et al.</i> , 2006)	
Tembetarine (35)	Roots	(Adesina, 2005)	
Benzophenanthridine alkaloids			
6-(2-Oxobutyl)-dihydrochelerythrine (82)	Roots	(Chaaib <i>et al.</i> , 2003)	
6-Acetonyldihydrochelerythrine (83)	Roots	(Chaaib <i>et al.</i> , 2003)	
6-Ethoxychelerythrine (84)	Roots bark	Torto et al., 1966	
6-Hydroxydihydrochelerythrine (85)	Roots	(Wangensteen et al.,	
		2017)	
Bis-Dihydrochelerythrinyl ether (86)	Roots	(Adesina, 2005)	
Arnottianamide (38)	Roots	(Adesina, 2005)	
Buesgenine (87)	Roots	(Wangensteen et al.,	
		2017)	
Chelerythrine (40)	Roots	(Torto <i>et al.</i> , 1966)	
Dihydroavicine (42)	Roots	(Adesina, 2005)	
Dihydrochelerythrine (88)	Stem bark	(Torto <i>et al.</i> , 1973)	
Fagaridine (89)	Stem bark	(Torto <i>et al.</i> , 1973)	
Fagaronine (34)	Roots and	(Queiroz et al., 2006)	
	stem bark		
Norchelerythrine (36)	Roots	(Queiroz <i>et al.</i> , 2006)	
Oxychelerythrine (90)	Roots	(Adesina, 2005)	
Coumarins			
4-Methoxycoumarin (91)	Fruits	(Tine <i>et al.</i> , 2017)	
6-Methylcoumarin (92)	Fruits	(Tine <i>et al.</i> , 2017)	
6,7-Dimethylesculetin (47)	Fruits	(Tine <i>et al.</i> , 2017)	
6,7,8-Trimethoxycoumarin (48)	Roots	(Wangensteen et al.,	
		2017)	

# Table 1.3 continued

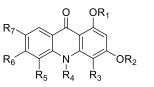
7-Methylcoumarin (93)	Fruits	(Tine <i>et al.</i> , 2017	
Bergapten (94)	Fruits and stem bark	(Adesina, 2005)	
Daphnetin-7-methylether (95)	Fruits and stem bark	(Tine <i>et al.</i> , 2017)	
Herniarin (96)	Fruits and stem bark	(Tine <i>et al.</i> , 2017)	
Isobergapten (97)	Fruits	(Tine <i>et al.</i> , 2017)	
Isopimpinellin (98)	Fruits and leaves	(Tine et al., 2017)	
Imperatorin (99)	Stem bark	(Adesina, 2005)	
Marmesin (100)	Fruits	(Adesina, 2005)	
Pimpinellin (101)	Stem bark	(Adesina, 2005)	
Psoralen (102)	Fruits	(Adesina, 2005)	
Scoparone (50)	Fruits, roots and bark	(Adesina, 2005; Misra et al., 2013	
Scopoletin (103)	Stem bark	(Adesina, 2005)	
Umbelliferone (104)	Stem bark	(Adesina, 2005)	
Xanthotoxin (51)	Fruits and stem bark	(Adesina, 2005)	
Lignans			
Asarinin (52)	Roots	(Adesina, 2005)	
Sesamin (54)	Roots	(Chaaib <i>et al.</i> , 2003)	
Xanthoxylol (105)	Roots	(Elujoba and Nagels, 1985)	
Triterpenes			
$\beta$ -Amyrin ( <b>106</b> )	Roots	(Adesina, 2005)	
$\beta$ -Sitosterol (9)	Fruits and roots	(Chaaib <i>et al.</i> , 2003; Misra <i>et al.</i> , 2013)	
Campesterol (107)	Roots	(Adesina, 2005)	
Lupeol (8)	Fruits and roots	(Chaaib <i>et al.</i> , 2003; Misra et al., 2013)	
Squalene (108)	Roots	(Adesina, 2005)	
Stigmasterol (10)	Fruits and roots	(Chaaib <i>et al.</i> , 2003; Adesina, 2005; Misra <i>et al.</i> , 2013)	

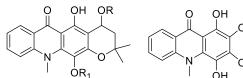
# Table 1.3 continued

Phenolic and flavonoids		
2-Hydroxymethylbenzoic acid	Roots	(Adesina, 2005)
(109)		
2,5-Dihydroxybenzoic acid	Leaves	(1 , 1 2000
(110)		(Isaac <i>et al.</i> , 2009
Burkinabin A (111)	Roots	(Ouattara et al., 2004;
		Queiroz et al., 2006)
Burkinabin B (112)	Roots	(Ouattara <i>et al.</i> , 2004)
Burkinabin C (113)	Roots	(Ouattara et al., 2004;
		Queiroz et al., 2006)
Caffeic acid (114)	Leaves and stem bark	(Asante <i>et al.</i> , 2009)
Chlorogenic acid (115)	Stem bark, roots	(Wangensteen <i>et al.</i> , 2017)
Diosmin (116)	Roots	(Adesina, 2005)
Gallic acid (117)	Leaves	(Asante <i>et al.</i> , 2009)
Hesperidin (59)	Roots	(Queiroz et al., 2006)
Kaempferol (118)	Stem bark	(Ogunbolude <i>et al.</i> , 2014)
Neochlorogenic acid (119)	Roots	(Wangensteen <i>et al.</i> , 2017)
para-Coumaric acid (120)	Leaves	(Asante et al., 2009)
para-Hydroxybenzoic acid	Roots	(Elujoba and Nagels, 1985)
(121)		
Quercetin (122)	Stem bark	(Ogunbolude <i>et al.</i> , 2014)
Rosmaniric acid (123)	Leaves	(Asante <i>et al.</i> , 2009)
Rutin (124)	Stem bark	(Ogunbolude <i>et al.</i> , 2014)
Syringic acid (125)	Leaves	(Elujoba and Nagels, 1985;
		Asante <i>et al.</i> , 2009)
Vanillic acid (126)	Leaves, roots	(Asante <i>et al.</i> , 2009;
		Elujoba and Nagels, 1985)
Quinoline alkaloids		
Acronycine (127)	Fruits and roots	(Adesina, 2005)
Atanine (128)	Ptericarps, Stem bark	(Torto <i>et al.</i> , 1973;
	and roots	Adesina, 2005)

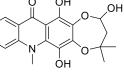
Table 1.3 continued

Dictamnine (129)	Roots	(Wangensteen <i>et al.</i> , 2017)
Fagarine (130)	Roots	(Adesina, 2005)
Skimmianine (55)	Roots	(Adesina, 2005)
Other compounds		
(-)- <i>p</i> -Synephrine ( <b>131</b> )	Roots	(Wangensteen <i>et al.</i> , 2017)
4-Methylthiocanthin-6-one	Fruits and roots	(Adesina, 2005)
(132)		
4'-(3"-Methylbut-2"-enyloxy)-	Roots	(Chaaib <i>et al.</i> , 2003)
3-phenylpropanol (133)		
4'-(4"-Hydroxy-3"-methyl-	Roots	(Chaaib <i>et al.</i> , 2003)
butyloxy)-2-phenylethyl (134)		
5-Methoxycanthin-6-one (135)	Fruits and roots	(Adesina, 2005)
6-Canthinone (136)	Roots	(Adesina, 2005)
7,8-Di- <i>O</i> -(3-methoxy-4-	Roots bark	(Chaaib <i>et al.</i> , 2003)
hydroxybenzoyl)-2,5-		
dihydroxycyclooctane-1,6-		
endoperoxyde (137)		
<i>p</i> -Mentha-1,8-dien-9-ol ( <b>138</b> )	Fruits	(Misra <i>et al.</i> , 2013)
Cuspidiol (139)	Roots	(Chaaib <i>et al.</i> , 2003)
Dihydrocuspidiol (140)	Roots	(Chaaib <i>et al.</i> , 2003)
Germacrone (141)	Roots	(Adesina, 2005)
Zantholic acid (142)	Fruits	(Wouatsa <i>et al.</i> , 2013b)





**68**  $R = R_1 = H$ 



69

**66**  $R_1 = R_3 = R_4 = R_5 = R_7 = H R_2 = CH_3 R_6 = OH$ **67**  $R_1 = R_4 = CH_3 R_2 = R_6 = H R_3 = R_5 = R_7 = OH$ 

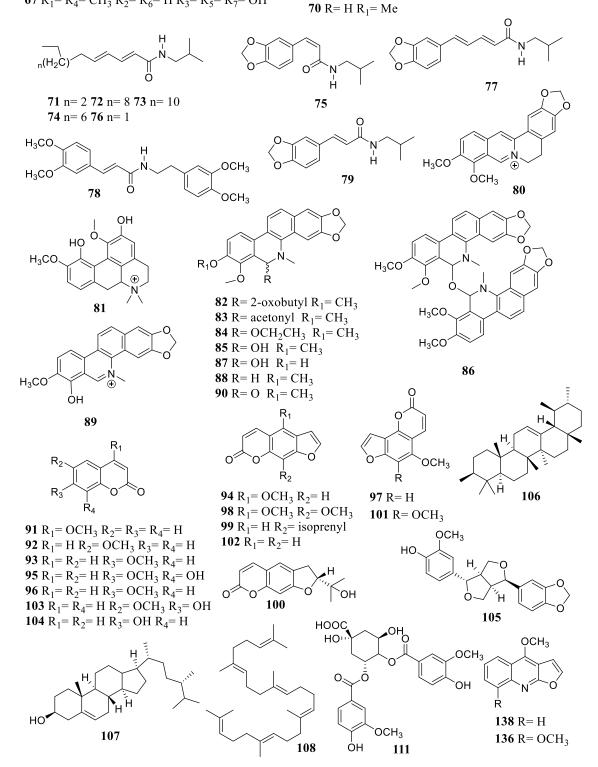


Figure 1.21 Identified and isolated compounds from Z. zanthoxyloides

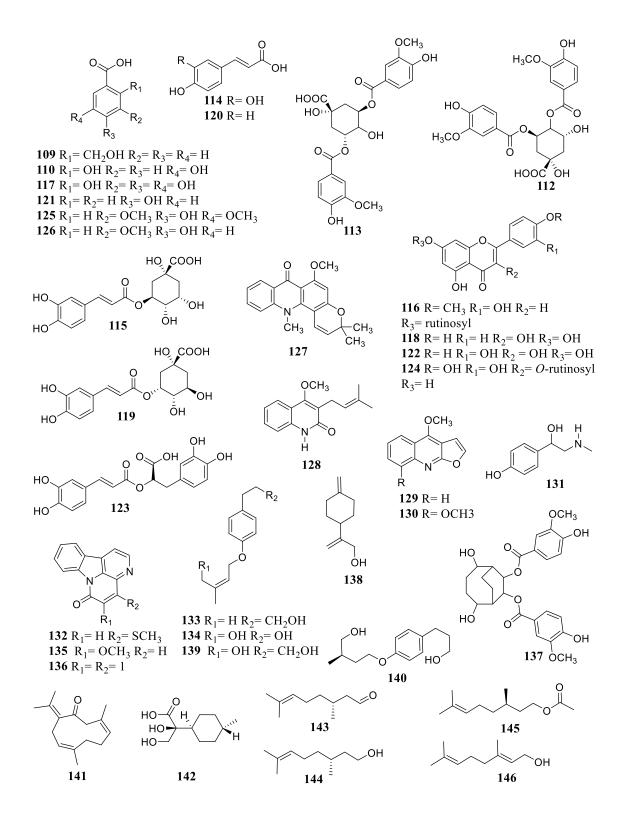


Figure 1.21 Continued

#### 1.6 Aim and objectives

Despite important medical and technological advances made in the development of new drugs, as well as considerable efforts in the treatment and the campaign against cancer and malaria, they remain two of the most commoncauses of death worldwide. About 200 million cases of malaria are reported every year giving rise to an estimated 445,000-500,000 deaths yearly (WHO, 2017). The burden of the disease is boosted by the resistance of the parasites to existing antimalarial drugs and by the resistance of the mosquito vector to insecticides in endemic areas particularly in sub-Saharan Africa, where 91% of deaths from malaria are reported. As for cancer, it is the second leading cause of death worldwide claiming about 8.8 million deaths in 2015 (WHO, 2018a). Globally, one in six deaths are due to cancer and this number is projected to almost double by the year 2030. Therefore, more research on alternative prevention and treatment are needed.

Traditional medicine is important and indispensable in the development and discovery of new drug leads. Indeed, screenings of medicinal plants based on their ethnopharmacological applications, especially in the treatment of tumours and malaria, have led to some remarkable discoveries in the past.

Thus, the aim of the present investigation was to identify phytochemicals with antimalarial and/or chemopreventive properties from five medicinal plants based on their folklore use in the Cameroonian community for the treatment of different types of fever, malaria and tumours.

This aim comprised following objectives:

• To obtain crude extracts from the selected plants and to evaluate their potent chemopreventive and antimalarial activities.

- To carry out bioassay directed isolation of active compounds.
- To characterise these bioactive compounds by spectroscopic means.
- To evaluate the biological activities of the isolated compounds.

# **Chapter 2 Materials and Methods**

# 2.1 Plant materials

Plants were collected from the Centre (Mount Eloundem, Mfoundi) and the West (Dschang Local Market) regions of the Republic of Cameroon, which is a tropical country located in sub-Saharan Africa at longitude 7.3697° N, and latitude 12.3547° E (Figure 2.1), with the assistance Mr Victor Nana, a botanist of the Cameroon National Herbarium (CNH), Yaoundé, Cameroon. Mr Nana also identified all the species collected by comparison with respective voucher specimens available at the CNH. Plant materials were collected during the months of June and November in 2015. Table 2.1 gives details on the different plants' parts collected, the voucher numbers, period and place of collection.

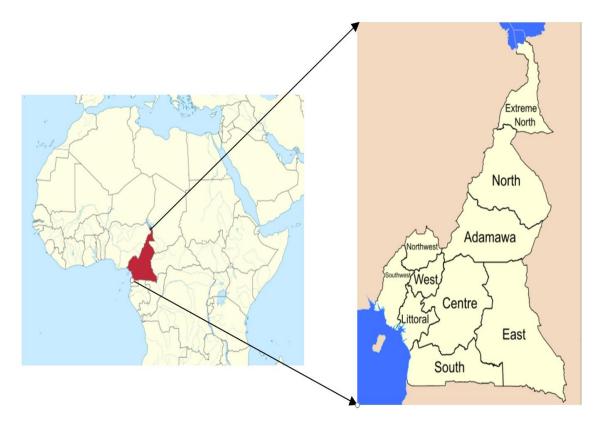


Figure 2.1 Map showing Cameroon geographical position and the different regions where the plants were collected

Table 2.1 List of medicinal	plants studied
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Plant name	Family	Voucher	Part	Period/	
		Number	collected	Region	
Croton oligandrus	Euphorbiaceae	6687/SFR	Stem bark	June 2015/	
Pierre ex Hutch			and leaves	Centre	
Justicia	Acanthaceae	37822/SFR	Leaves	June 2015/	
hypocrateriformis				Centre	
(Vahl) Milne-Redh					
Pseudospondias	Anacardiaceae	41437/SFR	Fruits,	June 2015/	
microcarpa (A. Rich.)			leaves and	Centre	
Engl.			stem bark		
Zanthoxylum lepreurii	Rutaceae	106669/SFR	Fruits	November	
Guill. and Perr.				2015/West	
Zanthoxylum	Rutaceae	21793/SFR	Fruits	November	
zanthoxyloides (Lam.)				2015/ West	
Zepern. and Timler					

### **2.2** Chemicals and reagents

#### 2.2.1 Chemicals

All the solvents were purchased from Fisher Scientific Ltd. (Loughborough, UK). Deionized water used in the present work was obtained from a MiliQ water system (Merck, Germany). Deuterated NMR solvents including chloroform (CDCl<sub>3</sub>), methanol (CD<sub>3</sub>OD), acetone (acetone-d<sub>6</sub>), pyridine (Pyr-d<sub>5</sub>), DMSO (DMSO-d<sub>6</sub>) and water (D<sub>2</sub>O) were purchased from Cambridge Isotope Laboratories Inc. (Tewksbury, USA). Precoated aluminium plates of Silica gel 60 PF<sub>254</sub> (0.2 mm thickness, 20 x 20 cm) for Thin Layer Chromatography (TLC) were sourced from Merck Ltd., Germany. Silica gel (70-230 mesh) and Silica Kieselgel 60H, purchased from Sigma-Aldrich Company Ltd. (Dorset, UK), were used for open column chromatography (CC) and vacuum liquid

chromatography (VLC), respectively. Sodium dodecyl sulphate (SDS), bovine hemin, sodium acetate buffer 3M, NaOH and 37% HCl solution were also purchased from Sigma-Aldrich Company Ltd.

## 2.2.2 Cell lines, cell culture media and reagents

Names and tissues of origin of each cell line used in this study are listed in Table 2.2. AREc32 cells were a generous gift of Prof Roland Wolf (University of Dundee, UK), while the other cell lines were obtained from Dr Andrews Evans cells bank (LJMU, Liverpool, UK).

Table 2.2 Cell lines used in bioassays

Name	Origin
AREc32	Human mammary MCF7-derived reporter cells
A549	Adenocarcinoma human alveolar basal epithelial cells
MCF7	Human breast adenocarcinoma cells
PC3	Human prostate cancer cell line
PNT2	Human normal prostate epithelium cells

Foetal bovine serum (FBS), penicillin/streptomycin, L-glutamine, Dulbecco's Modified Eagles Medium (DMEM), trypsin-EDTA and phosphate buffer saline (PBS), were purchased from Biosera (Neuville, France). Doxorubicin, gentamicin and trypan blue solution were obtained from Sigma Aldrich Company Ltd. (Dorset, UK). Luciferase assay kit reagent was from Promega Corp. (Loughborough, UK).

#### 2.3 Phytochemical work

#### **2.3.1 Plant preparation and Soxhlet extraction**

The plant materials were individually cut into small pieces; air dried at room temperature and powdered using a maize grinder into a coarse powder. The powdered plant materials were extracted using Soxhlet extraction.

Individual plant material was placed in a thimble made of thick filter paper and placed in the extraction chamber, extraction solvent (0.9 L) was poured in and the condenser was placed on the extraction chamber (Figure 2.2). The solvent was brought to boil using an Electrothermal<sup>TM</sup> electromantle (Fisher Scientific, Loughborough-UK). When solvent boiled, the extraction chamber was gradually filled with solvent and performed extraction until it reached the maximum level, when the solvent with the dissolved extracted components went back to the flask. This was one cycle of extraction. This cycle was repeated 10 times for each solvent used. In this way, the dissolved components remained in the flask, while clean solvent was evaporated and the plant material in the thimble was repeatedly extracted (Rostagno and Prado, 2013). The plant material was extracted, successively, with *n*-hexane, dichloromethane (DCM) and methanol (MeOH). The resulting extracts were concentrated to dryness under reduced pressure on a Cole-Palmer rotary evaporator (Stone, UK). Extracts were stored at 4°C and dissolved in appropriate solvent prior to usage.

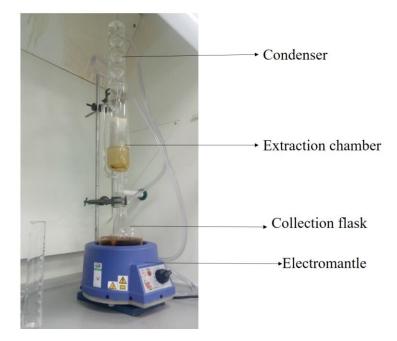


Figure 2.2 Soxhlet apparatus

# **2.3.2** Chromatographic techniques

2.3.2.1 Vacuum liquid chromatography

Vacuum liquid chromatography (VLC) was used for the rapid fractionation of active nonpolar and medium polarity extracts. The technique was used as described by Pelletier *et al.* (1986). The system consisted of a sintered glass Buchner filter funnel with an attachment to a side arm flask connected to a vacuum pump (Figure 2.3). Approximately the two third of the VLC funnel was dry-packed with Kieselgel 60H, the stationary phase, under vacuum and the appropriate solvent was allowed to flow through the system (Reid and Sarker, 2012). A portion of the *n*-hexane or DCM extract were dissolved in a little amount of appropriate solvent, mixed with normal silica gel (70-230 mesh), taken to dryness either by air drying or under reduced pressure using rotary evaporator and the mixture was loaded on the top of the funnel as a uniform thin layer. Vacuum was applied, and the column was eluted with a stepwise gradient of mobile phase consisting of an increasing amount of ethyl acetate in *n*-hexane (Hex/EA 0%, 10%, 20%, 40%, 60% and 80%, H1-H6) or MeOH in DCM (DCM/MeOH 0%, 2%, 6%, 10%, 15% and 25%, D1-D6) for the *n*-hexane or DCM extract respectively to obtained six different fractions. Collected VLC fractions were concentrated to dryness under reduced pressure and analysed by TLC to get an idea of their composition. Fractions with similar TLC profiles were mixed together.

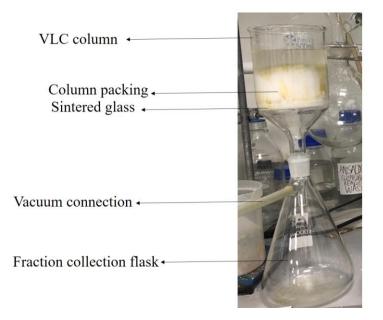


Figure 2.3 VLC system

# 2.3.2.2 Solid-phase extraction

Solid-phase extraction (SPE) was used for the fractionation of active methanolic extract and to clean-up DCM extract prior to preparative HPLC. The SPE is quite similar to VLC with the only difference being that a commercially available pre-packed SPE cartridge is used instead of a funnel prepacked with silica gel (Reid and Sarker, 2012) (Figure 2.4). A Strata C-18-E cartridge 20 g (Phenomenex, California-USA) was used for this study. A portion of the dried MeOH extract (2 g) was suspended in 10 mL of a 10% MeOH in water mixture and loaded into the cartridge, previously washed with MeOH (50 mL), followed by equilibration with water (100 mL). The cartridge was eluted with MeOH– water mixture of decreasing polarity to obtain four fractions (F1-F4): 20, 50, 80 % MeOH in water and 100% MeOH (200 mL each). All four fractions were evaporated to dryness using a combination of a rotary evaporator and a freeze-dryer. A portion of each resulting fraction was dissolved in MeOH to obtain a solution of 1 mg/mL and analysed by HPLC-UV/DAD. For the clean-up of DCM extract, 2 g of the sample were dissolved in 10 mL 30% MeOH in water solution and the cartridge was eluted with 80% MeOH in water solution.

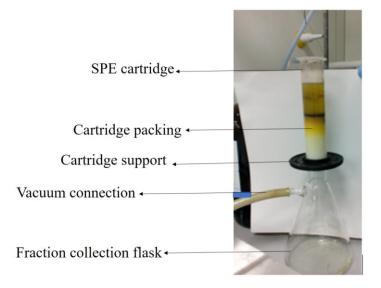


Figure 2.4 Solid Phase Extraction system

### 2.3.2.3 Thin layer chromatography

Thin layer chromatography (TLC) is a separation technique mainly used for the detection and monitoring of compounds through a separation process based on the criteria of adsorption, binding to silica and polarity (Gibbons, 2012). The technique was used in this study to screen the plant extracts for the presence of secondary metabolites, to determine the starting eluting solvent composition for column chromatography, to monitor fractions collection during chromatography, to check the purity of isolated compounds from CC and as isolation technique for the purification of compounds (preparative TLC). Precoated silica gel 60 PF<sub>254</sub> aluminium plates were used for this purpose. Samples were applied manually at the base, about 1 cm from the bottom edge of the TLC plate, using a glass capillary and developed in a chromatographic chamber using an appropriate solvent system. Multiple developments were sometime needed to improve resolution. After drying, the developed plates were visualised by irradiation with short-wave (254 nm) and long-wave (366 nm) using a Camag UV lamp (CAMAG, Hungerford-UK) and by spraying with a 1% anisaldehyde solution in aqueous H<sub>2</sub>SO<sub>4</sub> followed by heating to 105 °C in Sciquip oven 30S (Sciquip, Shrewsbury-UK) for 5 min (Figure 2.5).

In the case of preparative TLC, only a small part (1/10 part) of TLC plate was spread with the anisaldehyde reagent after having covered the remaining part with aluminium foil. Bands of interest were identified and scratched with the help of spatula. The compounds of interest were extracted from the collected silica gel by cold maceration with a mixture of 10% MeOH in DCM.

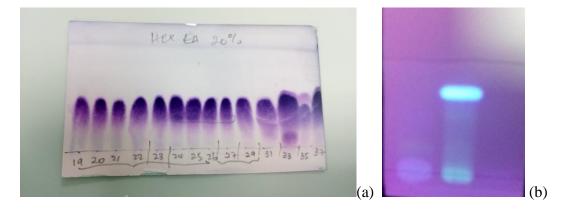


Figure 2.5 TLC plates of a) xylopic acid developed in Hex:EtOAC (8:2) and sprayed with 1% anisaldehyde and b) skimmianine developed in DCM:MeOH (9.2:0.8) visualised at 254 nm

## 2.3.2.4 Column chromatography

Column chromatography (CC) was used to isolate compounds from active *n*-hexane fraction. Silica gel (70-230 mesh) was used for this purpose. The sample and the column (Figure 2.6) were prepared in the same way as described for VLC, but in the case of CC the elution was carried out purely under the gravitational force, and no negative or positive pressure was applied. Wet packing technique was also used instead of dry packing as for the case of VLC. *n*-Hexane was used as packing solvent. Excess solvent was run through the column to allow the silica gel to settle well before introducing the sample on the top of the column. The column was eluted with a stepwise gradient of mobile phase of increasing polarity. The initial solvent composition was determined by TLC analysis. Depending on the column size, 10-50 mL of eluents were collected per fraction, and their contents were analysed by TLC.



Figure 2.6 CC system operated by manual fraction collection

### 2.3.2.5 High-performance liquid chromatography

High-performance liquid chromatography (HPLC) is a separation technique both used as analytical and preparative for the detection, separation, quantification and isolation of compound mixtures, including natural products.

## 2.3.2.5.1 Analytical high-performance liquid chromatography

Analytical HPLC was used to screen the plant extracts for the presence of secondary metabolites, to develop methods for isolation using preparative HPLC and to verify the purity of isolated compounds. The experiment was performed on an UPLC Dionex 3000 (Thermoscientific, UK) or Agilent 1260 infinity series (Agilent, UK) both equipped with a binary pump, an autosampler, a column chamber, a degasser and a UV/DAD detector from the same corresponding model and company. Samples were prepared in MeOH (1 mg/mL) and analysed on a Phenomenex Gemini-NX 5 U C18 column ( $150 \times 4.6$  mm, Phenomenex, USA) or a Hypersil 5 U C18 column ( $150 \times 4.6$  mm, Phenomenex, USA) with a gradient of 30-100% MeOH (containing 0.1% TFA), 70-0% H<sub>2</sub>O (containing 0.1% TFA) over 30 min or of 30-100% ACN, 70-0% H<sub>2</sub>O (containing 0.1% TFA). The column temperature was set at 25 °C. A volume of 10 µL of the prepared sample was injected and the chromatogram was monitored at 1 mL/min using variable UV/vis wavelengths comprising between 205 and 366 nm. Methods described above were always used to verify the purity of the collected fractions obtained from preparative HPLC analyses and as starting methods for the development of further methods (Table 2.3) to be used on preparative HPLC for isolation and purification of compounds.

### 2.3.2.5.2 Preparative high-performance liquid chromatography

Preparative HPLC was used for the isolation and purification of compounds from the DCM and MeOH fractions using developed methods from analytical HPLC. Table 2.3 below shows the different methods developed on the analytical HPLC and used on the prep-HPLC for the isolation of compounds. The experiment was carried out on an Agilent 1260 infinity series equipped with a binary pump, a degasser, a column chamber and a UV/DAD detector. An ACE Gemini-NX 5 U C18 column (150 × 21.2 mm, Hichrom Ltd, UK) maintained at 25 °C and monitor at a flow rate of 10 mL/min was used for isolation. Manual peak collection was carried on using UV/DAD detector and repeated chromatography were needed to isolate a good amount of compound. The samples were all prepared in MeOH. MeOH and  $H_2O$  used for HPLC analysed contained 0.1% TFA.

	Time slot (min)	%MeOH	%ACN	%H2O
Method A	0 - 30	30 - 100	-	70 - 0
	30 - 35	100	-	0
Method B	0 - 15	-	30 - 65	70 - 35
Method C	0 - 20	30 - 75	-	70 - 25
Method D	0 - 10	45 - 50	-	55 - 50
	10 - 30	50 - 60	-	50 - 40
Method E	0 - 30	30 - 65	-	70 - 35

Table 2.3 List of the different HPLC methods developed

### 2.3.3 Isolation of compounds

2.3.3.1 Isolation of compounds from Croton oligandrus

The air-dried and powdered stem bark (330.8 g) of *C. oligandrus* were extracted with *n*-hexane, DCM and MeOH to obtain 3.7, 2.1 and 7.4 g of beige oily, beige and brown extracts, respectively.

A portion of the *n*-hexane (3.3 g) and DCM (1.8 g) extracts were adsorbed on to normal silica gel (70-230 mesh) and fractionated using VLC to obtain six fractions each (H1-H6) and (D1-D6) for *n*-hexane and DCM extracts, respectively, following the VLC procedure as described earlier.

Fraction H3 (420.9 mg) of *n*-hexane extract was further purified using a CC over silica gel using a gradient of Hex-EtOAc 0-30% to give a mixture (18.2 mg) of **149** and **150**, and **147** (4.4 mg). Compounds **7** (16.1 mg) and **8** (19.0 mg) were obtained from H5, H6 and D4 by recrystallisation using Hex-EtOAc 15-30%. The mixture of **149** and **150** (3.2 mg) were also obtained from D2 and D3 of the DCM crude extract by recrystallisation with Hex-EtOAc 15%. D4, D5 and D6 showed similar TLC and HPLC profiles and were then mixed together for purification. The mixture of the two fractions (783.0 mg) was finally purified by preparative RP-HPLC using the method A (Table 2.3). Figure 2.7 showed the obtained chromatogram as well as the peaks corresponding to the different compounds (**148**, **151-162**) identified and characterised from the combined fractions.

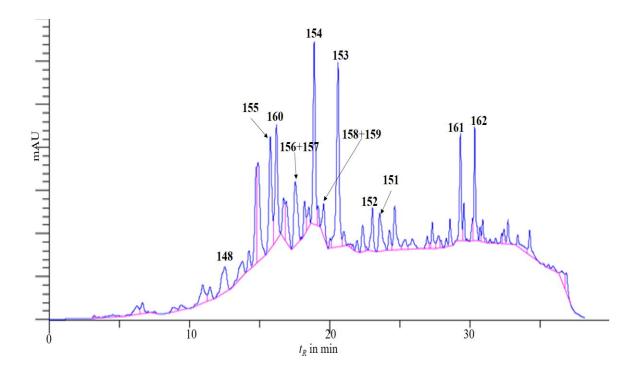


Figure 2.7 Chromatogram of *C. oligandrus* fractions D4+D6-method A, inj. vol. 130  $\mu$ L, sample conc. 182.7 mg/mL, monitored at 215nm

Table 2.4 Isolation of compounds from C. oligandrus

Extract	VLC	Mode of	Rf or tR	Visual aspect	Amount
containing	fraction	isolation	in min		in mg
the cpds	containing				
	the cpds				
<i>n</i> -hexane	H5, D4	-	0.47*	White powder	16.1
<i>n</i> -hexane	H6, D4	-	0.40*	White powder	19.0
<i>n</i> -hexane	H3	CC	0.56*	White powder	4.4
DCM	D4, D5 and	HPLC	12.51	White powder	8.3
	D6				
DCM, n-	H3, D2 and	CC	0.63*	White powder	21.4
hexane	D3				
DCM	D4, D5 and	HPLC	23.54	White powder	6.3
	D6				
DCM	D4, D5 and	HPLC	23.03	White powder	3.3
	D6				
DCM	D4, D5 and	HPLC	20.57	White powder	15.6
	D6				
DCM	D4, D5 and	HPLC	18.89	White powder	13.9
	D6				
DCM	D4, D5 and	HPLC	15.76	White powder	2.9
	D6				
DCM	D4, D5 and	HPLC	17.54	White powder	7.8
	D6				
DCM	D4, D5 and	HPLC	19.55	White powder	6.5
	D6				
DCM	D4, D5 and	HPLC	16.19	White powder	7.3
	D6				
DCM	D4, D5 and	HPLC	29.29	Yellow	4.1
	D6			powder	
DCM	D4, D5 and	HPLC	30.31	White powder	5.7
	D6				
	containing the cpds n-hexane n-hexane DCM DCM, n- hexane DCM DCM DCM DCM DCM DCM	containing the cpdsfraction containing the cpdsn-hexaneH5, D4n-hexaneH6, D4n-hexaneH3DCMD4, D5 and D6DCM, n-H3, D2 and hexaneDCMD4, D5 and D6DCMD4, D5 and D6	containing the cpdsfraction containing the cpdsisolationn-hexaneH5, D4-n-hexaneH6, D4-n-hexaneH3CCDCMD4, D5 and D6HPLCDCMD4, D5 and D6 <t< td=""><td>containing the cpdsfraction containing the cpdsisolation in minn-hexaneH5, D4-0.47*n-hexaneH6, D4-0.40*n-hexaneH3CC0.56*DCMD4, D5 and D6HPLC12.51DCM, n-H3, D2 and D6CC0.63*bexaneD3CC0.63*DCMD4, D5 and D6HPLC23.54DCMD4, D5 and D6HPLC23.03DCMD4, D5 and D6HPLC23.03DCMD4, D5 and D6HPLC20.57DCMD4, D5 and D6HPLC18.89DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC16.19DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCM<td< td=""><td>containing the cpdsfraction containing the cpdsisolationin minin minn-hexaneH5, D4-0.47*White powdern-hexaneH6, D4-0.40*White powdern-hexaneH3CC0.56*White powderDCMD4, D5 andHPLC12.51White powderDCM, n-H3, D2 andCC0.63*White powderhexaneD3CC0.63*White powderDCMD4, D5 andHPLC23.54White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.74White powderDCMD4, D5 andHPLC17.54White powderDCMD4, D5 andHPLC19.55White powderDCMD4, D5 andHPLC16.19White powderDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC20.31White powder</td></td<></td></t<>	containing the cpdsfraction containing the cpdsisolation in minn-hexaneH5, D4-0.47*n-hexaneH6, D4-0.40*n-hexaneH3CC0.56*DCMD4, D5 and D6HPLC12.51DCM, n-H3, D2 and D6CC0.63*bexaneD3CC0.63*DCMD4, D5 and D6HPLC23.54DCMD4, D5 and D6HPLC23.03DCMD4, D5 and D6HPLC23.03DCMD4, D5 and D6HPLC20.57DCMD4, D5 and D6HPLC18.89DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and D6HPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC15.76DCMD4, D5 and HPLCHPLC16.19DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCMD4, D5 and HPLCHPLC29.29DCM <td< td=""><td>containing the cpdsfraction containing the cpdsisolationin minin minn-hexaneH5, D4-0.47*White powdern-hexaneH6, D4-0.40*White powdern-hexaneH3CC0.56*White powderDCMD4, D5 andHPLC12.51White powderDCM, n-H3, D2 andCC0.63*White powderhexaneD3CC0.63*White powderDCMD4, D5 andHPLC23.54White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.74White powderDCMD4, D5 andHPLC17.54White powderDCMD4, D5 andHPLC19.55White powderDCMD4, D5 andHPLC16.19White powderDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC20.31White powder</td></td<>	containing the cpdsfraction containing the cpdsisolationin minin minn-hexaneH5, D4-0.47*White powdern-hexaneH6, D4-0.40*White powdern-hexaneH3CC0.56*White powderDCMD4, D5 andHPLC12.51White powderDCM, n-H3, D2 andCC0.63*White powderhexaneD3CC0.63*White powderDCMD4, D5 andHPLC23.54White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC20.57White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.76White powderDCMD4, D5 andHPLC15.74White powderDCMD4, D5 andHPLC17.54White powderDCMD4, D5 andHPLC19.55White powderDCMD4, D5 andHPLC16.19White powderDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC29.29YellowDCMD4, D5 andHPLC20.31White powder

\*TLC plate developed in Hex:EtOAc 40%

The MeOH extract was not processed further as it was found to be non-active. In addition, the same extraction procedure was carried out with 280.0 g of the dried leaves to obtain 6.9, 2.9 and 9.9 g of *n*-hexane, DCM and MeOH extracts, respectively. Only the *n*-hexane extract was found to be active. However, it was not processed further as preliminary <sup>1</sup>H NMR revealed a mixture of **7**, **8** and fatty acids.

# 2.3.3.2 Isolation of compounds from Justicia hypocrateriformis

The air-dried and powdered leaves (450.5 g) of *J. hypocrateriformis* were extracted with *n*-hexane, DCM and MeOH to obtain 13.6, 10.2 and 41.8 g of dark green, brown green and dark brown extracts, respectively. Preliminary screening revealed MeOH as the only active extract.

The MeOH extract (4 g) was dissolved in MeOH and cleaned using SPE eluted with MeOH/H<sub>2</sub>O 50% to remove chlorophyll. After drying, 2 g of the obtained extract was subjected to standard SPE procedure described above. F1 (668.8 mg) was adsorbed on to normal silica gel and subjected to CC using of a gradient of EtOAc:MeOH 0-30%. Forty-one fractions of 50 mL were collected and grouped based on their TLC profiles to eight main subfractions (F1A-F1H). A white crystalline precipitate identified as sucrose (**60**) was obtained from F1H (CC fractions 30-41, collected with 24-30% MeOH in EtOAc). Compound **170-171** were obtained by PTLC of CC fractions 17-20 (EtOAc:MeOH 13%). The Plate was developed in DCM:MeOH 10%.

Fractions F2 (428.6 mg) and F3 (361.8 mg) were subjected to preparative RP-HPLC using method C and D (Table 2.3) respectively. Chromatograms of the fractions are shown on Figure 2.8. Seven compounds (**163-169**) were identified and characterised from the

collected peaks. Overall, ten compounds were isolated and characterised from the leaves of *J. hypocrateriformis* (Table 2.5).

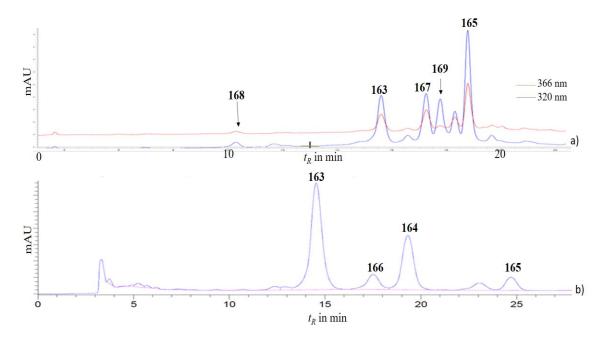


Figure 2.8 Chromatogram of *J. hypocrateriformis* fractions, a) F2, method C, injection vol. 500  $\mu$ L, sample conc. 25 mg/mL, monitored at 320 and 366 nm b) F3, method D, injection vol. 100  $\mu$ L, sample conc. 50 mg/mL, monitored at 205 nm

Cpds	SPE fractions	Mode of	$\mathbf{R}_{\mathbf{f}}$ or $t_{\mathbf{R}}$	Visual aspect	Amount
	containing the	isolation	in min		in mg
	compounds				
163	F2, F3	HPLC	15.53	Yellow powder	8.4
164	F3	HPLC	19.34	Yellow powder	2.9
165	F2, F3	HPLC	24.71	Yellow powder	2.2
166	F3	HPLC	17.50	Yellow powder	3.3
167	F2	HPLC	16.75	Yellow powder	2.6
168	F2	HPLC	27.01	Yellow powder	4.4
169	F2	HPLC	17.75	Brown powder	1.6
170+171	F1	PTLC* of CC	0.37	Brown powder	3.4
		pooled fractions			
		17-20			

\* PTLC developed in DCM:MeOH 10%.

Stem bark (276.0 g), fruits (262.0 g) and leaves (437.0 g) of *P. microcarpa* were individually extracted to yield 2.2, 4.1 and 13.4 g of *n*-hexane; 1.0, 0.9 and 2.7 g of DCM; and 5.9, 7.1 and 12.5 g of MeOH extract, respectively. Only the stem bark DCM and MeOH extracts, and the leaves MeOH extract were subjected to chromatographic separation.

The bark DCM extract was cleaned up, while the MeOH extract was subjected to standard SPE prior to isolation of their phytochemicals by preparative RP-HPLC. F2 and F3 obtained from the SPE of the MeOH extract were found to have similar chromatograms after analytical HPLC and combined to a single fraction. The DCM extract (597.6 mg) and the MeOH fractions F2+F3 (194.3 mg) were chromatographed using methods A and E (Table 2.3), respectively. Both extracts afforded compounds **103**, **148** and **175** (Table 2.7) from the peaks collected.

The leaves MeOH extract was cleaned up and subjected to standard SPE method as described for *J. hypocarteriformis*. F2 (198.8 mg) and F3 (328.6 mg) were found to have similar analytical HPLC profiles and were mixed together for separation of their constituents. Separation was performed using HPLC method C (Table 2.3). Four compounds (**164**, **166**, **172** and **173**) were identified and characterised from the obtained chromatogram (Figure 2.10). F1 and F4 were not processed further. Table 2.6 lists all the compounds isolated and characterised from *P. microcarpa*.

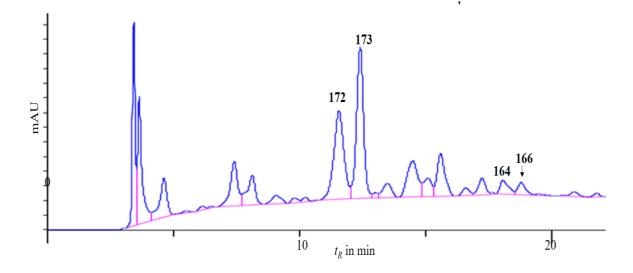


Figure 2.9 Chromatogram of *P. microcarpa* leaves fractions F2 + F3, method C, injection vol. 120  $\mu$ L, sample conc. 75 mg/mL, monitored at 230 nm

Cpds	Part of plant	Extract	SPE/VLC	Mode of	$t_R$ in	Visual	Amount
	containing	containing	containing	isolation	min	aspect	in mg
	the cpds	the cpds	the cpds	(method)			
103	Stem bark	DCM	-	HPLC (A)	4.55	Brown	3.1
		МеОН	F2 and F3	HPLC (E)	13.28	powder	
148	Stem bark	DCM	-	HPLC (A)	4.28	Beige	3.6
		МеОН	F2 and F3	HPLC (E)	13.00	powder	
164	Leaves	MeOH	F2 and F3	HPLC (C)	18.67	Brown	1.9
						powder	
166	Leaves	MeOH	F2 and F3	HPLC (C)	19.23	Brown	2.1
						powder	
172	Leaves	MeOH	F2 and F3	HPLC (C)	11.20	Yellow	6.1
						powder	
173	Leaves	МеОН	F2 and F3	HPLC (C)	14.69	Brown	2.4
						powder	
174	Stem bark	DCM	-	HPLC (A)	4.78	Brown	4.3
		МеОН	F2 and F3	HPLC (E)	13.33	powder	

Table 2.6 Isolation of compounds from P. microcarpa

## 2.3.3.4 Isolation of compounds from Zanthoxylum leprieurii

The air-dried and powdered fruits (546.8 g) of *Z. leprieurii* were extracted with *n*-hexane, DCM and MeOH to obtain 32.7, 21.1 and 54.3 g of brown oily, light maroon and dark maroon extracts, respectively.

The brown oily *n*-hexane extract (20 g) was subjected to CC over silica gel 60 (230-400 mesh, 6.5 cm x 50 cm) using a gradient system of Hex:EtOAc (0-50%) and DCM-MeOH (10-30%) as eluents. Fifty-three sub-fractions (ca. 125 mL each) were collected and pooled on the basis of their analytical TLC profile to six main fractions A-F. Fraction B (3.8 g, pooled sub-fractions 9-20) was further chromatographed using silica gel 60 (230-400 mesh) CC with increasing amounts of EtOAc in *n*-hexane as eluent to afford **175** (120.5 mg) and **176** (230.2 mg). Fraction C (1.9 g, pooled sub-fractions 22-32) was subjected to repeated CC as previously described to yield **178** (7.4 mg), **179** (80.7 mg), along with a mixture of sterols. A white granular precipitate giving single pink spot on TLC and identified as **177** (43.1 mg) was obtained from fraction F (2.1 g, pooled sub-fractions 48-53).

Fractions D4 (2.46 g), D5 (533.5 mg) and D6 (892.3 mg) obtained by VLC of the DCM extract (7.11 g) were mixed together based on their TLC profiles. The mixture obtained (2 g) was cleaned up using SPE, dried, resuspended in MeOH and subjected to preparative RP-HPLC using HPLC method C (Table 2.3) for compounds isolation. From the peaks collected, compounds **114**, **180-182** were identified and characterised.

The MeOH extract was found to be inactive and was not purified. Table 2.7 lists all the compounds isolated from *Z. leprieurii* and their methods of isolation.

Table 2.7 isolation of compounds from Z. leprieurii

Cpds	Extract	VLC	Mode of isolation	R <sub>f</sub> or	Visual	Amount
	containing	fractions		$t_R$ in	aspect	in mg
	the cpds	containing		min		
		the cpds				
114	DCM	D4, D5 and	HPLC	10.62	Brown	4.4
		D6			powder	
175	<i>n</i> -hexane	-	CC of CC pooled	0.47*	White	120.5
			fractions 9-20		powder	
176	<i>n</i> -hexane	-	(Hex:EtOAc 10-	0.38*	White	230.2
			20%) eluted with		powder	
			EtOAc in <i>n</i> -			
			hexane (5-15%)			
177	<i>n</i> -hexane	-	Precipitates in CC	0.12*	White	43.1
			pooled fractions		powder	
			48-53			
			(DCM:MeOH			
			15%)			
178	<i>n</i> -hexane	-	CC of CC pooled	0.18*	White	7.4
			fractions 22-32		powder	
179	<i>n</i> -hexane	-	(Hex:EtOAc 20-	0.22*	White	80.7
			30%) eluted with		powder	
			EtOAc in <i>n</i> -			
			hexane			
			(1-20%)			
180	DCM	D4, D5 and	HPLC	9.41	Brown	3.6
		D6			powder	
181	DCM	D4, D5 and	HPLC	7.70	Brown	1.3
		D6			powder	
182	DCM	D4, D5 and	HPLC	9.42	Yellow	3.3
		D6			powder	
			  / and annoved with 1	L	l	

\*TLC developed in Hex:EtOAC 25% and sprayed with 1% anisaldehyde solution

The air-dried ground fruits (350.0 g) of *Z. zanthoxyloides* were extracted to give 34.6 g, 3.9 g and 19.6 g of *n*-hexane, DCM and MeOH extract, respectively.

The DCM extract (3.0 g) was subjected to VLC to obtained six fractions D1-D6 as describe above. Fractions D4 (336.0 mg), D5 (223.2 mg) and D6 (129.9 mg) were mixed based on their TLC profiles, cleaned up and subjected to RP-HPLC for separation. Compounds **54**, **55**, **79**, **128**, **184** and **187** were identified and characterised from the fractions collected from the obtained chromatogram (Figure 2.10).

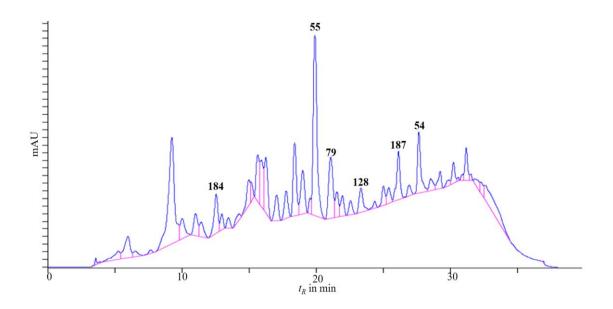


Figure 2.10 Chromatogram of Z. zanthoxyloides DCM fractions D4-D6, method A, injection vol.  $300 \,\mu$ L, sample conc.  $50 \,\text{mg/mL}$ , monitored at 220 nm

The MeOH extract was subjected to standard SPE and four fractions F1-F4 were obtained. The procedure was repeated twice to obtain a good amount of each fraction. F2 (704.8 mg) was subjected to preparative HPLC analysis (Figure 2.11a) using HPLC method B (Table 2.3) to yield **183** (3.5 mg) and **184** (3.6 mg) having the retention times ( $t_R$ ) 4.35 and 5.95 min, respectively. Fraction F2-B (5.7 mg) collected at  $t_R$  = 5.28 min was further purified through preparative TLC to afford **189** (3.2 mg, EtOAc:MeOH 7:3, R<sub>f</sub> 0.41). Fraction F2-D (3.6 mg) collected at  $t_R = 6.55$  min was also purified by preparative TLC to obtain **190** (1.9 mg, EtOAc:MeOH 7:3, R<sub>f</sub> 0.46). Preparative TLC of fraction F2-E (3.8 mg) collected at  $t_R = 7.22$  min provided **188** (1.8 mg, EtOAc:MeOH 7:3, R<sub>f</sub> 0.41). Purification of fraction F2-F (10.2 mg) collected at  $t_R = 7.61$  min using TLC with a mixture of EtOAc:MeOH (7:3) as eluent afforded more of **188** (2.5 mg, R<sub>f</sub> 0.41) and **186** (3.6 mg, R<sub>f</sub> 0.28).

F3 (845.8 mg) was also analysed by preparative HPLC using method A (Figure 2.11b). Compounds **54**, **55**, **59**, **128**, **183**, **185**, **187** and **191** were identified and characterised from the collected fractions.

Fractions F1 and F4 were not processed further. F1 was found to be rich in sugar while F4 were found to contain the same compounds as the DCM extract. Table 2.8 groups all the characterised compounds from *Z. zanthoxyloides*.

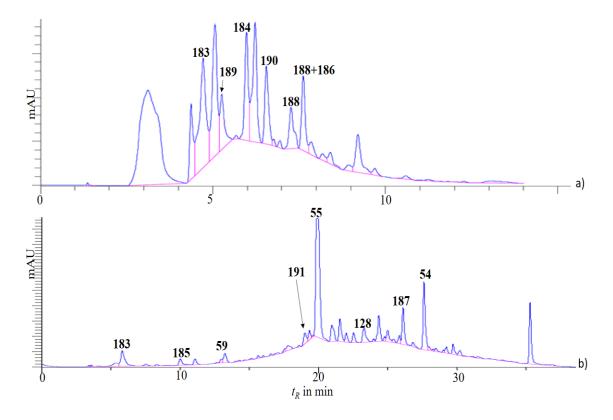


Figure 2.11 Chromatogram of *Z. zanthoxyloides* a) F2, method B, inj vol. 280  $\mu$ L, sample conc. 182.5 mg/mL, monitored at 254 nm b) F3, method A, inj vol. 120  $\mu$ L, sample conc. 100 mg/mL, monitored at 230 nm

Table 2.8 Isolation of compounds from Z. zanthoxyloides

Cpds	Extract (s)	VLC/SPE	VLC/SPE Mode of isolation		Visual	Amount
	containing	fraction (s)	(method)	$t_R$ in	aspect	in mg
	the cpds	containing		min		
		the cpds				
54	DCM,	D4-D6, F3	HPLC (A)	27.35	Brown	20.7
	MeOH				powder	
55	DCM,	D4-D6, F3	HPLC (A)	19.90	White	36.1
	MeOH				needles	
59	МеОН	F3	HPLC (A)	13.14	Yellow	1.9
					powder	
79	DCM	D4-D6	HPLC (A)	20.43	Brown	5.4
					powder	
128	DCM,	D4-D6, F3	HPLC (A)	24.01	Yellow	3.9
	MeOH				powder	
183	MeOH	F2, F3	HPLC (A and B)	4.88	Yellow	11.5
					powder	
184	DCM,	D4-D6, F2	HPLC (B)	12.42	Beige	3.6
	MeOH				powder	
185	MeOH	F3	HPLC (A)	10.00	Brown	1.2
					powder	
186	МеОН	F2	1. HPLC (B)	0.28	Yellow	3.6
			2. PTLC of HPLC		powder	
			fraction ( $t_R$ 7.61)			
			in EtOAc:MeOH			
			7:3			
187	DCM,	D4-D6, F3	HPLC (A)	25.85	Yellow	11.2
	MeOH				powder	
188	MeOH	F2	1. HPLC (B)	0.41	White	4.3
			2. PTLC of HPLC		liquid	
			fraction ( $t_R$ 7.61)			
			in EtOAc:MeOH			
			7:3			

Table 2.8 continued

189	MeOH	F2	HPLC (B)	0.41	White liquid	3.2
190	MeOH	F2	1. HPLC (B)	0.46	White liquid	1.9
			2. PTLC of HPLC fraction ( $t_R$			
			7.61) in EtOAc:MeOH 7:3			
191	MeOH	F3	HPLC (A)	23.77	Yellow	1.7
					powder	

### 2.3.4 Identification and characterisation of isolated compounds

Structure elucidation of all the isolated compounds was carried out by spectroscopic analyses including 1D (<sup>1</sup>H and <sup>13</sup>C NMR) and 2D NMR (<sup>1</sup>H-<sup>1</sup>H COSY, DEPT 145, <sup>1</sup>H-<sup>13</sup>C HSQC-DEPT, <sup>1</sup>H-<sup>13</sup>C HMBC, <sup>1</sup>H-<sup>1</sup>H NOESY and <sup>1</sup>H-<sup>1</sup>H TOCSY) and mass spectrometry. Some compounds were also characterised by UV, FT-IR spectroscopy and polarimetry. Spectroscopic techniques are useful means for assessing the purity and the determination of the structures of natural substances (Boiteau *et al.*, 1964).

#### 2.3.4.1 Nuclear magnetic resonance

Nuclear magnetic resonance (NMR) is a useful spectroscopic technique to obtain structural information about a given molecule. NMR spectra are plots of signals arising from absorption of radio frequency due to excitement of a nucleus or simultaneous excitement of two nuclei (Breitmaier and Sinnema, 1993; Crews *et al.*, 1998). Only <sup>1</sup>H and <sup>13</sup>C nuclei were excited during experiments. Experiments consist of 1D and 2D experiments. 1D experiments used in this study include proton (<sup>1</sup>H) and carbon (<sup>13</sup>C) NMR while 2D NMR experiments are <sup>1</sup>H-<sup>1</sup>H COSY (Correlation SpectroscopY), DEPT 135° (Distortionless Enhancement by Polarization Transfer), <sup>1</sup>H-<sup>1</sup>H NOESY (Nuclear Overhauser Effect SpectroscopY) and <sup>1</sup>H-<sup>1</sup>H TOCSY (TOtal Correlation SpectroscopY), HMBC (<sup>1</sup>H-<sup>13</sup>C Heteronuclear Multiple Bond Correlation Spectroscopy) and HSQC (<sup>1</sup>H-<sup>13</sup>C Heteronuclear Single Quantum correlation). NOESY was used to determine the relative stereochemistry of the molecule while TOCSY was used to identify protons belonging the same spin system for the structure elucidation of glycosylated molecules (Agrawal, 1992; Reggelin *et al.*, 1992). Experiments were performed on either a Bruker AMX 600 or a Bruker AMX 300 instrument (Bruker, Germany). Chemical shifts ( $\delta$ ) are given in part by millions (ppm) and the coupling constants *J* in Hertz (Hz).

2.3.4.2 Ultraviolet-visible and Fourier transform infrared spectroscopies

Ultraviolet-visible (UV) and Fourier Transform Infrared (FT-IR) spectroscopies were recorded on Analytik Jena Specord 210 (Jena, Germany) and Agilent Cary 630 FT-IR (Agilent, UK) spectrophotometers respectively. Both methods are useful for the detection of functional groups in a molecule. Wavelengths scan for the UV was between 200-500 nm. Quartz cells were used for samples and blanks. The results were reported in nm as the wavelength of the maxima.

# 2.3.4.3 Optical rotation

Optical rotation was determined using Bellingham- Stanley ADP660 polarimeter with the sodium D line (589.3 nm) as the source of light. The measurements were made at 25 °C in a 10 cm cell. The results were calculated with the following formula:  $[\alpha]^{25}_{D}=1000\times\alpha/1.c$  (in which  $\alpha$ = observed rotation, 1 = cell length in dm and c = concentration in g/100 mL).

#### 2.3.4.4 Mass spectrometry

Mass spectrometry (MS) is a technique useful in determination of the molecular weight of compounds, the possible molecular formula and fragmentation pattern (Hoffmann *et al.*, 1996). High resolution mass spectra (HR MS) were recorded at the National Mass Spectrometry Facility (NMSF) (Swansea, UK) on a Xevo G2-S ASAP or LTQ Orbitrap XL1 spectrometers. Low and high resolution MS analyses were also performed at LJMU. HR MS was run on an Agilent 6200 Series Accurate-Mass Time-of-Flight (TOF) LC/MS using electro spray ionisation (ESI) in positive ion mode connected to an Agilent autosampler injection system. The analyte was prepared in MeOH and the mass spectrum was recorded relating to their mass to charge (*m*/*z*) at a capillary potential of 3,500 V. Low resolution analyses were performed on a Waters LCT Premier<sup>TM</sup> ESI-TOF Mass Spectrometer.

#### 2.4 Bioactivity studies

#### 2.4.1 Samples preparation for screening

Stock solutions of assayed samples were prepared in DMSO 100 mg/mL for crude extracts and fractions, and 100 mM for pure isolated compounds and stored at 4 °C. Prior to use, the stock solution was further diluted in the medium to final working concentrations ranging from 0-1000  $\mu$ g/mL for extract and, 0-250  $\mu$ M for pure compounds on the day of assay.

## 2.4.2 Cell culture

Dulbecco's Modified Eagles Medium (DMEM), Foetal bovine serum (FBS), penicillin/Streptomycin, and L-glutamine were warmed at 37 °C before making up a complete media. Penicillin/streptomycin (100 units/mL, 4 mL), L-glutamine (4 mL) and 50 mL of FBS were added into 500 mL of DMEM in a sterilised environment. The media was mixed and stored in a fridge at 4°C. Growth media was pre-warmed at 37°C for 20 min before each use. In the case of AREc2 cell line, 4 mL of 100 mg/mL of geneticin was also added to the growth medium. All cell lines were cultured at 37 °C in 95 % air and 5 % CO<sub>2</sub> in 25 or 75 cm<sup>2</sup> tissue culture flask (Corning, Staffordshire, UK) containing appropriate medium for the cell line.

When cells were 80-100 % confluent, they were passaged or used for the desired experiment. Firstly, the media was removed from the flasks and cells were washed with PBS twice to remove the residual medium and dead cells, as they are non-adherent to the flask surface. Small amount of pre-warmed trypsin-EDTA (0.2-0.5 mL) was added and incubated for 1-2 min at room temperature, then excess of trypsin was removed and the cells were incubated for an additional 2-4 min at 37 °C. The flasks were tapped gently to dislodge the cells and were visualised under a microscope.

After the cells were completely disassociated from the flask, the complete growth media was added into the flask. The medium was pipetted up and down several times to break up any cell clumps. The cell suspension was split into two fresh flasks and were labelled with new passage number and kept in an incubator at 37°C.

For the assays, after reaching 80-90% confluence, the cells were seeded into 96 well plates or 6 well plates at density  $1.2 \times 10^4$  in a corresponding working volume of 180  $\mu$ L/well and allowed to grow for 24 h before each experiment commenced.

87

#### 2.4.3 Cell viability assays

Trypan blue assay using a haemocytometer was used during cell counting to determine cell numbers for setting up assays. Equal amount of trypan blue and cell solution were mixed by pipetting up and down for few times, and 10  $\mu$ L of the obtained solution was transferred in each quadrant of the haemocytometer. Viable cells are typically round and refractile and exclude the stain, whilst non-viable cells are stained dark blue.

3-(4,5-dimethylthlthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay based on the method described by Mosmann (1983) was used to assess the cytotoxicity of extracts and pure isolated compounds on the different cell lines used in this study. The MTT is converted by the NADPH present in viable cells to a purple formazan crystal (Figure 2.12) which has an absorbance maximum near 570 nm. When cells died, they lose the ability to convert the MTT into formazan and therefore, colour formation serves as a useful convenient manner of viable cells only.

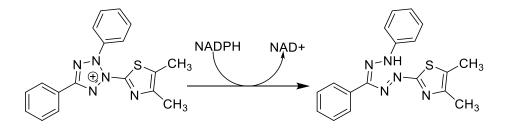


Figure 2.12 MTT transformation to formazan

Cells were seeded into 96 well plates at density  $1.2 \times 10^4$  cells/well. After 24 h incubation, the medium was removed and replaced with fresh medium containing assayed sample (extract or pure compound) at the desired concentration. The final concentration of DMSO in each well was no more than 0.1%. A set of untreated control cells was included in each experiment, as well as a set of cells treated with 0.1% DMSO. Following

incubation with the extracts for 24 h (48 h for pure compounds except with AREc32 cell line), 20  $\mu$ L of MTT solution (prepared in PBS at the concentration of 5 mg/mL) was added to each well. After 2 h of incubation at 37 °C, the medium was discarded and replaced with 100  $\mu$ L of DMSO. The OD<sub>570</sub> was determined with a microplate reader (CLARIO Star Microplate reader, BMG Labtech, UK). The percentage of cell viability was determined as percentage of control cells [(absorbance of treated cells/absorbance of untreated cells) × 100].

#### 2.4.4 Luciferase reporter gene assay

The ability of the extracts, fractions and isolated compounds to activate the expression of ARE driven genes which are commonly activated by Nrf2 were evaluated by cell-based luciferase assay with a cell line which contains an ARE-driven luciferase reporter gene (AREc32, modified MCF-7 cell line). Luciferases are a class of oxidative enzymes found in several species that enable the host to bioluminesce in the presence of luciferin (Figure 2.13). The emission can be read using a luminometer and is directly correlated to the level of ARE activation (Wang *et al.*, 2006).

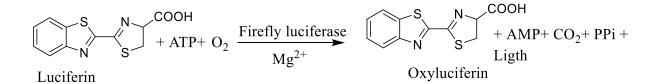


Figure 2.13 Formation of oxyluciferin

The Steady-Glo Luciferase assay kit from Promega was used for this purpose according to the manufacturer instructions. Briefly, AREc32 cells were seeded into 96 well plates at density  $1.2 \times 10^4$  cells/well and treated for 24 h with the concentration of extract or

pure compound causing no more than a 10% reduction of cell viability as determined by the MTT assay. Wells were then washed twice with 100  $\mu$ L of PBS and 20  $\mu$ L of luciferase reporter lysis buffer (Promega, USA) was added to each well followed by a freeze thaw cycle (24 h) to achieve complete cell lysis. The 20  $\mu$ L cell lysate was then aspirated and dispensed into a white opaque 96 well plates. 80  $\mu$ L of Luciferase reporter substrate was added to each well and the luminescence immediately measured using a plate reader (ClarioStar microplate reader, BMG Labtech, UK). Levels of luciferase activity were expressed relative to the basal level of luciferase activity in control (no treatment) and presented as a fold increase (relative to control).

### 2.4.5 Haem polymerisation assay

The potential antimalarial activity of plant extracts was evaluated by the method described by Hussain *et al.* (2011). Briefly, varying concentrations (0-2 mg/mL in 10% DMSO) of the extracts were incubated with 300  $\mu$ M of hematin (freshly dissolved in 0.1 M NaOH), 10 mM oleic acid and 10  $\mu$ M HCl. The reaction volume was adjusted to 1000  $\mu$ L using 500 mM sodium acetate buffer, pH 5. Chloroquine diphosphate was used as a positive control. The samples were incubated overnight at 37 °C with regular shaking. After incubation, samples were centrifuged (14,000 x g, 10 min, at 21 °C) and the hemozoin pellet repeatedly washed with sonication (30 min, at 21 °C; FS100 bath sonicator; Decon Ultrasonics Ltd.) in 2.5% (w/v) SDS in phosphate buffered saline followed by a final wash in 0.1 M sodium bicarbonate, pH 9.0, until the supernatant was clear. After the final wash, the supernatant was removed, and the pellets were resuspended in 1 mL of 0.1 M NaOH before determining the hemozoin content by measuring the absorbance at 405 nm using a 1 cm quartz cuvette. The results were recorded as % inhibition (I%) of haem polymerization/crystallization compared to

90

positive control (chloroquine) using the following formula:  $I\% = [(AB-AA)/AB] \times 100$ , where AB: absorbance of blank; AA: absorbance of test samples. All the solutions were pre-warmed at 60°C before initial mixing.

#### 2.5 Statistical analysis

All bioassay experiments were carried out in triplicate with a minimum of three replicates per assayed concentration. Data were expressed as means  $\pm$  SEM (standard error of mean). The graphs were plotted using Excel Microsoft office 2016. SEM is shown on the graph as an error bar. IC<sub>50</sub>, LC<sub>10</sub> and LC<sub>50</sub> where calculated using Graph Prism 7.0 software. *P* value were calculated using unpaired student *t*-test and considered significant when  $P \le 0.01$ .

# **Chapter 3 Results and Discussion**

# 3.1 Yield of extraction

Different parts of the following medicinal plants viz. *Croton oligandrus* Pierre ex Hutch, *Ruspolia hypocrateriformis* (Vahl) Milne-Redh, *Pseudospondias microcarpa* (A. Rich.) Engl., *Zanthoxylum lepreurii* Guill. and Perr. and *Zanthoxylum zanthoxyloides* (Lam.) Zepern. and Timler (Figure 3.1) were collected, dried at room temperature, ground and successively extracted with a Soxhlet apparatus using *n*-hexane, DCM and MeOH. Eighteen extracts were obtained (Table 3.1). As the three solvents used have different polarity index, the constituents of the plant were gradually extracted according to their affinity with each solvent. The highest yield percentage of extraction was that of *Z. zanthoxyloides* with 23.9 % recovery percentage while *P. microcarpa* bark yielded the lowest extraction percentage of 3.3 %.



Figure 3.1 Pictures showing the different plant parts collected

a) *Croton oligandrus* Pierre ex Hutch stem bark; b) *Ruspolia hypocrateriformis* Vahl Milne-Redh leaves; c) *Pseudospondias microcarpa* (A. Rich.) Engl. stem bark; d) *Pseudospondias microcarpa* (A. Rich.) Engl. leaves; e) *Zanthoxylum lepreurii* Guill. and Perr. fruits; f) *Zanthoxylum zanthoxyloides* (Lam.) Zepern. and Timler fruits.

Plants	Powder weight (g)	Extra	ct weig	Yield (%)	
		Hexane	DCM	MeOH	
C. oligandrus bark	330.8	3.7	2.1	7.4	4.0
R. hypocrateriformis leaves	450.0	13.6	10.2	41.8	14.6
<i>P. microcarpa</i> bark	276.6	2.2	1.1	5.9	3.3
P. microcarpa leaves	437.3	13.4	2.7	12.5	6.5
Z. leprieurii fruits	546.8	32.7	21.1	54.3	19.8
Z. zanthoxyloides fruits	350.0	59.58	5.9	18.25	23.9

Table 3.1 Yield of extraction of the different plants collected

## **3.2 Preliminary screening**

## **3.2.1 Luciferase activity**

The extracts obtained from the collected plant materials were screened for their potential cancer chemopreventive properties using the AREc32 cell line. AREc32 cells are stably transformed variant of the MCF-7 cells containing a luciferase reporter gene under the control of antioxidant promoters, and therefore, is an excellent cell model for the identification of potent activator of the Nrf2 signalling pathway (Wang *et al.*, 2006).

## 3.2.1.1 AREc32 cells viability

The cell viability of the AREc32 cells in the presence of each extract was first assessed by the MTT assay with the aim of determing the threshold dose nontoxic for the cells and the suitable dosage for the luciferase assay. The cells were incubated in the presence of the extracts and the effect of those extracts on cell viability measured after 24 h and compared to that of control (untreated cells). Results are shown on Figure 3.2. The DCM Extract of *P. microcarpa* bark was found to be the most toxic to the AREc32 cells killing nearly 50% of cells at 62.5  $\mu$ g/mL while the MeOH Extract of *Z. zanthoxyloides* fruits was found to be the least toxic with only 70% of cells death occurring at 1 mg/mL. JHL and PML were found to have a cytotoxic effect on AREc32 cells at very low concentrations. The concentration causing no more than 10% cells death (LC<sub>10</sub>), was chosen as the assayed concentration for the luciferase assay (Table 3.2) as no significant cell death was observed below that concentration.

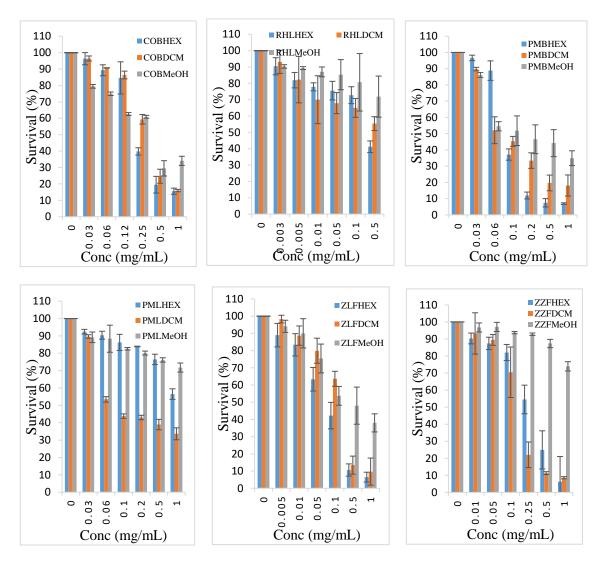


Figure 3.2 Effect of tested extracts on the viability of AREc32 cells.

Cells were treated for 24 h with 0.1% DMSO (control), or with various concentrations (0 – 1 mg/mL) of extracts. Cell viability was determined by MTT assay. Each measured parameter is plotted as mean ± SEM of three independent experiments (5 replicates/ experiment). COB: *C. oligandrus* bark; RHL: *R. hypocrateriformis* leaves; PMB: *P. microcarpa* bark; PML: *P. microcarpa* leaves; ZLF: *Z. leprieurii* fruits; ZZF: *Z. zanthoxyloides* fruits, HEX: *n*-Hexane extract; DCM: DCM extract; MeOH: MeOH extract.

Extract	Least toxic concentration (LC10 in µg/mL)						
	COB	JHL	PMB	PML	ZLF	ZZF	
HEX	60	4	60	60	4	5	
DCM	60	3	30	30	5	50	
МеОН	10	5	30	30	5	500	

Table 3.2 Concentrations of extracts causing no more than 10% toxicity

Cells were treated for 24 h with various concentrations (0 - 1 mg/mL) of extracts. Cell viability was determined by MTT assay. LC<sub>10</sub> was determined using GraphPad Prism software of three independent experiments (5 replicates/ experiment). COB: *C. oligandrus* bark; RHL: *R. hypocrateriformis* leaves; PMB: *P. microcarpa* bark; PML: *P. microcarpa* leaves; ZLF: *Z. leprieurii* fruits; ZZF: *Z. zanthoxyloides* fruits, HEX: *n*-Hexane extract; DCM: DCM extract; MeOH: MeOH extract.

### 3.2.1.2 Luciferase assay

Luciferase assay was carried out as describe in the experimental section (Chapter 2, 2.4.4). The results are shown on Figure 3.3. Among the tested extracts, COBHEX, COBDCM, ZZFDCM and ZZFMeOH extracts were found to induce luciferase activity. ZZFMeOH was the most active with a 36-fold induction follow by its DCM extract with 33-fold induction. The activities of *Z. zanthoxyloides* DCM and MeOH extracts were similar to that of *t*BHQ (50  $\mu$ M) used as positive control. *t*BHQ is a well characterized chemopreventive compound which exert its activity by inducing Nrf2-regulated genes (Wang *et al.*, 2006). The fact that only DCM and MeOH extracts induced high levels of the luciferase activity (greater than 30-fold) might be linked to presence of polar compounds in the plant.

Several *Zanthoxylum* species are traditionally used in Asia and Central America to treat and prevent cancer (Epifano *et al.*, 2011). Phytochemical studies of *Z. zanthoxyloides* have led to the isolation of alkaloids with good cytotoxicity against a panel of cancer cell lines (Wouatsa *et al.*, 2013). It is interesting to note that *Z. leprieuri* extracts did not show any significant increase of the luciferase activity compared to the closely related species *Z. zanthoxyloides*. A comparative study carried out by Wouatsa *et al.* (2013a) has revealed that the two plants only have few phytochemicals in common.

In contrast to *Z. zanthoxyloides*, for which polar extracts were active, only nonpolar and medium polar extracts of *C. oligandrum* showed good luciferase induction. COBHEX and COBDCM extracts were found to induce the luciferase activity by 18 and 21-fold, respectively. *Croton* is a rich source of diterpenes (Xu *et al.*, 2018), and some isolated diterpenes from plants have been reported to have cancer chemopreventive activity (Dhanasekaran *et al.*, 2009; Endringer *et al.*, 2014).

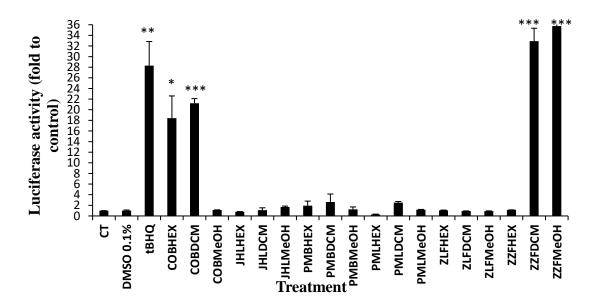


Figure 3.3 Luciferase activity of the screened extracts

AREc32 cells were seeded in 96-well plates at  $1.2 \times 10^4$  cells/well. After 24 h, *t*BHQ (50  $\mu$ M) and the extracts were added to the medium. The cells were then incubated for another 24 h and assayed for luciferase activity as detailed in Chapter 2, 2.4.4. The value of luciferase activity of untreated cells (CT) was set at 1. Values shown are mean  $\pm$  SEM of three experiments. \*p < 0.1; \*\*p < 0.01, \*\*\*p < 0.001 significantly increased versus control. DMSO 0.1%: (cells treated with 0.1% DMSO medium).

The most active crude extracts were fractionated using VLC for *n*-hexane and DCM extracts, and SPE for MeOH extract. For each *n*-hexane or DCM extract, six fractions were obtained and four fractions only for MeOH extract. The effect of the fractions on

the viability of AREc32 cells was assessed by the MTT assay to determine their least toxic concentration (Concentrations causing no more than 10% cells death), and the obtained concentration (Table 3.3) was used as dosage to evaluate the luciferase activity of the fractions.

Extract	Least toxic concentration (LC <sub>10</sub> in µg/mL)						
	F1	F2	F3	F4	F5	F6	
COBHEX	5	5	10	1	0.5	1	
COBDCM	5	10	0.5	5	0.5	5	
ZZFDCM	0.5	1	1	5	5	0.1	
ZZFMeOH	5	1	0.5	5	-	-	

Table 3.3 Concentrations of fractions causing no more than 10% toxicity

Cells were treated for 24 h with various concentrations (0 - 1 mg/mL) of extracts. Cell viability was determined by MTT assay. LC<sub>10</sub> was determined using GraphPad Prism software of three independent experiments (5 replicates/ experiment). F1-F6 are fractions obtained from VLC and SPE of active crude extracts COB: *C. oligandrus* bark; ZZF: *Z. zanthoxyloides* fruits, HEX: *n*-Hexane extract; DCM: DCM extract; MeOH: MeOH extract.

The level of induction of the luciferase activity by each fraction tested is depicted in Figure 3.4. Overall, fractionation considerably reduced the activity, as most of the fractions did not show significant fold induction of the luciferase activity as the mother crude extract. The only plausible explanation could be that the activities observed from the corresponding crude extract were due to synergistic action of the different compounds present in the plant. In addition, the fractionation techniques used, VLC and SPE separated the constituents of the plant based on their affinity with the eluent used. Thus, each fraction is a mixture of several compounds with similar polarity in different proportions conferring a unique phytochemical profile. Moreover, plant extracts contain complex mixtures of compounds in which most of the time a single constituent is not responsible for the overall activity (Williamson, 2001).

Among the tested fractions, COBHEX F1 and ZZFMeOH F3 showed induction of 3.24and 2.4-fold, respectively. The activity of ZZFMeOH F3 was comparable to that of *t*BHQ 6  $\mu$ M (2.5- fold induction) while COBHEX F1 was more active than *t*BHQ 6  $\mu$ M. The remaining fractions showed fold induction below 2. According to McMahon and coworkers (2014), chemicals causing fold increase of luciferase activity greater than 2 are considered as active.

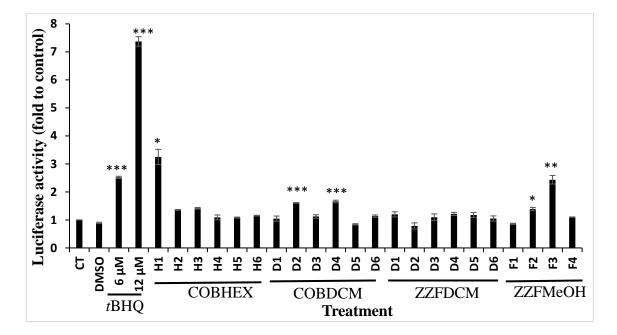


Figure 3.4 Luciferase activity of the fractions

AREc32 cells were seeded in 96-well plates at  $1.2 \times 10^4$  cells/well. After 24 h, *t*BHQ (6 and 12  $\mu$ M) and the fractions were added to the medium. The cells were then incubated for another 24 h and assayed for luciferase activity as detailed in Chapter 2, 2.4.4. The value of luciferase activity of untreated cells (CT) was set at 1. Values shown are mean  $\pm$  SEM of three experiments. \*p < 0.01; \*\*p < 0.001, \*\*\*p < 0.0001 significantly increased versus control. DMSO, cells treated with 0.1% DMSO medium.

## 3.2.2 Haem polymerisation assay

Hemozoin, also called malaria pigment, is essential for the survival of the malaria parasite in red blood cells (Coronado *et al.*, 2011). The pigment is a polymer of haem units, which is biocrystallised during haem detoxification in the erythrocytic life of the parasite. Inhibition of the haem polymerisation is therefore targeted for the search of antimalarial substances. Several screening techniques have been developed for the purpose. In the present study, we have performed the assay using the method published by Hussain *et al.* (2011) as described in the experimental section. Chloroquine diphosphate was used as positive control. Results are shown in Table 3.4.

Extract	IC <sub>50</sub> ( $\mu$ g/mL) ± SEM					
	HEX	DCM	МеОН			
СОВ	$180.0\pm6.0$	$164.8 \pm 53.0$	>1			
JHL	$206.7 \pm 52.0$	>1	$170.3 \pm 77.9$			
РМВ	$73.9\pm25.8$	2.5 ± 1.5	4.0 ± 1.7			
PML	> 1	>1	$13.0 \pm 9.0$			
ZLF	> 1	$45.8 \pm 25.0$	> 1			
ZZF	> 1	>1	> 1			
Chloroquine	$0.43\pm0.08$					

Table 3.4 Haem polymerisation inhibitory concentration (IC<sub>50</sub> in  $\mu$ g/mL)

IC<sub>50</sub> was determined using GraphPad Prism of three independent experiments (3 replicates/experiment). COB: *C. oligandrus* bark; RHL: *R. hypocrateriformis* leaves; PMB: *P. microcarpa* bark; PML: *P. microcarpa* leaves; ZLF: *Z. leprieurii* fruits; ZZF: *Z. zanthoxyloides* fruits, HEX: *n*-Hexane extract; DCM: DCM extract; MeOH: MeOH extract.

Among the five plants screened, *P. microcarpa* extracts were found to be the most active. Inhibition of the haem polymerisation was observed for the DCM and MeOH extracts of the stem bark with IC<sub>50</sub> values  $2.5 \pm 1.5$  and  $4.0 \pm 1.7 \,\mu$ g/mL, respectively. MeOH extract produced 50% inhibition at 13.0  $\mu$ g/mL. The PMB hexane extract was less active (IC<sub>50</sub>  $73.9 \pm 25.8 \,\mu$ g/mL), while PML hexane and MeOH were found to have no effect at the highest concentration tested. The IC<sub>50</sub> obtained for PMB DCM and MeOH extracts were comparable to the result obtained when the stem bark ethanol extract (IC<sub>50</sub> 1.13 $\pm$  0.16  $\mu$ g/mL) was screened *in vitro* against multi-drug resistant *Plasmodium falciparum* K1 strain (Malebo *et al.*, 2009). The IC<sub>50</sub> of the MeOH extract was close to the result published by Mbatchi et *al.* (2006) (IC<sub>50</sub> 26  $\pm$  10 µg/mL) when the ethanol extract was tested in *vitro* against human erythrocytes infected by FcM29 originated from Cameroon. *Z. zanthoxyloides* fruits extracts were found to be not active at the highest concentration tested. However, previous screenings have demonstrated a moderate antimalarial activity for the root and trunk bark extracts (Kassim *et al.*, 2005; Gansane *et al.*, 2010).

This is the first report on the evaluation of the antimalarial activity *C. oligandrus* and *J. hypocrateriformis*. Overall, extracts with  $IC_{50} < 200 \ \mu g/mL$ , i.e. COBHEX, COBDCM, JHLMeOH, PMBHEX, PMBDCM, PMBMeOH, PMLMeOH and ZLFDCM were selected for further studies.

#### 3.3 Characterisation and structure elucidation of isolated compounds

#### **3.3.1 Phytochemistry of** *Croton oligandrus*

The *n*-hexane and DCM extracts of *C. oligandrus* exhibited good antimalarial and chemopreventive properties. CC and RP-HPLC analyses of the *n*-hexane and DCM extracts led to the isolation and characterization of eighteen compounds, which were identified as  $3-\beta$ -*O*-acetyl aleuritolic acid (7), lupeol (8), vanillin (147), *trans*-ferulic acid (148), a mixture of ferulate derivatives: cluytyl ferulate (149) and hexacosanoyl ferulate (150), crotocorylifuran (151), crotonzambefuran A (152), megalocarpoidolide D (153), 12-*epi*-megalocarpoloide D (154), crotonolin E (155), the epimeric mixtures of crotonolin A (156) and crotonolin B (157), and crotonolin C (158) and crotonolin D (159), crotonolin F (160), 7-*a*-hydroxydehydroabietic acid (161) and 7-oxo-dehydroabietic acid (162) (Figure 3.5). This is the first report on the occurrence of the compounds 147-162 in *C. oligandrus*. Compounds 154-160 are new clerodane diterpenes from any natural

source. Diterpenes are the most abundant secondary metabolites found in the genus *Croton*, representing about 85% of all the isolated metabolites with clerodane being the most abundant subclass (Xu *et al.*, 2018).

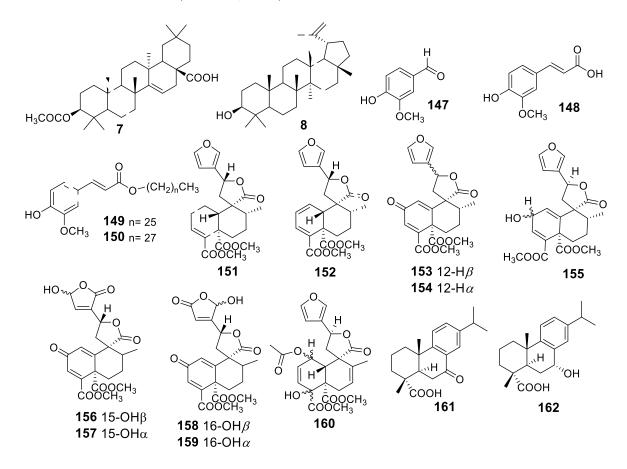


Figure 3.5 Structures of isolated compounds from the bark of C. oligandrus

3.3.1.1 Structure elucidation of  $3-\beta$ -O-acetyl aleuritolic acid (7)

Compound **7** was obtained as a white powder, which precipitated in a mixture of Hex-EtOAc (7:1). Its molecular formula  $C_{32}H_{50}O_4$  was deduced from the ESI-MS spectrum in positive ion mode (Figure 3.6) by the sodiated molecular ion peak at m/z 521 [M+Na]<sup>+</sup>. Its <sup>1</sup>H NMR spectrum (Figure 3.7, Table 3.5), showed seven methyl singlets at  $\delta_H$  0.88, 0.90, 0.96, 0.97, 0.97, 0.98, 0.99, 1.95 and 2.06; a doublet of doublets at  $\delta_H$  5.52 (J = 3.4, 7.9 Hz) indicating the presence of an olefinic proton (Misra and Khastgir, 1970) and

another doublet of doublets at  $\delta$  4.48 (J = 5.3, 10.9 Hz), attributable to an oxymethine proton. All these preliminary observations suggested the presence of a triterpenoid skeleton in the molecule. This was further confirmed by the presence of thirty-two carbon signals in the <sup>13</sup>C NMR spectrum of 7 (Figure 3.8, Table 3.5) corresponding to eight methyls, ten methylenes, five methines including an oxymethine at  $\delta_{\rm C}$  80.9 (C-3) and an olefinic methine at  $\delta_{\rm C}$  116.7 (C-15), nine quaternary carbons including an olefinic carbon at  $\delta_{\rm C}$  160.6 (C-14) and two carbonyls signals at  $\delta_{\rm C}$  184.0 and 170.9 indicating the presence of a carboxylic acid and an acetoxy function, respectively. The signals at  $\delta_{C}$  160.5 and 116.8 could be assigned to the double-bonded carbons (C=C), which are typical of that of C14-C15 double bond signals of a taraxerane skeleton (Rasool et al., 1989). The position of the double bond could be confirmed by the different <sup>1</sup>H-<sup>13</sup>C long-range correlations (Figure 3.8) observed in the HMBC spectrum. The methyl signal at  $\delta_{\rm H}$  2.06 was attributed to the methyl of the acetoxy group based on its correlation to the carbonyl at  $\delta_{\rm C}$  170.9 observed in the HMBC. The acetoxy group could be placed on C-3 of the triterpene skeleton as there was a  ${}^{3}J$  correlation from the 3-oxymethine ( $\delta_{\rm H}$  4.48) to the acetoxy carbonyl ( $\delta_{\rm C}$  170.9), observed in the HMBC spectrum. The assignment of all <sup>1</sup>H and <sup>13</sup>C NMR signals (Table 3.5) of **7** was achieved by a combination of COSY, HSQC and HMBC spectral analyses, and in comparison, with the published data for  $3-\beta$ -O-acetyl aleuritolic acid (Mahato and Kundu, 1994; Prabowo et al., 2013). Thus, compound 7 was identified conclusively as  $3-\beta$ -O-acetyl aleuritolic acid. Acetyl aleuritolic acid (7) was first isolated from the bark of Aleurites montana (Euphorbiaceae) and is widespread in all genera of the Euphorbiaceae including the genus Croton (Misra and Khastgir, 1970; Salatino *et al.*, 2007).

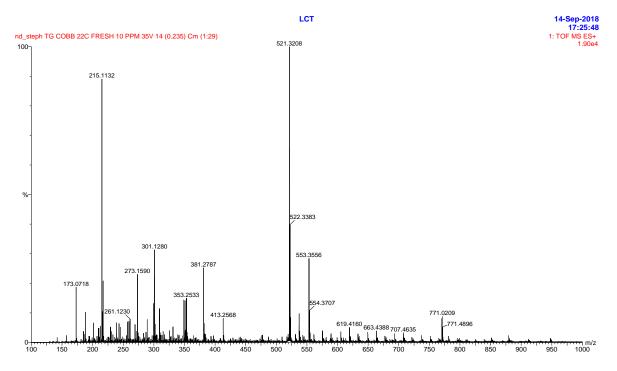


Figure 3.6 ESI-MS spectrum of 7

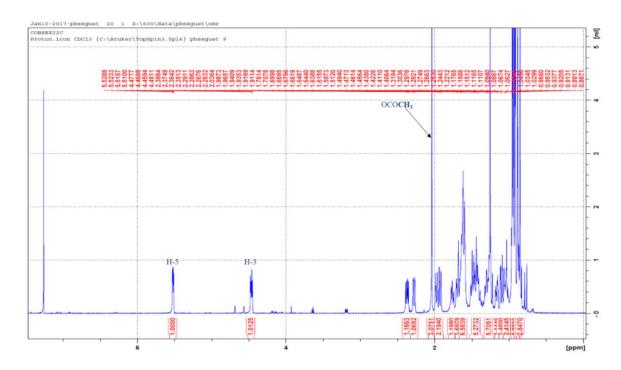


Figure 3.7  $^{1}$ H NMR (600 MHz, CDCl<sub>3</sub>) of **7** 

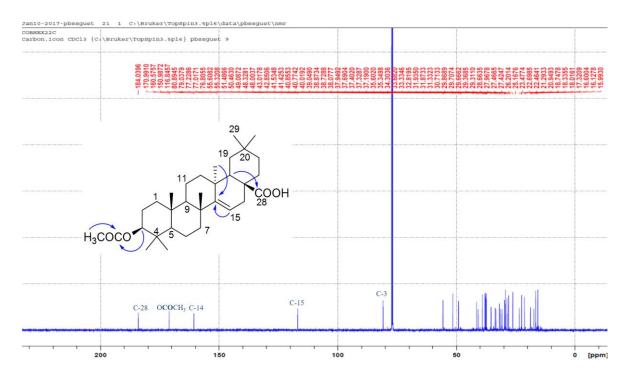


Figure 3.8 <sup>13</sup>C NMR (150 MHz, CDCl<sub>3</sub>) and key HMBC correlations of 7

Position	$\delta_{\rm H} { m m} (J { m in}$	$\delta_{\rm C}$	Position	$\delta_H m (J$	$\delta_{\rm C}$	Position	δ <sub>H</sub> m (J	$\delta_{\rm C}$
	Hz)			in Hz)			in Hz)	
1	1.05	37.7	11	1.52	17.3	20	29.7	-
	1.65			1.67		21	33.6	1.09
2	1.65	23.4	12	1.65	33.3	22	1.43	30.7
3	4.48 dd	80.9		1.76			1.71	
	(5.3,							
	10.9)							
4	-	37.4	13	-	37.3	23	0.88 s	27.9
5	0.92	55.6	14	-	160.6	24	0.99 s	16.5
6	1.20	18.7	15	5.52 dd	116.7	25	0.96 s	15.6
				(3.4, 7.9)				
	1.65		16	1.95	31.4	26	0.98 s	26.1
7	1.32	40.8		2.39		27	0.97 s	22.4
	2.00		17	-	51.3	28	-	184.0
8	-	39.0	18	2.29	41.5	29	0.97 s	31.9
9	1.45	49.0	19	1.14	35.3	30	0.90 s	28.7
10	-	37.9		1.30				
CH <sub>3</sub> CO	-	170.9	OCCH <sub>3</sub>	2.06 s	21.3			

Table 3.5  $^1\text{H}$  (600 MHz, CDCl\_3) and  $^{13}\text{C}$  NMR (150 MHz, CDCl\_3) data of 7

3.3.1.2 Structure elucidation of cluytyl ferulate (**149**) and hexacosanoyl ferulate (**150**) as a mixture

Compounds 149 and 150 were obtained as a mixture and as a white powder. These were identified as the main constituents of the *n*-hexane extract along with acetyl aleuritolic acid (7) and lupeol (8). The HR ESI-MS spectrum in negative ion mode (Figure 3.9) of the white powder revealed a mixture composition of two homologous substances showing [M-H] peaks at m/z 557.4571 and 585.4881, with a relative abundance of 52 and 45%, respectively, corresponding to the formulas C<sub>36</sub>H<sub>61</sub>O<sub>4</sub> and C<sub>38</sub>H<sub>65</sub>O<sub>4</sub>, respectively, calculated for  $[C_{36}H_{61}O_4]^-$ , 557.4575 and  $[C_{38}H_{65}O_4]^-$ , 585.4888, respectively. The <sup>1</sup>H NMR spectrum (Figure 3.10, Table 3.6) exhibited signals for a methoxy group ( $\delta_{\rm H}$  3.95, s), two *trans* olefinic protons ( $\delta_{\rm H}$  7.63 and 6.31, J =16.0 Hz), and three aromatic hydrogens forming an ABX system ( $\delta_{\rm H}$  6.93, d, J = 8.1 Hz; 7.09, d, J = 1.8 Hz and 7.10 , dd, J = 1.8, 8.1 Hz) that could be assigned to a ferulic acid moiety (Balde *et al.*, 1991). The signals at  $\delta_{\rm H}$  4.21 (2H, t, J = 6.7 Hz), 1.27-1.71 (2H, brs, m) and 0.90 (3H, t, J = 6.9Hz) could be attributed to an oxymethylene, a long saturated aliphatic chain and a terminal methyl functionality, respectively, suggesting the presence an aliphatic alcohol moiety in the molecule. The <sup>13</sup>C NMR spectrum (Figure 3.11, Table 3.6) confirmed the presence of a saturated fatty alcohol moiety, an ester linkage and a ferulic moiety. Assignment of the <sup>1</sup>H and <sup>13</sup>C NMR data of **149** and **150** (Table 3.6) were confirmed by the <sup>1</sup>H-<sup>13</sup>C longrange correlations (Figure 3.11) observed in the HMBC spectrum. All assignments of <sup>1</sup>H and <sup>13</sup>C NMR signals of these compounds are in good agreement with those published for cluytyl ferulate and hexacosenoyl ferulate, respectively (Wandji et al., 1990; Balde et al., 1991). Thus, compounds 149 and 150 were identified as cluytyl ferulate (149) and hexacosenovl ferulate (150), respectively.

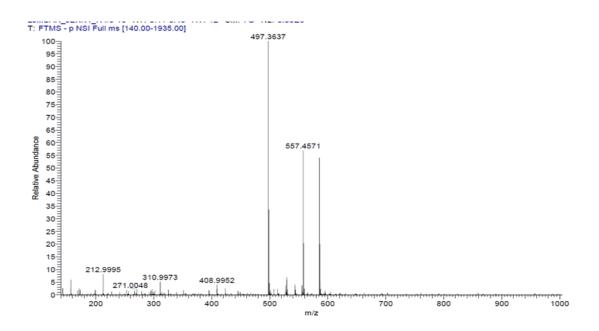


Figure 3.9 ESI-MS spectrum of 149 and 150

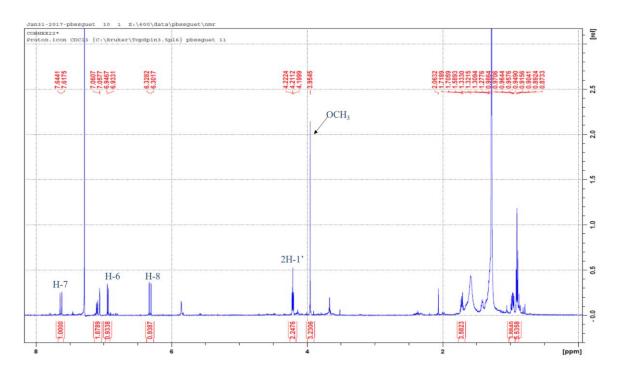


Figure 3.10  $^{1}$ H NMR (600 MHz, CDCl<sub>3</sub>) of **149** and **150** 

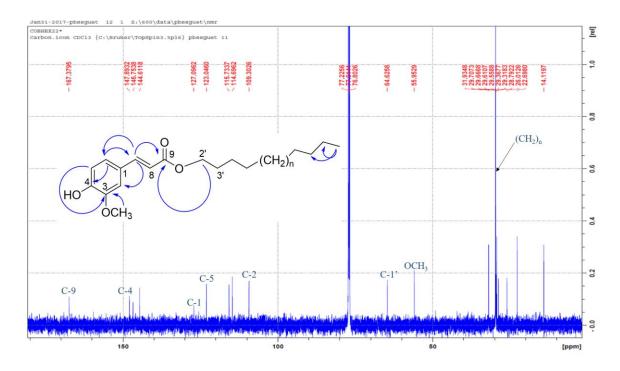


Figure 3.11  $^{13}$ C NMR (150 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **149** and **150** 

$\delta_{\rm H}$ m ( <i>J</i> in Hz)	$\delta_{C}$	Position	$\delta_{\rm H}$ m ( <i>J</i> in Hz)	$\delta_{C}$
-	127.1	1'	4.20 t (6.7)	64.6
7.06 d ( <i>1.8</i> )	109.3	2'	1.42 m	26.0
-	146.7	3'	1.71 m	28.8
-	147.9	4'-23'/25'	1.27 brs	29.3
7.10 dd ( <i>1.8</i> , <i>8.1</i> )	123.0	24'/26'	1.27 brs	31.9
6.93 d (8.1)	114.7	25'/27'	1.27 brs	22.7
7.63 d ( <i>16.0</i> )	144.6	26'/28'	0.90 t (6.9)	14.1
6.31 d ( <i>16.0</i> )	115.7	OCH <sub>3</sub>	3.95 s	55.9
-	167.4			
	- 7.06 d (1.8) - - 7.10 dd (1.8, 8.1) 6.93 d (8.1) 7.63 d (16.0)	-       127.1         7.06 d (1.8)       109.3         -       146.7         -       147.9         7.10 dd (1.8, 8.1)       123.0         6.93 d (8.1)       114.7         7.63 d (16.0)       144.6         6.31 d (16.0)       115.7	-       127.1       1'         7.06 d (1.8)       109.3       2'         -       146.7       3'         -       147.9       4'-23'/25'         7.10 dd (1.8, 8.1)       123.0       24'/26'         6.93 d (8.1)       114.7       25'/27'         7.63 d (16.0)       144.6       26'/28'         6.31 d (16.0)       115.7       OCH <sub>3</sub>	-       127.1       1'       4.20 t (6.7)         7.06 d (1.8)       109.3       2'       1.42 m         -       146.7       3'       1.71 m         -       147.9       4'-23'/25'       1.27 brs         7.10 dd (1.8, 8.1)       123.0       24'/26'       1.27 brs         6.93 d (8.1)       114.7       25'/27'       1.27 brs         7.63 d (16.0)       144.6       26'/28'       0.90 t (6.9)         6.31 d (16.0)       115.7       OCH <sub>3</sub> 3.95 s

Table 3.6  $^{1}$ H (600 MHz, CDCl<sub>3</sub>) and  $^{13}$ C NMR (150 MHz, CDCl<sub>3</sub>) data of **149** and **150** 

#### 3.3.1.3 Structure elucidation of crotocorylifuran (151)

Compound 151 was obtained as a white amorphous powder. Its molecular formula  $C_{22}H_{26}O_7$  was determined by ESI-MS analysis (Figure 3.12) in positive ion mode, where the sodiated ion peak was observed at m/z 425 [M+Na]<sup>+</sup>. Its <sup>1</sup>H NMR spectrum (Figure 3.13, Table 3.7) indicated the presence of a secondary methyl group at  $\delta_{\rm H}$  1.02 (3H, d, J = 6.4 Hz), two methoxy singlet functions at  $\delta_{\rm H}$  3.71 and 3.76, characteristic signals of a  $\beta$ -substituted furan ring at  $\delta_{\rm H}$  6.41 (1H, brd, J = 0.9 Hz), 7.44 (1H, ov) and 7.45 (1H, brs) (Tchissambou *et al.*, 1990), an oxymethine proton at  $\delta_{\rm H}$  5.40 (1H, t, J = 8.5 Hz) and an olefinic proton belonging to a trisubstituted double bond at  $\delta_{\rm H}$  6.84 (1H, dd, J = 4.3, 7.5Hz) were also observed. The <sup>13</sup>C NMR spectrum (Figure 3.14, Table 3.8) of **151** showed 22 carbon resonances suggesting a diterpene skeleton. The <sup>13</sup>C NMR and HSQC DEPT analyses supported the observations made on the <sup>1</sup>H NMR spectrum and also indicated the presence of five methylenes at  $\delta_{C}$  19.2 (C-1), 26.4 (C-2), 32.3 (C-6), 27.9 (C-7) and 42.4 (C-11) and two methines at  $\delta_{\rm C}$  40.2 (C-8) and 51.9 (C-10), which on the HSQC spectrum correlated with the proton signals appearing at  $\delta_{\rm H}$  1.58 (m) and 1.76 (dd, J =2.4, 13.2 Hz), respectively. All these data together with the different correlations observed in the COSY and HMBC spectra suggested a neo-clerodane (ex ent-clerodane)-type diterpene skeleton (Li et al., 2016). The <sup>1</sup>H and <sup>13</sup>C NMR data (Tables 3.7-3.8) of **151** were similar to those of crotocorylifuran (Tchissambou et al., 1990). The absolute configuration of the C-12 carbon was determined using the NOESY experiment. For a C-12R configuration, a correlation would be seen between H-12 and 3H-17, while a correlation seen between H-12 and H-1 $\alpha$  will imply C-12S configuration (Bautista *et al.*, 2012; Ndunda et al., 2016). For compound 151, a correlation observed in the NOESY spectrum between the H-12 and H-1 $\alpha$  proton ( $\delta_{\rm H}$  1.92) suggested a C-12S configuration. All the <sup>1</sup>H and <sup>13</sup>C NMR data of **151** supported by the <sup>1</sup>H-<sup>13</sup>C long-range correlations observed (Figure 3.14) led to its identification as crotocorylifuran, a clerodane diterpene first isolated from the trunk bark of *Croton haumanianus* (Tchissambou *et al.*, 1990).

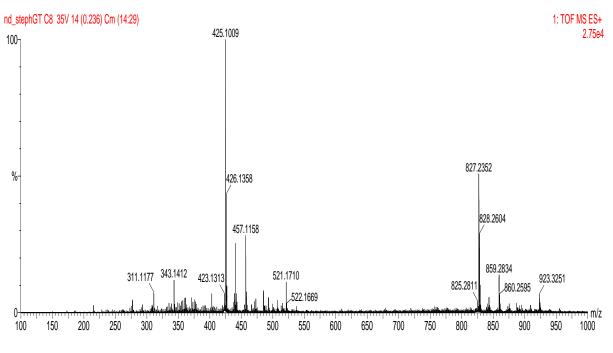


Figure 3.12 ESI-MS spectrum of 151

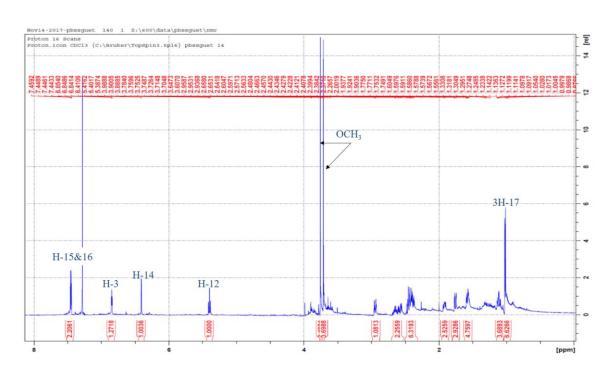


Figure 3.13 <sup>1</sup>H NMR (600 MHz, CDCl<sub>3</sub>) of compound 151

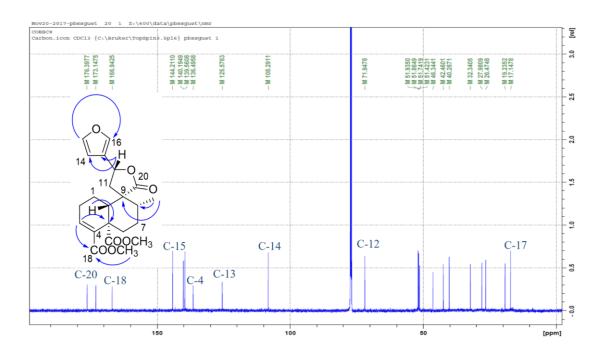


Figure 3.14<sup>13</sup>C NMR (150 MHz, CDCl<sub>3</sub>) and key HMBC correlations of 151

# 3.3.1.4 Structure elucidation of megalocarpoidolide D (153)

Compound **153** was isolated as a white amorphous powder,  $[\alpha]_D^{25}$ -88.8 (c 0.001, MeOH). Its molecular formula C<sub>22</sub>H<sub>22</sub>O<sub>8</sub> was determined from the *pseudo*molecular ion peak at m/z 437 [M+Na]<sup>+</sup> from its ESI-MS spectrum (Figure 3.15) obtained in positive ion mode. The IR spectrum of **153** displayed absorption stretching bands at 1715, 1767 and 1663 cm<sup>-1</sup> that could be attributed to the carbonyl of ester, lactone and ketone groups, respectively. The <sup>1</sup>H NMR spectrum (Figure 3.16, Table 3.7) revealed characteristic signals of a  $\beta$ -substituted furan ring at  $\delta_H$  6.45 (m, H-14), 7.47 (br dd, J = 1.5, 3.1 Hz, H-15) and 7.54 (brs, H-16), two olefinic protons at  $\delta_H$  6.47 (d, J = 1.3 Hz) and 6.78 (d, J = 1.3 Hz), an oxymethine at  $\delta_H$  5.55 (dd, J = 5.2, 11.1 Hz), as well as three methyl signals including two methoxy singlets at  $\delta_H$  3.65 and 3.84 and a doublet methyl at  $\delta_H$  1.17 (d, J = 6.4 Hz). Its <sup>13</sup>C NMR spectrum (Figure 3.17, Table 3.8) exhibited the resonances of 22 carbons suggesting a diterpene skeleton. The carbon signals appearing at  $\delta_C$  108.1 (C-14),

123.5 (C-13), 140.5 (C-16) and 144.6 (C-15) and correlating on the HSQC spectrum (Figure 3.18) with the protons at  $\delta_{\rm H}$  6.47 (d, J = 1.3 Hz) and 6.78 (d, J = 1.3 Hz) could be attributed to those of the substituted furan ring. One oxymethine at  $\delta_C$  71.2 as well as seven quaternary carbons including three carbonyl ester groups at  $\delta_{\rm C}$  165.3, 166.3 and 172.1 and a ketone carbonyl at  $\delta_{\rm C}$  185.7 were also observed in the <sup>13</sup>C NMR spectrum. Assignment of the <sup>1</sup>H and <sup>13</sup>C NMR data of **153** supported by its 2D NMR correlations (Figures 3.18-3.21) suggested an ent-clerodane diterpene skeleton (Li et al., 2016). The HMBC spectrum revealed that the two carbonyls at  $\delta_{C}$  165.3 and 166.3 correlating with the methoxy signals at  $\delta_{\rm H}$  3.84 and 3.65, respectively, could be at C-18 and C-19, while the carbonyl at  $\delta_{\rm C}$  172.1 could be attributed to that of a lactone moiety and assigned the position C-20. The other carbonyl at  $\delta_{\rm C}$  185.8 could be identified to that of a conjugated carbonyl and assigned to position C-2 by comparison of the NMR data of 153 with those published for megalocarpoidolide D (Ndunda et al., 2016). All the <sup>1</sup>H and <sup>13</sup>C NMR data of **153** and the <sup>1</sup>H-<sup>13</sup>C long-range correlations (Figure 3.16) were in good agreement with those of the  $\beta$ -substituted furancelerodane, megalocarpoidolide D, isolated from the roots of Croton megalacarpoides (Ndunda et al., 2016). Furthermore, analysis of the NOESY spectrum (Figure 3.20) showed a strong correlation between H-12 and H-1 confirming the  $\beta$ -orientation of H-12 (Bautista *et al.*, 2012). Thus, compound **153** was identified as megalocarpoidolide D.

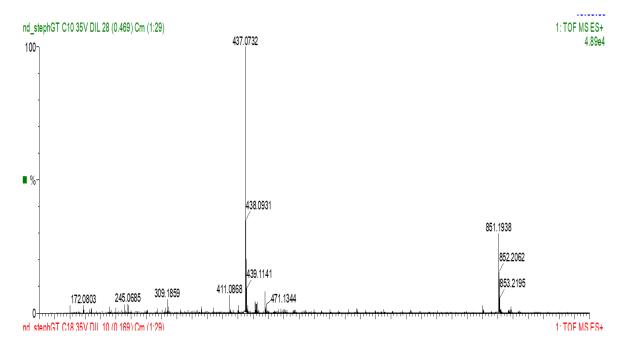


Figure 3.15 ESI-MS spectrum of 153

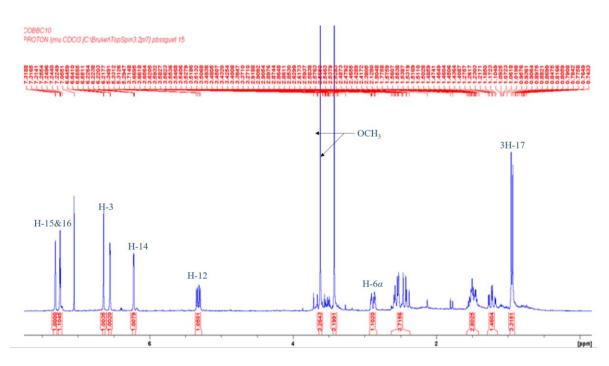


Figure 3.16 <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>) of **153** 

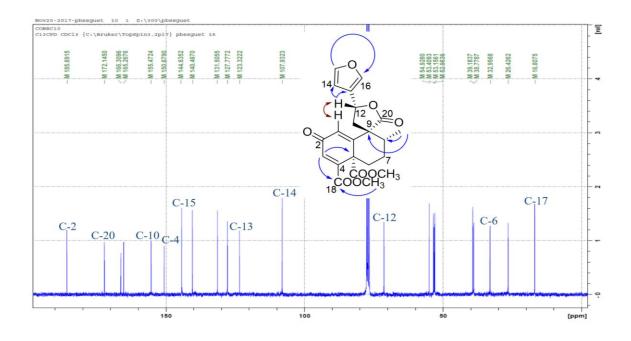


Figure 3.17 <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) and key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **153** 

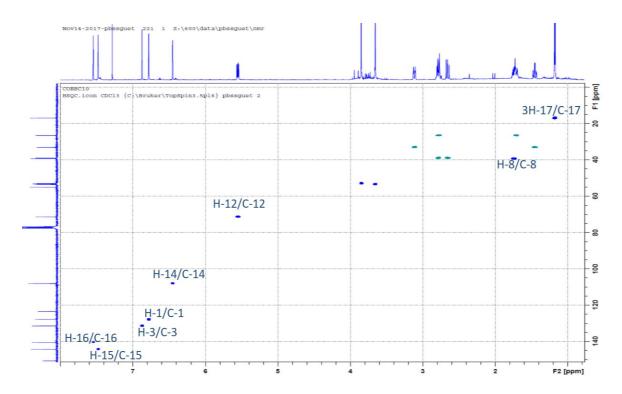


Figure 3.18 HSQC spectrum of 153

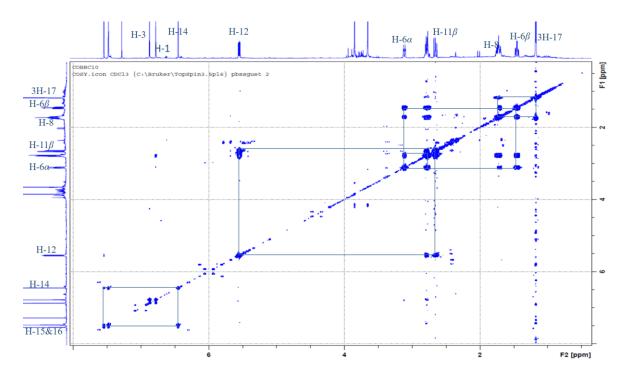


Figure 3.19 <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **153** 

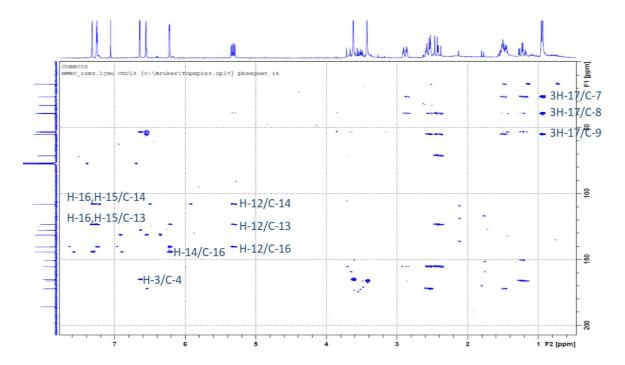


Figure 3.20 HMBC spectrum of 153

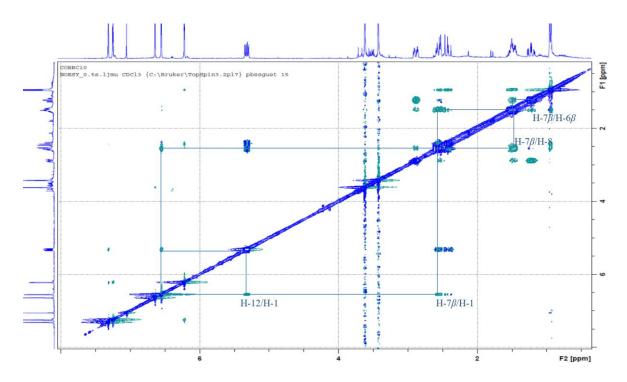


Figure 3.21 <sup>1</sup>H-<sup>1</sup>H NOESY spectrum of **153** 

# 3.3.1.5 Structure elucidation of 12-epi-megalocarpoloide D (154)

Compound **154** was isolated as a white amorphous powder,  $[\alpha]_{D}^{25}$  +81.7 (c 0.0018, MeOH). Its molecular formula C<sub>22</sub>H<sub>22</sub>O<sub>8</sub> was determined from the *pseudo*molecular ion peak at *m*/*z* 432.1650 calculated for C<sub>22</sub>H<sub>22</sub>O<sub>8</sub>NH<sub>4</sub> [M+NH<sub>4</sub>]<sup>+</sup>, 432.1653 from its HR ESI-MS spectrum obtained in positive ion mode. The IR spectrum of **154** quite identical to that of **153**, also displayed absorption stretching bands at 1715, 1767 and 1663 cm<sup>-1</sup> attributable to the carbonyl of ester, lactone and ketone groups, respectively. The <sup>1</sup>H and <sup>13</sup>C NMR spectra (Figures 3.22-3.23) of compound **154** as well as its 2D NMR spectra including COSY, HSQC and HMBC were all similar to those of **153** with only slight differences in terms of chemical shifts (Tables 3.7-3.8). Analysis of its <sup>1</sup>H, <sup>13</sup>C, COSY, HSQC and HMBC spectral data concluded its core structure to be identical to that of **153**. However, in the NOESY spectrum (Figure 3.24), a strong correlation observed between H-12 and H-17 suggested an  $\alpha$ -orientation of H-12 and thus, established a 12*R* 

configuration for **154** opposite to the 12*S* configuration found for **153** (Ndunda *et al.*, 2016). Therefore, the structure of compound **154** supported by its NOESY and HMBC key correlations (Figure 3.23) was established to be the C-12*R* epimer of **153** and named 12*-epi*-megalocarpoidolide D, a new clerodane from natural source.

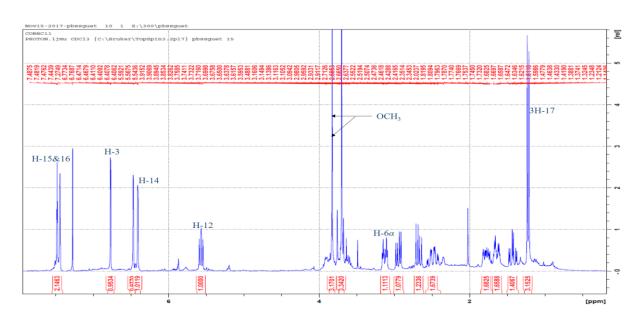


Figure 3.22  $^{1}$ H NMR (300 MHz, CDCl<sub>3</sub>) of **154** 

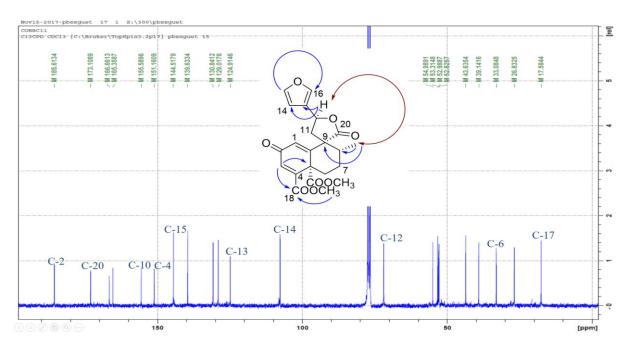


Figure 3.23 <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) and key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **154** 

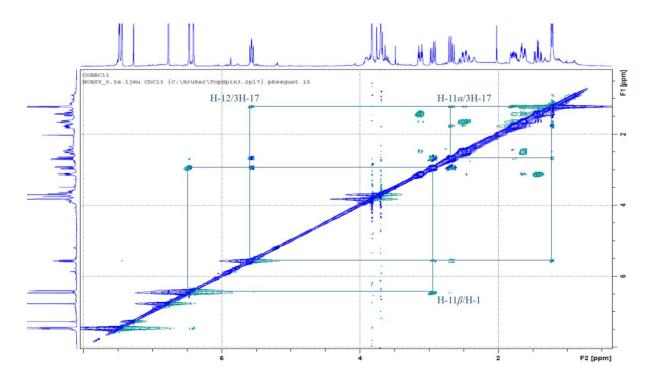


Figure 3.24 <sup>1</sup>H-<sup>1</sup>H NOESY spectrum of **154** 

# 3.3.1.6 Structure elucidation of crotonolin A (156) and crotonolin B (157) as a mixture

Compounds **156** and its 15-epimer **157** were isolated as a white amorphous powder mixture ( $[\alpha]_D^{25}$  0.0 (c 0.009, MeOH) with molecular formula of C<sub>22</sub>H<sub>22</sub>O<sub>10</sub> as determined from their HR ESI-MS spectrum obtained in the negative ion mode, where the peak at *m*/*z* 445.1142 [M-H]<sup>-</sup> could be calculated 445.1140 for C<sub>22</sub>H<sub>21</sub>O<sub>10</sub>. Their FT-IR spectrum showed absorption bands at 3465, 1775, 1712 and 1661 cm<sup>-1</sup> assignable to the stretch signals of hydroxyl and the carbonyls of ester, lactone and ketone groups, respectively. The <sup>1</sup>H (Figure 3.25) and <sup>13</sup>C NMR (Figure 3.26) spectra of the mixture of **156** and **157** similar to those of compound **153**, also suggested a clerodane skeleton (Li *et al.*, 2016). The only difference between the <sup>1</sup>H NMR of the mixture of **156** and **157** and that of compound **153** was the absence of the signals corresponding to the *β*-substituted furan ring protons in the <sup>1</sup>H NMR spectrum of **156** and **157**. Those furan ring protons were replaced by two methines doublets at  $\delta_H$  6.19 and 7.40 showing cross peak correlations

in the HSQC spectrum with carbon signals at  $\delta_{\rm C}$  99.3 and 149.8, respectively. These two methines, in the HMBC spectrum showed a strong correlation with a deshielded carbon signal at  $\delta_{\rm C}$  170.9. The above NMR data were similar to those of the unsaturated yhydroxy- $\alpha,\beta$ -unsaturated- $\gamma$ -lactone moiety present in salvidinin B, a clerodane diterpene isolated from Salvia divinorum (Shirota et al., 2006). The hemiacetal carbon C-15 was then attributed to the signal at  $\delta_{\rm C}$  99.3, the olefinic carbon C-14 at  $\delta_{\rm C}$  149.8, the carbonyl C-16 at  $\delta_{\rm C}$  170.9 and C-13 at  $\delta_{\rm C}$  135.0. These attributions were supported by the <sup>3</sup>J correlation between the proton signal at  $\delta_{\rm H}$  7.40 (H-14) and the carbon C-12, and correlations between protons H-12 and H-11 $\beta$  with C-13 observed in the HMBC experiment. However, the signals attributed to protons H-14 and H-15 were not broad singlets like in the case of similar molecular (Blas et al., 2004; Shirota et al., 2006; Maldonado et al., 2016). In addition, the two protons were not showing any correlation in the COSY experiment and the coupling constants of two doublets were different (6.5 and 9.4 Hz). Therefore, it was clear that two broad singlets instead of doublets were present; this led to the conclusion that the obtained powder was a mixture of  $15\alpha$ -OH and  $15\beta$ -OH isomers. Furthermore, analysis of the NOESY spectrum showed a strong correlation between H-12 and H-1 suggesting a C-12S configuration. In addition, the optical inactivity of the mixture indicated that it was a racemic mixture. Based on their <sup>1</sup>H and <sup>13</sup>C NMR (Table 3.7-3.8) and the <sup>1</sup>H-<sup>13</sup>C long-range correlations observed in the HMBC spectrum (Figure 3.26), compounds 156 and 157 were identified as 2,16-dioxo-15β-hydroxy-18,19-dimethoxycarbonyl-15,16-epoxy-ent-cleroda-1(10),3,13-trien-

20,12-olide and 2,16-dioxo-15 $\alpha$ -hydroxy-18,19-dimethoxycarbonyl-15,16-epoxy-*ent*-cleroda-1(10),3,13(16),14-tetraen-20,12-olide, two new clerodane diterpenes and were given the trivial names crotonolins A (**158**) and B (**159**), respectively.

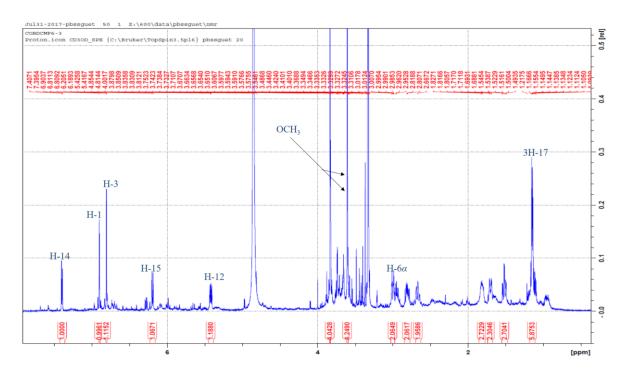


Figure 3.25  $^1\text{H}$  NMR (600 MHz, CD<sub>3</sub>OD) of **156** and **157** 

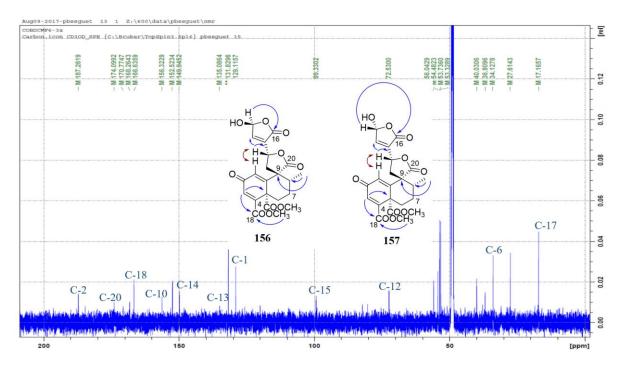


Figure 3.26 <sup>13</sup>C NMR (150 MHz, CD<sub>3</sub>OD) and key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **156** and **157** 

Compounds 158 and its epimer 159 were isolated as a white amorphous powder mixture  $([\alpha]_D^{25} - 16.5 (c 0.0002, MeOH))$  with molecular formula of  $C_{22}H_{22}O_{10}$  as determined from their HR-ESI-MS spectra obtained in the positive ion mode, where the sodiated ion peak was observed at m/z 469.1120 [M+Na]<sup>+</sup>, calculated 469.1111 for C<sub>22</sub>H<sub>22</sub>O<sub>10</sub>Na. Their FT-IR spectrum showed absorption bands at 3465, 1775, 1712 and 1670 cm<sup>-1</sup> that could be assigned to the stretch signals of hydroxyl and the carbonyls of ester, lactone and ketone groups, respectively. Their <sup>1</sup>H NMR and <sup>13</sup>C NMR spectra (Figures 3.27-3.28) similar with those of compounds 156 and 157, also suggested a clerodane skeleton containing a  $\gamma$ -hydroxy- $\alpha,\beta$ -unsaturated- $\gamma$ -lactone moiety (Shirota *et al.*, 2006). However, the observed chemical shifts corresponding to the proton and carbon resonances of the lactones units in 158 and 159 were different from those observed for 156 and 157. The lactone units 158 and 159 were identified as 16-hydroxy-13-en-15,16-olides from their NMR data (Blas et al., 2004; Shirota et al., 2006; Maldonado et al., 2016). In this case, the combination of the different signals and correlations observed in the 1D and 2D NMR spectra confirmed the suggested structure elements. The resonances at  $\delta_{\rm C}$  163.6 could be attributed to the  $\beta$ -carbon (C-13),  $\delta_{\rm H}$  6.22 and  $\delta_{\rm C}$  119.2 to the  $\alpha$ -methine (C-14),  $\delta_{\rm C}$  169.1 to the lactone carbonyl (C-15), and  $\delta_{\rm H}$  6.28 and  $\delta_{\rm C}$  96.7 to the hemiacetalic methine (C-16). The configuration of C-12 carbon in the compounds 158 and 159 were determined by NOESY experiment (Figure 3.28). A strong correlation observed in this spectrum between the H-12 protons and the H-1 protons suggested a C-12S orientation for both molecules. The <sup>1</sup>H and <sup>13</sup>C NMR data (Tables 3.7-3.8) of **158** and **159**, in addition to the key HMBC correlations (Figure 3.28) observed led to their characterisation as 2,15dioxo-16\beta-hydroxy-18,19-dimethoxycarbonyl-15,16-epoxy-ent-cleroda-

1(10),3,13(16),14-tetraen-20,12-olide and 2,15-dioxo-16α-hydroxy-18,19dimethoxycarbonyl-15,16-epoxy-*ent*-cleroda-1(10),3,13(16),14-tetraen-20,12-olide respectively, two clerodane diterpenes not previously described and were given the trivial names crotonolins C (**158**) and D (**159**), respectively.

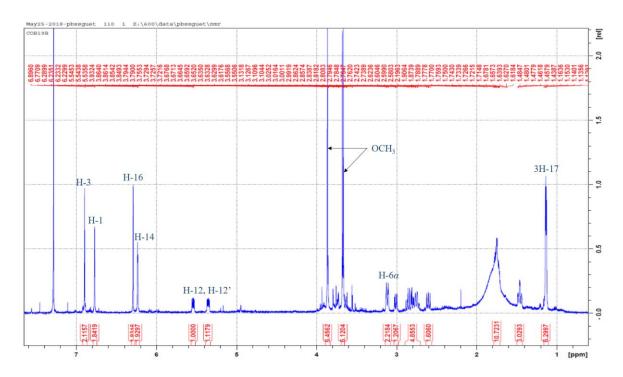


Figure 3.27  $^{1}$ H NMR (600 MHz, CDCl<sub>3</sub>) of **158** and **159** 

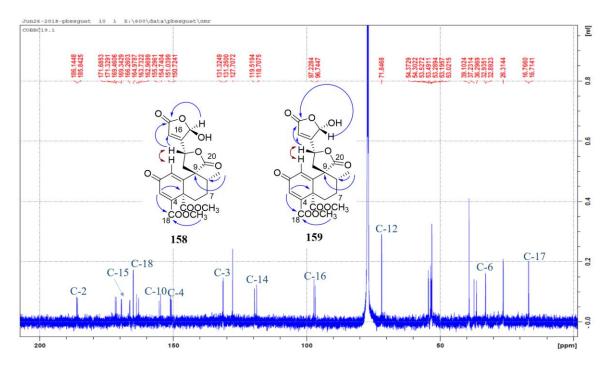


Figure 3.28 <sup>13</sup>C NMR (150 MHz, CDCl<sub>3</sub>) and key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **158** and **159** 

Compound 160 was isolated as a white amorphous powder,  $\left[\alpha\right]_{D}^{25}$  +58.4 (c 0.00025, MeOH. Its molecular formula C<sub>24</sub>H<sub>26</sub>O<sub>10</sub> was determined from its HR ESI-MS spectra obtained in the positive ion mode where the peak at m/z 497.1443 [M+Na]<sup>+</sup> was calculated for C<sub>24</sub>H<sub>26</sub>O<sub>10</sub>Na, 497.1418 Compound **160** could be identified as a  $\beta$ -substituted furan clerodane from its <sup>1</sup>H, <sup>13</sup>C NMR and HSQC spectra (Figures 3.29-3.31) which showed some similar features with those of previously identified clerodanes (Tchissambou et al., 1990). The <sup>1</sup>H NMR spectrum of **160** showed a couple of olefinic protons at  $\delta_{\rm H}$  5.88 (H-3) and 5.89 (H-2), which showed scalar coupling in the COSY spectrum, an oxymethine proton at  $\delta_{\rm H}$  5.21 (H-1) as well as a vinyl methyl proton resonance at  $\delta_{\rm H}$  1.84 (3H, t, J =1.2 Hz, H-17), which showed correlations in the HMBC spectrum (Figure 3.31) with the olefinic carbons at  $\delta_{\rm C}$  126.0 (C-7) and 129.3 (C-8) and the quaternary carbon at  $\delta_{\rm C}$  51.9 (C-9). The <sup>13</sup>C NMR spectrum revealed characteristic signals of r-lactone ( $\delta_C$  175.7, C-20; 71.8, C-12) and a  $\beta$ -substituted furan moiety ( $\delta_C$  124.8, C-13; 108.1, C-14; 144.2, C-15; 139.6, C-16). The signals at  $\delta_C$  172.0, 170.1 and 170.7 were attributed to the carbonyls of two methyl esters (C-18 and C-19) and an acetoxy function respectively. In the HMBC spectrum,  ${}^{3}J$  long-range  ${}^{1}H{}^{-13}C$  correlations could be observed between the oxymethine proton at  $\delta_{\rm H}$  5.21 (H-1) and the carbonyl of the acetoxy at  $\delta_{\rm C}$  170.7 suggesting an acetylation on that position C-1. Correlations were also observed between the carbon  $\delta_C$ 77.9 (C-4) and the protons at  $\delta_{\rm H}$  5.21 (H-1), 3.14 (H-10) and 2.36 (H-6). In the NOESY spectrum, a strong correlation observed between H-12 and 3H-17 suggested that H-12 is on the same side as 3H-17. The correlation observed between H11 $\beta$ /H10 established the  $\beta$ -orientation of H-10. The NOESY experiment was not helpful to determine the position of the 1-acetoxy and 4-OH groups. The <sup>1</sup>H and <sup>13</sup>C NMR data of **160** (Tables 3.7-3.8), supported by its key HMBC correlations (Figure 3.30) led to its characterisation as 1\*acetoxy-4\*-hydroxy-18,19-dimethoxycarbonyl-15,16-epoxy-ent-cleroda-2,7,13(16),14tetraen-20,12-olide, a new *ent*-clerodane diterpene not previously described and given the trivial name crotonolin F.

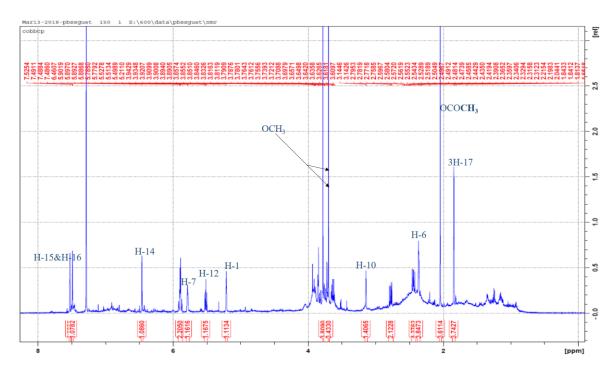


Figure 3.29  $^{1}$ H NMR (600 MHz, CDCl<sub>3</sub>) of **160** 

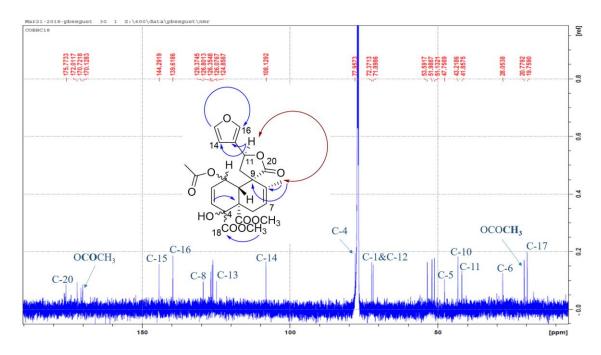


Figure 3.30 <sup>13</sup>C NMR (150 MHz, CDCl<sub>3</sub>) and key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **160** 

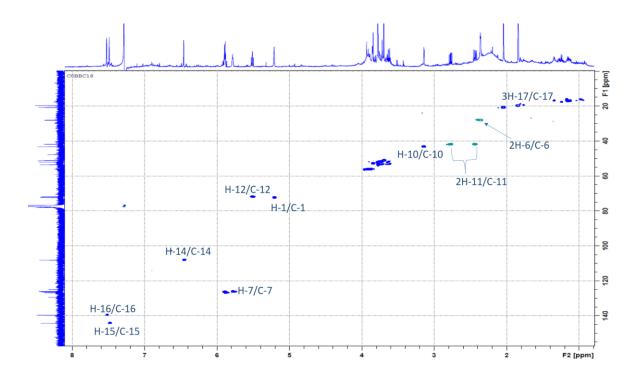


Figure 3.31 HSQC spectrum of 160

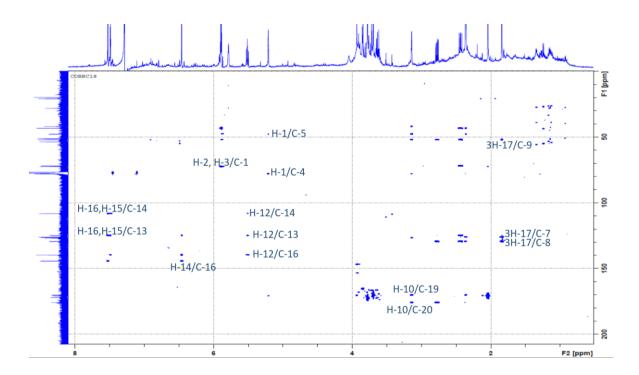


Figure 3.32 HMBC spectrum of 160

Compound 161 was obtained as a yellow powder and assigned the molecular formula of  $C_{20}H_{26}O_3$  on the basis of the *pseudo* molecular ion peak at m/z 315.1963 [M + H]<sup>+</sup> calculated for  $[C_{20}H_{26}O_3 + H]^+$ , 315.1966 from the HR ESI-MS spectrum. Its <sup>1</sup>H NMR spectrum (Figure 3.33; Table 3.7) indicated the presence of four methyls including two singlets at  $\delta_H$  1.38 and 1.29 and two doublets at  $\delta_H$  1.26 (J = 1.7 Hz) and 1.27 (J = 1.7Hz). Four methylenes and five methines including three aromatic methines at  $\delta_{\rm H}$  7.32 (d, J = 8.2 Hz), 7.42 (dd, J = 2.0, 8.2 Hz) and 7.89 (d, J = 2.0 Hz) as well as a septuplet methine at  $\delta_{\rm H}$  2.95, which coupled with the two methyl doublets in the COSY spectrum, were also observed. These observations were supported by the different correlations observed on the HSQC-DEPT. The <sup>13</sup>C NMR spectrum (Figure 3.34; Table 3.8) also indicated the presence of seven quaternary carbons including two carbonyls at  $\delta_{\rm C}$  198.7 and 181.4, which could be attributed to ketone and carboxylic acid functions respectively. In the HMBC spectrum, long-range correlations could be observed between the methine proton at  $\delta_{\rm H}$  2.95 (H-14) and the carbons at  $\delta_{\rm C}$  132.6 (C-12), 125.1 (C-16), 23.8 (C-15) and 23.7 (C-17). The proton at  $\delta_{\rm H}$  2.51 (H-6 $\alpha$ ) showed correlations with the carbons at  $\delta_{\rm C}$ 198.7 (C-7) and 37.2 (C-10), while the H-6 $\beta$  proton at  $\delta_{\rm H}$  2.76 showed correlations with the carbons on position C-4 ( $\delta_{C}$  46.3), C-5 ( $\delta_{C}$  43.6), C-7( $\delta_{C}$  198.7), C-10 ( $\delta_{C}$  37.2) and C-19 ( $\delta_{\rm C}$  16.2). The <sup>1</sup>H and <sup>13</sup>C NMR data (Tables 3.7-3.8), supported by the key HMBC correlations (Figure 3.33), were in good agreement with those published for 7oxodehydroabietic acid (Tanaka et al., 1997). Therefore, compound 161 was identified as 7-oxo-dehydroabietic acid.

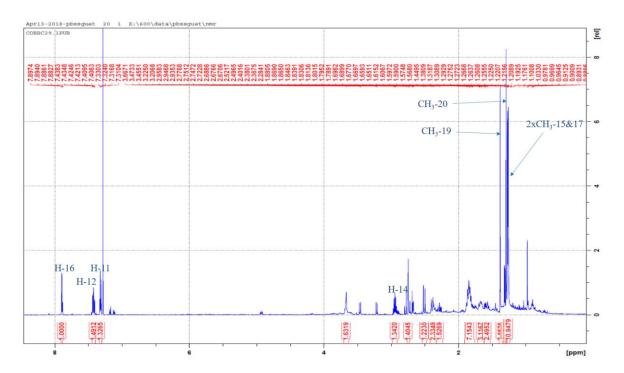


Figure 3.33 <sup>1</sup>H NMR spectrum (600 MHz, CDCl<sub>3</sub>) of 161

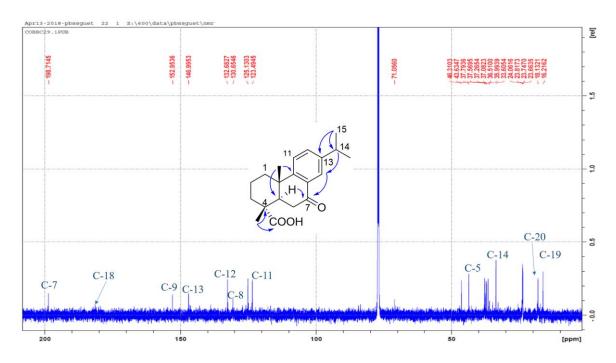


Figure 3.34 <sup>13</sup>C NMR (600 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **161** 

Position						$\delta_{\rm H} (J \text{ in Hz})$					
	151	152	153	154	155	<b>156/157</b> <sup>a</sup>	158	159	160	161	162
1	1.92 m	6.00 dd	6.47 d	6.47 d (1.0)	6.04 brt	6.91 d ( <i>1.2</i> )	6.76 s	6.76 s	5.21 m	1.84 m	1.49 m
		(3.5, 9.3)	(1.3)		(1.9)						
	2.65 m	-	-	-	-	-	-	-	-	-	2.31 m
2	2.43 m	6.20 dd	-	-	4.87 t	-	-	-	5.89 ov	1.67 m	1.83 m
		(3.1, 4.9)			(2.4)						
	2.58 m	-	-	-	-	-	-	-	-	2.39 m	-
3	6.84 dd	7.00 d	6.78 d	6.78 d (1.0)	6.70 dd	6.80 d ( <i>1.2</i> )	6.88 s	6.88 s	5.88 ov	1.84 m	1.62 m
	(4.3, 7.5)	(5.2)	(1.3)		(1.8, 2.3)						
						-	-	-	-	-	1.83 m
5						-	-	-	-	2.74	2.48 m
6	1.12 ddd	1.32 m	1.45 ddd	1.44 ddd	1.31 m	1.51 ddd (4.0,	1.45 ddd	1.45 ddd	2.36 m	2.51 d	1.72 m
	(3.6, 13.4,		(3.7, 13.3)	(13.5, 9.3,		13.5 17.5)	(3.2, 13.1,	(3.2, 13.1,		(15.2)	
	16.9)			17.7)			16.9)	16.9)			
	2.95 dt (3.3,	2.95, dt	3.11 dt	3.13 dt	3.04 dt	3.00 m	3.10, dt ( <i>3</i> . <i>1</i> ,	3.10, dt ( <i>3</i> . <i>1</i> ,	5.78 brt	2.76 d	2.12 m
	6.4)	(3.7,	(3.0, 6.1)	(13.3, 3.3)	(3.3, 13.3)		13.0)	13.0)	(3.4)	(15.2)	
		13.0)									
7	1.58 m	1.56 m	1.71 m	1.65 m	1.57 dd	1.66 m	1.74 m	1.74 m	-	-	4.79 d
					(3.6,						(3.4)
					14.2)						
	2.43 m	2.28 m	2.78 m	2.49 ddd	2.40 m	2.66 m	2.77 m	2.77 m	-	-	-
				(4.2, 13.9,							
				17.1)							
8	1.58 m	1.70 m	1.75 m	1.78 m	1.69 m	1.82 m	1.78 m	1.78 m	-	-	-

# Table 3.7 <sup>1</sup>H NMR data of compounds **151-162**

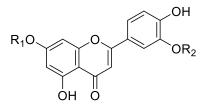
Table 3.7	continued										
Position						$\delta_{\rm H} (J \text{ in Hz})$					
	151	152	153	154	155	<b>156/157</b> <sup>a</sup>	158	159	160	161	162
10	1.76 dd	2.89 t	-	-	-	-	-	-	3.14 m		-
	(2.4, 13.2)	(3.2)									
11	2.43 m	2.48 m	2.78 m	2.69 dd (8.2,	2.62 dd	2.80 m	2.59 dd	2.81 m	2.43 dd	7.32 d	7.18 d
				14.3)	(8.8, 14.3)		(11.2, 14.6)		(9.4,	(8.2)	(8.3)
									14.1)		
	-	-	2.65 dd	2.94 dd (6.3,	2.94 dd	2.93 m	2.99 dd (5.3,		2.77 dd	-	-
			(11.1,	14.2)	(6.2, 14.3)		14.6)		(7.9, 14.1)		
			14.5)								
12	5.40 t (8.5)	5.48 t	5.55 dd	5.57 t (7.0)	5.53 dd	5.72 dd (5.5,	5.34 dd (5.3,	5.53 dd (6.7,	5.50 m	7.42 dd	7.12 dd
		(8.7)	(5.2, 11.1)		(6.4, 8.7)	11.1)	11.2)	10.7)		(2.0, 8.2)	(1.9, 8.2)
14	6.41 brd	6.45 d	6.45 m	6.41 m	6.44 d	7.40/7.39, brs	6.22 s	6.22 s	6.46 brs	7.89 d	7.20 d
	(0.9)	(0.9)			(0.9)					(2.0)	(1.9)
15	7.44 ov	7.49 brt	7.47 dd	7.49 brd	7.49 dd	6.18 /6.20, brs	-	-	7.48 brt	2.95 m	2.86 m
		(1.6)	(1.5, 3.1)	(1.6)	(0.9, 1.6)				(1.6)		
16	7.45 brs	7.51 brs	7.54 brs	7.45 m	7.48 brs		6.28 s	6.28 s	7.52 brs	1.26 d	1.23 d
										(6.4)	(6.4)
17	1.02 d (6.4)	1.15 d	1.17 d	1.23 d (6.7)	1.17 d	1.15 d (5.8)	1.12 d (6.5)	1.11 d (6.3)	-	1.27 d	1.23 <i>d</i>
		(6.9)	(6.4)		(6.7)					(6.4)	(6.4)
19						-	-	-	-	1.38 s	1.28 s
20						-	-	-	-	1.29 s	1.16 s
18-OCH <sub>3</sub>	3.71 s	3.74 s	3.84 s	3.84 s	3.76 s	3.71 s	3.85 s	3.85 s	3.78 s	-	-
19-OCH <sub>3</sub>	3.76 s	3.62 s	3.65 s	3.71 s	3.71 s	3.60 s	3.66 s	3.65 s	3.69 s	-	-
COCH <sub>3</sub>						-	-	-	2.04 s	-	-

Position	$\delta_{\rm C}$										
	151	152	153	154	155	156,157	158	159	160	161	162
1	19.2	132.7	127.8	129.1	129.4	129.0	127.7	127.7	72.3	18.4	37.8
2	26.4	125.9	185.8	185.7	64.4	187.2	186.1	185.8	126.8	37.0	18.6
3	140.1	135.5	131.4	130.9	136.2	131.7	131.3	131.2	126.3	36.5	36.5
4	136.5	136.3	150.7	151.3	156.1	152.4	150.7	151.0	77.9	46.3	47.1
5	46.3	47.0	53.5	55.1	48.3	54.3	53.4	53.2	47.7	43.6	39.8
6	32.3	31.9	33.1	33.2	31.4	33.9	32.9	32.9	28.0	37.7	31.1
7	27.9	28.1	26.5	27.2	27.2	27.5	26.4	26.4	126.0	198.7	68.3
8	40.2	43.1	39.7	43.7	41.7	40.0	39.1	39.1	129.3	130.6	135.9
9	51.4	50.7	55.0	53.6	53.3	55.9	54.3	54.3	51.9	152.9	146.7
10	51.9	50.0	155.4	155.7	151.9	156.2	155.2	154.7	43.2	37.2	37.5
11	42.4	41.8	38.9	39.2	40.0	36.9	37.1	36.3	41.8	123.4	124.3
12	71.9	72.1	71.2	72.0	71.8	72.4	71.5	71.5	71.8	132.6	126.7
13	125.5	125.4	123.5	125.0	125.8	135.0	163.7	162.9	124.8	146.9	146.7
14	108.2	108.3	108.1	107.8	107.9	149.8	119.5	118.7	108.1	125.1	128.3
15	144.2	144.5	144.4	144.6	144.2	99.3	169.3	169.4	144.2	33.6	33.6
16	139.5	139.4	140.5	139.7	139.3	170.9	96.7	97.2	139.6	23.8	23.9
17	17.1	17.5	16.9	17.7	17.8	17.1	17.1	17.1	19.7	23.7	23.9
18	166.9	166.9	165.3	165.5	166.2	166.7	164.9	164.9	172.0	181.4	182.2
19	173.0	173.4	166.3	166.7	170.7	168.4	166.3	166.2	170.0	16.2	16.4
20	176.3	176.6	172.1	173.1	174.8	174.0	171.1	171.6	175.7	23.6	24.1
18-OCH <sub>3</sub>	51.7	51.8	53.0	52.9	52.0	53.3	53.0	53.0	53.5		
19-OCH <sub>3</sub>	51.8	51.7	53.2	53.1	52.6	53.7	53.5	53.2	51.1		
COCH <sub>3</sub>									170.7		
COCH <sub>3</sub>									20.7		

Table 3.8 <sup>13</sup>C NMR data of compounds **151-162** 

#### 3.3.2 Phytochemistry of Ruspolia hypocrateriformis

Combination of CC, SPE, preparative TLC and RP-HPLC analyses of the methanol extract of the leaves of R. hypocrateriformis afforded ten compounds including six glycosylated flavonoids, a quindoline, two pyrrolidine alkaloids and a disaccharide. The compounds were identified luteolin-7-O- $\beta$ -D-apiofuranosyl- $(1 \rightarrow 2)$ - $\beta$ -Das xylopyranoside (163), chrysoeriol-7-O- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside chrysoeriol-7-O-[4"'-O-acetyl- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)]- $\beta$ -D-xylopyranoside (164).(165), luteolin-7-O- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (166), chrysoeriol-7-*O*- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (167),luteolin 7-O-[*b*-Dglucopyranosyl- $(1\rightarrow 2)$ - $\alpha$ -L-rhamnosyl- $(1\rightarrow 6)$ ]- $\beta$ -D-glucopyranoside 10H-(168),quindoline (169), secundallerones B and C (170-171) and sucrose (60) (Figure 3.35).



**163**  $R_1 = \beta$ -D-apiofuranosyl-(1<sup>'''</sup> $\rightarrow$ 2<sup>''</sup>)- $\beta$ -D-xylopyranosyl  $R_2 = H$  **164**  $R_1 = \beta$ -D-apiofuranosyl-(1<sup>'''</sup> $\rightarrow$ 2<sup>''</sup>)- $\beta$ -D-xylopyranosyl  $R_2 = CH_3$  **165**  $R_1 = 4^{'''}$ -*O*-acetyl- $\beta$ -D-apiofuranosyl-(1<sup>'''</sup> $\rightarrow$ 2<sup>''</sup>)- $\beta$ -D-xylopyranosyl  $R_2 = CH_3$  **166**  $R_1 = \alpha$ -L-rhamnopyranosyl-(1<sup>'''</sup> $\rightarrow$ 2<sup>''</sup>)- $\beta$ -D-xylopyranosyl  $R_2 = H$  **167**  $R_1 = \alpha$ -D-rhamnopyranosyl-(1<sup>'''</sup> $\rightarrow$ 2<sup>''</sup>)- $\beta$ -D-xylopyranosyl  $R_2 = CH_3$ **168**  $R_1 = \beta$ -D-glucopyranosyl-(1<sup>''''</sup> $\rightarrow$ 2<sup>'''</sup>)- $\alpha$ -L-rhamnosyl-(1<sup>''''</sup> $\rightarrow$ 6<sup>''</sup>)- $\beta$ -D-glucopyranosyl  $R_2 = H$ 

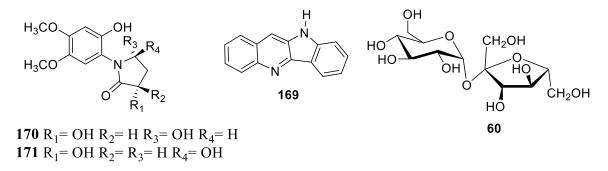


Figure 3.35 Structures of isolated compounds from the leaves of R. hypocrateriformis

3.3.2.1 Structure elucidation of luteolin-7-*O*- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (**163**)

Compound 163 was isolated as a yellow amorphous powder. Its molecular formula  $C_{25}H_{26}O_{14}$  was determined by the peak at m/z 551.1392 calculated for  $[C_{25}H_{26}O_{14}+H]^+$ , 551.1395 and consistent with thirteen RDB, from its HR ESI-MS spectrum (Figure 3.36) in positive mode ion. The <sup>1</sup>H NMR spectrum (Figure 3.38; Table 3.9) revealed 17 peaks belonging to a glycosylated molecule bearing two sugar moieties. The aglycone part of the compound was identified as luteolin by the presence of an ABX system resonating at  $\delta_{\rm H}$  7.26 (d, J = 8.2 Hz), 7.49 (dd, J = 2.1, 8.2 Hz) and 7.87 (d, J = 2.1 Hz), characteristic of a 3,4-dihydroxylated B ring; an AX system appearing as two doublet signals at  $\delta_{\rm H}$  6.92 and 6.95 (J = 2.0 Hz), typical of a *meta*-dihydroxylated A ring, and one singlet at  $\delta_{\rm H}$  6.89 attributable to the H-3 proton of the luteolin C ring. The sugar moieties were identified by the presence of two anomeric proton signals at  $\delta_{\rm H}$  5.58 (d, J = 7.4 Hz) and 6.52 (d, J =0.9 Hz). 2D NMR experiments, mainly COSY and HSQC (Figures 3.39-3.40), helped to assign the resonance signals belonging to the sugar moieties. COSY and TOCSY were useful to construct each sugar moiety and to identify proton signals belonging to the same sugar. This led to the identification of two pentose residues, one as  $\beta$ -D-apiofuranoside and the second as and  $\beta$ -D-xylopyranose by comparison of their NMR spectroscopic data with those reported in the literature (Koffi et al., 2013; Tagousop et al., 2017). Analysis of the HMBC spectrum (Figure 3.41) revealed that the two sugars were linked together.  ${}^{3}J$  long-range correlations were observed between the proton H-1" ( $\delta_{\rm H}$  6.52) of the apiose and the carbon C-2" of the xylose ( $\delta_{\rm C}$  77.0 ppm), and between the proton H-2"  $(\delta_H 4.49)$  of the xylose and the carbon C-1''' ( $\delta_C 110.5$ ) of the apiose. Other long-range correlations observed between the proton H-1'' ( $\delta_{\rm H}$  5.58) of the xylose and the carbon C-7 ( $\delta_{\rm C}$  163.3) of the luteolin aglycone and between the proton H-8 ( $\delta_{\rm H}$  6.95) of the aglycone and the anomeric carbon ( $\delta_{\rm C}$  100.3, C-1") of the xylose confirmed the attachment of the 132

diglycoside unit on the carbon C-7 of the aglycone. All the <sup>1</sup>H and <sup>13</sup>C NMR data of **163** (Tables 3.9-3.10), as well as the key HMBC correlations (Figure 3.37) identified **163** as luteolin 7-O-[ $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)]- $\beta$ -D-xylopyranoside, a glycosylated flavone previously isolated from two Acanthaceae species i.e. *Graptophyllum grandulosum* and *Justicia secunda* (Koffi *et al.*, 2003; Tagousop *et al.*, 2017).

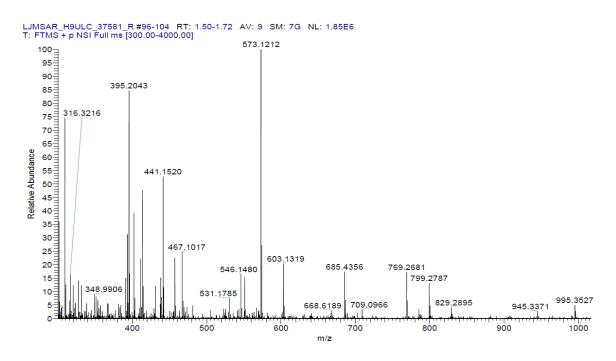


Figure 3.36 ESI-MS spectrum of 163

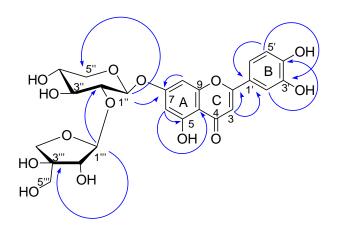


Figure 3.37 Key HMBC correlations of 163

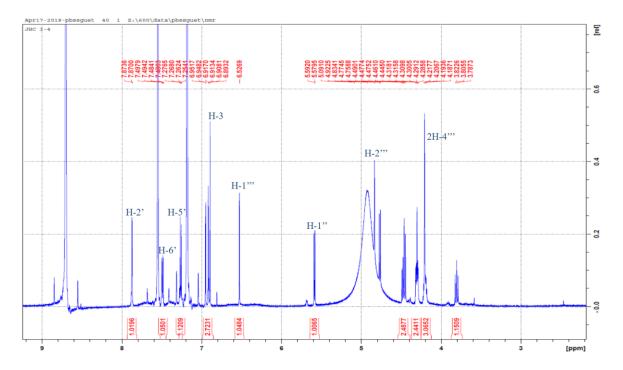


Figure 3.38  $^{1}$ H NMR (600 MHz, Pyr-d<sub>5</sub>) of **163** 

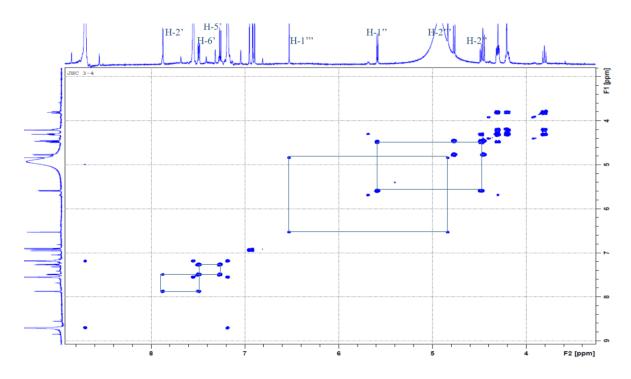


Figure 3.39 <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **163** 

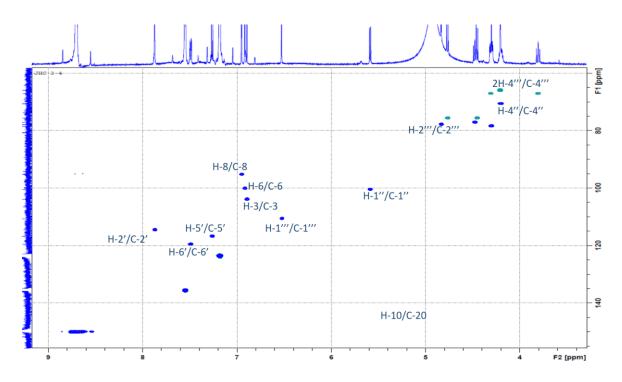


Figure 3.40 HSQC spectrum of 163

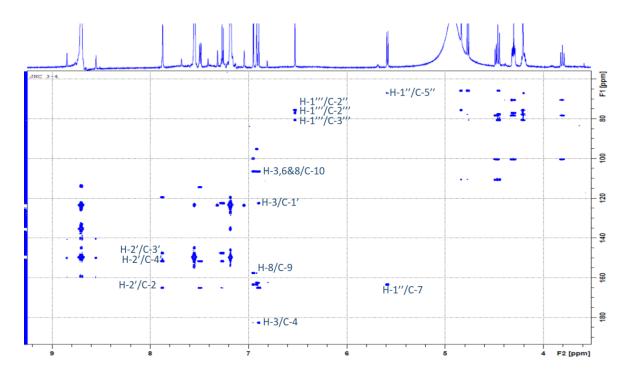


Figure 3.41 HMBC spectrum of 163

3.3.2.2 Structure elucidation of chrysoeriol-7-*O*- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (**164**)

Compound **164** was isolated as a yellow amorphous powder. Its molecular formula  $C_{26}H_{28}O_{14}$  was determined from the sodiated ion peak at m/z 587 [M+Na]<sup>+</sup> in its ESI-MS spectrum in positive mode ion. Its 1D and 2D NMR spectra similar to those of **163** suggested a luteolin diglycoside skeleton. The only difference was the presence of an additional signal on the <sup>1</sup>H NMR (Figure 3.42, Table 3.9) of **164** of a singlet attributable to a methoxy group at  $\delta_H$  3.78. This was confirmed by the cross peak correlation observed on its HSQC spectrum between this signal and the carbon resonating at  $\delta_C$  55.8. In the HMBC spectrum, a <sup>3</sup>J long-range correlation was observed between the signal of the methoxy proton and a carbon resonating at  $\delta_C$  148.5 corresponding to the carbon C-3' of the luteolin skeleton. Therefore, the aglycone in compound **164** was identified as 3'-methoxyluteolin or chrysoeriol. All the <sup>1</sup>H and <sup>13</sup>C NMR data of **164** (Tables 3.9-3.10), and the key long-range correlations observed in the HMBC spectrum (Figure 3.42) led to its identification as chrysoeriol-7-*O*- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside or granduloside A, a flavonoid previously isolated from *Graptophyllum grandulosum* (Tagousop *et al.*, 2017).

3.3.2.3 Structure elucidation of chrysoeriol-7-O-[4'''-O-acetyl- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)]- $\beta$ -D-xylopyranoside (**165**)

Compound **165** was also isolated as a yellow amorphous powder. Its molecular formula  $C_{28}H_{30}O_{15}$  was determined by the peak at m/z 629 [M+Na]<sup>+</sup> in its ESI-MS spectrum in positive mode ion. Its 1D and 2D NMR spectra similar to those of **164** also suggested a chrysoeriol diglycoside skeleton. The only difference was the presence of an additional

signal on the <sup>1</sup>H NMR (Figure 3.43, Table 3.9) of **165** of a singlet at  $\delta_{\rm H}$  1.93 (3H) attributable to the methyl protons of an acetoxy function. This was confirmed by the cross peak correlation observed on its HSQC spectrum between this methyl signal and the carbon resonating at  $\delta_{\rm C}$  19.2. These observations were supported in the HMBC spectrum (Figure 3.43), by the <sup>2</sup>*J* correlations observed between the methyl signal at  $\delta_{\rm H}$  1.93 and the deshielded carbon signal at  $\delta_{\rm C}$  171.1 attributable the carbonyl of the acetoxy function. Another correlation observed between the 2H-4<sup>\*\*\*</sup> protons ( $\delta_{\rm H}$  4.08) of the apiose and the carbonyl carbon ( $\delta_{\rm C}$  171.1) established an acetylation of the alcohol function on position C-4<sup>\*\*\*</sup> of the apiose moiety. Thus, all the <sup>1</sup>H and <sup>13</sup>C NMR data of **167** (Tables 3.9-3.10), in addition to the key HMBC correlations observed (Figure 3.43) led to its identification as chrysoeriol-7-*O*-[4<sup>\*\*\*</sup>-*O*-acetyl- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)]- $\beta$ -D-xylopyranoside or granduloside B, previously isolated from *Graptophyllum grandulosum* (Tagousop *et al.*, 2017).

3.3.2.4 Structure elucidation of luteolin-7-*O*- $\alpha$ -D-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -Dxylopyranoside (**166**)

Compound **166** was isolated as a yellow amorphous powder. Its molecular formula  $C_{26}H_{28}O_{14}$  was confirmed from its HR ESI-MS spectrum (Figure 3.44) in positive mode ion, where the sodiated ion peak was observed at m/z 587.1378, calculated for  $[C_{26}H_{28}O_{14}+Na]^+$ , 587.1377, consistent with thirteen RDB. The <sup>1</sup>H NMR spectrum (Figure 3.45, Table 3.9) of **166** revealed a glycosylated flavone skeleton with the aglycone moiety clearly identifiable as luteolin. The only difference between the <sup>1</sup>H NMR spectrum of **166** with that of **163** was the resonance signals attributable to the sugar moieties. The <sup>1</sup>H NMR spectrum of **166** exhibited characteristic peaks belonging to two anomeric protons at  $\delta_H$  5.62 (d, J = 7.5 Hz) and 6.38 (d, J = 1.6 Hz) and a doublet methyl at  $\delta_H$  1.78 <sup>137</sup>

(J = 6.1 Hz) showing a cross peak correlation in the HSQC spectrum (Figure 3.46) with the carbon signal resonating at  $\delta_{\rm C}$  18.6 suggesting a rhamnose as one of the glycosylic unit. Complete assignment of the protons and carbons of the sugar units was achieved by analysis of COSY, HSQC, HMBC and TOCSY spectra of 166. The sugars were identified  $\beta$ -D-xylopyranose and  $\alpha$ -L-rhamnose with their anomeric protons resonating at  $\delta_{\rm H}$  5.62 (d, J = 7.5 Hz) and 6.38 (d, J = 1.6 Hz) respectively, by comparison of their chemical shifts with those reported in the literature (Osterdahl, 1979; Beier and Mundy, 1980; Agrawal, 1992). The HMBC spectrum (Figure 3.47) showed a correlation between the proton H-1" ( $\delta_{\rm H}$  5.62) of the xylose and the carbon C-7 ( $\delta_{\rm C}$  163.2) of the aglycone, confirming its direct attachment to the luteolin skeleton. Other correlations observed between the proton H-1" ( $\delta_{\rm H}$  6.38) of the rhamnose and the carbon C-2" of the xylose  $(\delta_C 77.4)$  and between the proton H-2"  $(\delta_H 4.47)$  of the xylose and the carbon C-1"  $(\delta_C$ 102.3) of the rhamnose confirmed attachment of the rhamnosyl unit on the carbon C-2" of the xylose. Thus, all the <sup>1</sup>H and <sup>13</sup>C NMR data of **166** (Tables 3.9-3.10), in additition to the keys 2D correlations observed (Figure 3.45) led to its characterisation as luteolin-7-O- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside, a new flavone diglycoside from natural source and was given the trivial name justicialoside A.

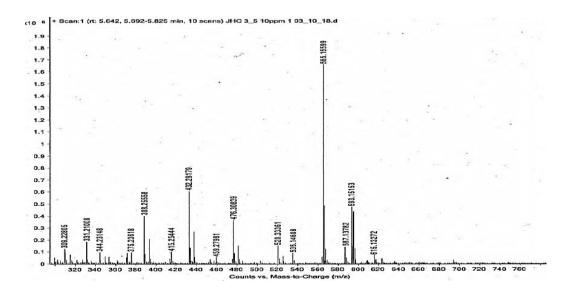


Figure 3.44 ESI MS spectrum of 166

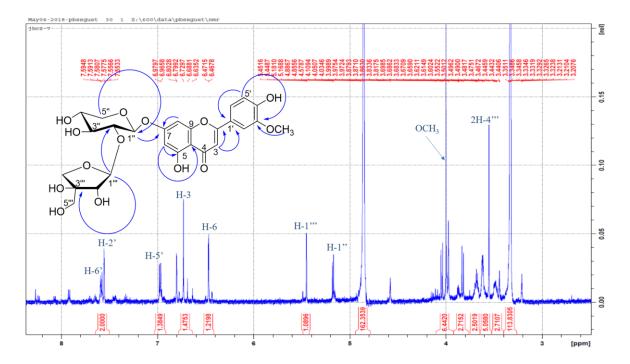


Figure 3.42 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of 164

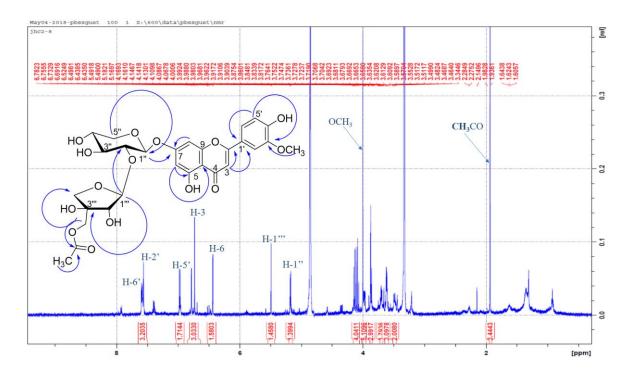


Figure 3.43  $^{1}$ H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **165** 

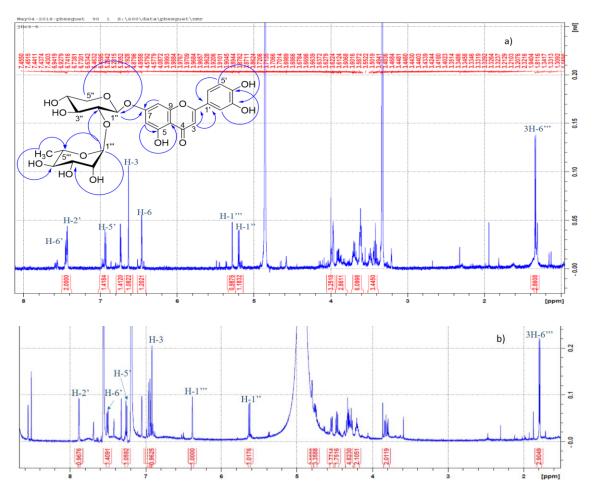


Figure 3.45 <sup>1</sup>H NMR (600 MHz, a-CD<sub>3</sub>OD, b-Pyr-d<sub>5</sub>) and key HMBC correlations of **166** 

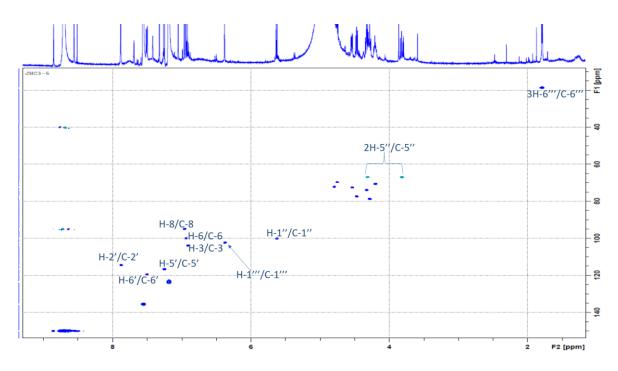


Figure 3.46 HSQC spectrum of 166

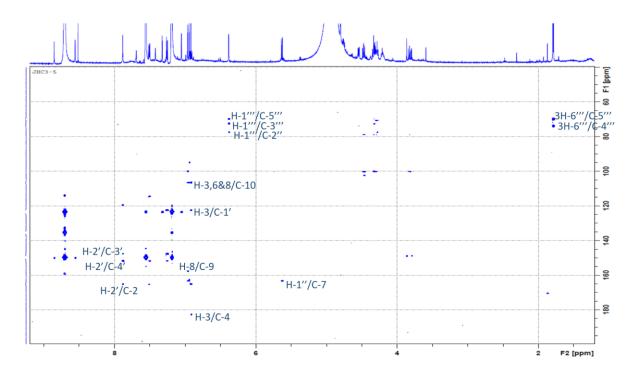


Figure 3.47 HMBC spectrum of 166

3.3.2.5 Structure elucidation of chrysoeriol-7-*O*- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (**167**)

Compound **167** was isolated as a yellow amorphous powder. Its molecular formula  $C_{27}H_{30}O_{14}$  was determined from its HR ESI-MS spectrum in positive mode ion by the peak at m/z 601.1533 calculated for  $[C_{27}H_{30}O_{14}+Na]^+$ , 601.1528 consistent with thirteen RDB. Its 1D and 2D NMR spectra, similar to those of **166**, also suggested a luteolin diglycoside skeleton. The only difference was the presence of an additional signal in the <sup>1</sup>H NMR (Figure 3.48, Table 3.9) of **167** of a singlet attributable to a methoxy group at  $\delta_H$  3.99 (3H). This was confirmed by the cross peak correlation observed on its HSQC spectrum between this signal and the carbon resonating at  $\delta_C$  55.4. In the HMBC spectrum (Figure 3.48), a cross peak correlation observed between the signal of the methoxy proton and a carbon resonating at  $\delta_C$  147.7 confirmed its position on the C-3' carbon of the luteolin skeleton. Thus, all the <sup>1</sup>H and <sup>13</sup>C NMR data (Tables 3.9-3.10), and the key

HMBC correlations of **167** (Figure 3.47) led to its identification as chrysoeriol-7-*O*- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside, a new flavone diglycoside from natural source and was given the trivial name justicialoside B.

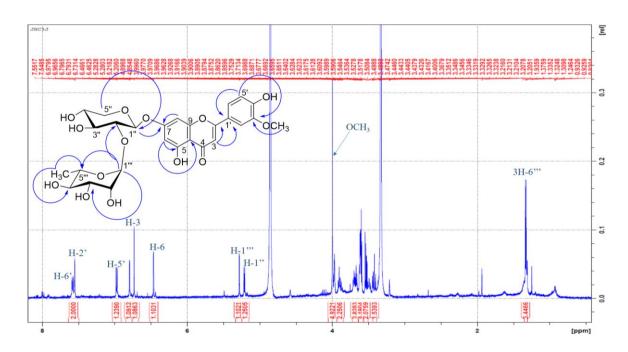


Figure 3.48 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of 167

3.3.2.6 Structure elucidation of luteolin 7-*O*-[ $\beta$ -D-glucopyranosyl-(1 $\rightarrow$ 2)- $\alpha$ -L-rhamnosyl-(1 $\rightarrow$ 6)]- $\beta$ -D-glucopyranoside (**168**)

Compound **168** was isolated as a yellow amorphous powder. Its molecular formula  $C_{33}H_{40}O_{20}$  was determined by the peak at m/z 779.2001 calculated for  $[C_{33}H_{40}O_{20}+Na]^+$ , 779.2005 and consistent with fourteen RDB, from its HR NSI-MS spectrum in positive mode ion. Its <sup>1</sup>H NMR spectrum (Figure 3.49, Table 3.9) revealed that **168** was a glycosylated luteolin. There were three sugar moieties recognizable by the three anomeric proton signals appearing at  $\delta_H$  4.69 (d, J = 7.9 Hz), 4.72 (d, J = 1.1 Hz) and 5.27 (d, J = 7.1 Hz). One of the sugar moieties could be identified as rhamnose by the presence of the characteristic methyl doublet at  $\delta_H$  1.19 (d, J = 6.1 Hz) (Osterdahl, 1979; Beier and

Mundy, 1980; Agrawal, 1992). TOCSY and COSY spectra of **168** were used to identify proton signals belonging to the same sugar moiety and for the construction of each sugar unit based on the direct <sup>1</sup>H-<sup>1</sup>H coupling of their respective protons. Their corresponding carbon resonances was assigned using the HSQC spectrum. The two other sugar moieties could be identified as  $\beta$ -D-glucopyranoses based on their chemical shifts (Beier *et al.*, 1980). In the HMBC spectrum, correlations were observed between the anomeric proton at  $\delta_{\rm H}$  5.27 (H-1<sup>''</sup>) and the carbon signal at  $\delta_{\rm C}$  163.0 attributable to the carbon C-7 of the luteolin aglycone. Other correlations could also be observed between the anomeric protons at  $\delta_{\rm H}$  4.69 (H-1<sup>'''</sup>) and 4.72 (H-1<sup>''''</sup>) with the carbons at  $\delta_{\rm C}$  82.0 (C-2<sup>''</sup>) and 66.1 (C-6<sup>''</sup>), respectively, of the glucopyranose unit attached to the aglycone. Thus, all the <sup>1</sup>H and <sup>13</sup>C NMR data of **168** (Tables 3.9-3.10), and the key HMBC correlations (Figure 3.49) confirmed its identification as luteolin 7-*O*-[ $\beta$ -glucopyranosyl-(1→2)- $\beta$ -rhamnosyl-(1→6)]- $\beta$ -glucopyranoside, a glycosylated flavone previously isolated from the leaves of *Justicia secunda* (Koffi *et al.*, 2013).

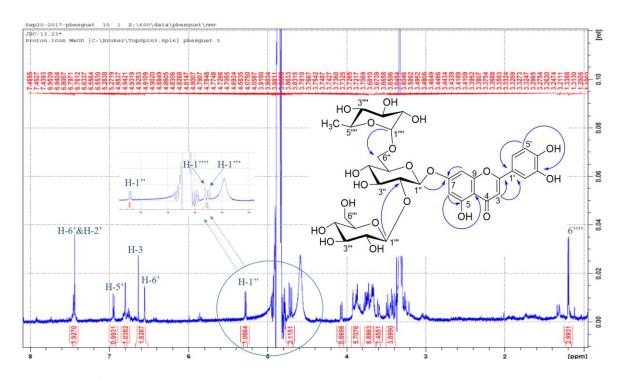


Figure 3.49 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **168** 

Compounds **170** and its diastereoisomer **171** were isolated as brown amorphous powder mixture with molecular formula of  $C_{12}H_{15}NO_6$  determined from their HR ESI-MS spectrum (Figure 3.50) obtained in negative ion mode by the peak at m/z 268.0828 calculated for  $[C_{12}H_{15}NO_6-H]^-$ , 268.0827. The <sup>1</sup>H NMR spectrum (Figure 3.51, Table 3.11) of the mixture showed two set of signals each attributable to four methines including two aromatic singlets suggesting a 1,2,4,6-tetrasubstituted benzene ring and two oxymethines, one methylene and two methoxyls. These attributions were supported by the different correlations observed on the HSQC-DEPT spectrum of the mixture as well as the chemical shift of the corresponding carbons. The <sup>1</sup>H and <sup>13</sup>C NMR spectra of **170** and **171** were quite similar to those published for secundellarones B and C, two diastereoisomeric pyrrolidone alkaloids isolated from the leaves of *J. secunda* (Theiler *et al.*, 2014). In addition, their <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.11) were in good agreement with the data published for secundallerone B and C (Theiler *et al.*, 2014). Thus, compounds **170** and **171** were identified as secundarellone B and C, respectively.

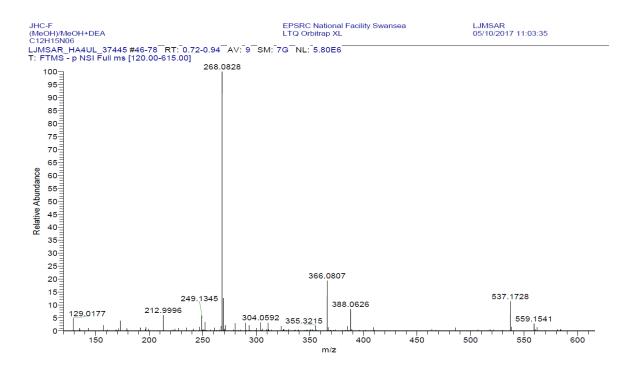


Figure 3.50 ESI-MS spectrum of 170 and 171

Position	$\delta_{\rm H} { m m} (J { m in} { m Hz})$									
	<b>163</b> <sup>a</sup>	<b>164</b> <sup>a</sup>	<b>165</b> <sup>b</sup>	<b>166</b> <sup>a</sup>	<b>167</b> <sup>b</sup>	<b>168</b> <sup>b</sup>				
3	6.89 s	6.93 s	6.76 s	6.91 s	6.73 s	6.63 s				
6	6.92 d (2.0)	6.94 d (2.1)	6.43 d (2.1)	6.94 d (2.1)	6.46 d (2.1)	6.55 d (2.0)				
8	6.95 d (2.0)	7.11 d (2.1)	6.78 d (2.1)	6.96 d (2.1)	6.79 d (2.1)	6.79 d (2.0)				
2'	7.87 d (2.1)	7.58 d (2.2)	7.56 d (2.0)	7.88 d (2.2)	7.55 d ( <i>1.9</i> )	7.43 d ( <i>1.6</i> )				
5'	7.26 d (8.2)	7.25 d (8.3)	6.97 d (8.2)	7.25 d (8.3)	6.97 d (8.4)	6.94 d (8.1)				
6'	7.49 dd (2.1, 8.2)	7.60 dd (2.2, 8.3)	7.58 dd (2.0, 8.2)	7.50 dd (2.2, 8.3)	7.58 dd (1.9, 8.4)	7.45 dd (1.6, 8.1)				
1"	5.58 d (7.4)	5.60 d (7.5)	5.17 d (7.3)	5.62 d (7.5)	5.21 d (7.3)	5.27 d (7.1)				
2"	4.49 m	4.47 dd (7.6, 9.0)	3.70 m	4.47 dd (7.6, 9.0)	3.69 m	3.76 m				
3"	4.30 ov	4.29 m	3.61 m	4.28 m	3.61 m	3.69 m				
4"	4.21 m	4.19 m	3.61 m	4.75 m	3.60 m	3.48 m				
5"	3.80 t ( <i>10.6</i> )	3.78 m	3.48 m	3.80 m	3.49 m	3.72 m				
	4.30 ov	4.30 m	3.97 dd (4.0, 11.8)	4.31 m	3.98 m					
6"						3.66 m				
						4.05 m				
1'''	6.52 d ( <i>1.8</i> )	6.52 d ( <i>1</i> .2)	5.49 d (1.4)	6.38 d ( <i>1.6</i> )	5.28 d (1.5)	4.69 d (7.9)				
2'''	4.83 brs	4.84 brs	3.87 ov	4.80 m	3.96 m	3.25 m				
3'''	-	-	-	4.54 dd (3.3, 9.3)	3.61 m	3.42 m				
4'''	4.21 s	4.21 brs	4.08 brs	4.32 m	3.41 t (9.6)	3.39 m				

## Table 3.9 <sup>1</sup>H NMR data of compounds **163-168**

Position	$\delta_{\rm H} { m m} (J { m in} { m Hz})$								
	<b>163</b> <sup>a</sup>	<b>164</b> <sup>a</sup>	<b>165</b> <sup>b</sup>	<b>166</b> <sup>a</sup>	<b>167</b> <sup>b</sup>	<b>168</b> <sup>b</sup>			
5'''	4.45 d (9.6)	4.43 d (9.5)	3.87 ov	4.75 m	3.89 m	3.30 m			
	4.77 d (9.6)	4.76 d (9.5)	4.13 ov						
6'''		-		1.78 d (6.1)	1.32 d (6.2)	3.59 m			
						3.67 m			
1''''						4.72 d (1.1)			
2''''						3.91 m			
3''''						3.73 m			
4''''						3.35 m			
5''''						3.65 m			
6''''						1.19 d (6.1)			
OCH <sub>3</sub>		3.78 s	4.00 s	-	3.99 s				

a- ran in Pyr-d<sub>5</sub>; b- ran in CD<sub>3</sub>OD

Position				δ <sub>C</sub>		
	<b>163</b> <sup>a</sup>	<b>164</b> <sup>a</sup>	<b>165</b> <sup>b</sup>	<b>166</b> <sup>a</sup>	<b>167</b> <sup>b</sup>	<b>168</b> <sup>b</sup>
1	-	-	-	-	-	-
2	165.2	164.7	165.3	165.1	165.0	165.7
3	106.5	104.0	103.5	104.0	103.2	102.9
4	182.6	182.5	183.1	182.5	182.1	182.9
5	162.6	157.6	161.3	162.4	160.9	161.5
6	100.3	100.3	99.9	99.9	99.5	99.1
7	163.3	163.0	162.6	163.2	162.5	163.0
8	95.3	95.3	94.2	95.0	94.5	95.1
9	157.6	157.6	157.6	157.4	156.8	157.6
10	106.5	106.6	105.7	106.5	105.1	105.8
1'	122.3	122.3	122.0	122.3	121.5	122.2
2'	114.4	110.1	109.7	114.1	109.5	113.0
3'	147.5	148.5	148.0	147.6	147.7	149.9
4'	151.6	152.4	150.9	151.0	150.2	145.7
5'	116.6	116.7	115.4	116.5	115.3	115.9
6'	119.3	121.1	120.7	119.4	120.5	119.3
1"	100.3	100.3	99.3	100.2	99.0	98.5
2"	77.0	77.0	76.6	77.4	77.6	82.0
3"	78.3	78.2	69.7	78.6	70.7	75.7
4"	70.5	70.4	76.8	70.5	77.2	69.5
5"	67.0	66.7	65.7	67.0	65.6	76.7
6"						66.1
1'''	110.5	110.4	108.9	102.3	101.2	104.0
2""	77.6	77.7	77.2	72.1	70.7	74.6
3'''	80.6	80.7	nd	72.4	69.7	76.4
4'''	65.8	65.8	66.9	74.0	72.6	69.7
5'''	75.6	75.4	74.1	69.9	68.6	76.7
6'''				18.6	16.9	60.7
1''''						100.9
2''''						70.7
3''''						71.2
4''''						72.6
5''''						68.4
6''''						16.4
OCH <sub>3</sub>		55.8	55.0		55.4	
CH <sub>3</sub> CO			19.2			
COCH <sub>3</sub>			171.1			

Table 3.10 <sup>13</sup>C NMR data of compounds **165-170** 

Position	170		171	
	δ <sub>H</sub> m ( <i>J</i> )	δ <sub>C</sub>	δ <sub>H</sub> m ( <i>J</i> )	δ <sub>C</sub>
1	-	-	-	-
2	-	176.0	-	177.4
3	5.30 t ( <i>6.0</i> )	81.8	5.42 d (6.4)	83.4
4	1.80 ddd (5.9, 8.3, 14.3)	39.6	2.28 m	39.6
	2.91 ddd (6.2, 8.5, 14.3)		2.42 dd (7.9, 8.4)	
5	4.36 t (8.4)	69.5	4.67 t (8.2)	69.2
1'	-	115.9	-	116.5
2'	-	148.8	-	148.9
3'	6.61 s	101.7	6.63 s	102.0
4'	-	151.4	-	151.5
5'	-	142.5	-	142.6
6'	6.86 s	116.3	6.81 s	115.0
4'-OCH3	3.74 s	56.6	3.76 s	56.6
5'-OCH3	3.83 s	56.7	3.81 s	57.3

Table 3.11 <sup>1</sup>H and <sup>13</sup>C NMR data of **170** and **171** 

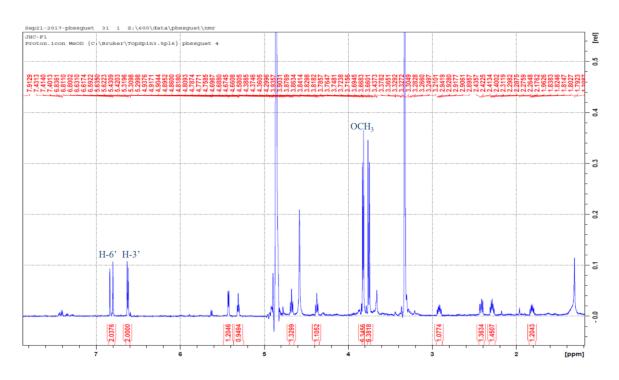


Figure 3.51 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) of **170** and **171** 

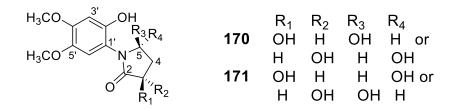
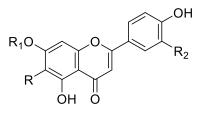


Figure 3.52 Key <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations **170** and **171** 

#### 3.3.3 Phytochemistry of Pseudospondias microcarpa

Chromatographic separation of the phytochemical constituents of the methanol and DCM extracts of the stem bark and leaves of *P. microcarpa* led to isolation of seven compounds (Figure 3.53). The compounds were characterised by spectroscopic means and by comparison of their spectral data with those of previously isolated compounds from our laboratory or with the literature. They were identified as *trans*-ferulic acid (**148**), chrysoeriol-7-*O*- $\beta$ -D-apiofuranosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (**164**), luteolin-7-*O*- $\alpha$ -L-rhamnopynosyl-(1 $\rightarrow$ 2)- $\beta$ -D-xylopyranoside (**166**), isovetexin (**172**), apigenin 7-O- $\beta$ -D-neohesperidoside (**173**), scopoletin (**103**) and pithecellobiumol B (**174**). All the isolated compounds belong to the class of phenolic compounds and their derivatives. To the best of our knowledge, this is the first report on the phytochemical studies of the genus *Pseudospondias* and the species *P. microcarpa*. The genus *Pseudospondias* derived from *Spondioidae* tribe of the Anarcadiaceae family (Mitchel and Daly, 2015), known to be a rich source of phenolic compounds including flavonoids, phenolic acid derivatives and tannins (Umadevi *et al.*, 1988; Sameh *et al.*, 2018).



**164** R= H R<sub>1</sub>=  $\beta$ -D-apiofuranosyl-(1" $\rightarrow$ 2")- $\beta$ -D-xylopyranosyl R<sub>2</sub>= OCH<sub>3</sub> **166** R= H R<sub>1</sub>=  $\alpha$ -L-rhamnopyranosyl-(1" $\rightarrow$ 2")- $\beta$ -D-xylopyranosyl R<sub>2</sub>= OH **172** R=  $\beta$ -D-glucopyranosyl R<sub>1</sub>= H R<sub>2</sub>= H **173** R= H R<sub>1</sub>=  $\alpha$ -L-rhamnosyl-(1" $\rightarrow$ 2")- $\beta$ -D-glucopyranosyl R<sub>2</sub>= H

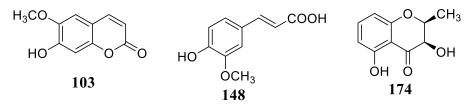


Figure 3.53 Structures of isolated compounds from P. microcarpa

#### 3.3.3.1 Structure elucidation of scopoletin (103)

Compound **103** was isolated as a light brown powder. Its molecular formula  $C_{10}H_8O_4$  was determined by the peak at m/z 215 [M+Na]<sup>+</sup>, from its ESI-MS spectrum obtained in positive mode ion. Its <sup>1</sup>H NMR spectrum (Figure 3.54, Table 3.12) showed characteristic signal of H-3 ( $\delta_H$  6.20, d, J = 9.4 Hz) and H-4 ( $\delta_H$  7.90, d, J = 9.4 Hz) protons of the pyrone ring of coumarins (Kayser and Kolodziej, 1995), two aromatic singlet methines at  $\delta_H$  6.77 and 7.19 consistent with a tetrasubstituted aromatic ring and a methoxy signal at  $\delta_H$  3.80. In the HMBC spectrum (Figure 3.54), long-range correlations were observed between the olefinic proton at  $\delta_H$  7.90 (H-1') and the carbons at  $\delta_C$  109.9 (C-2) and 149.8 (C-6) and between the methoxy proton at  $\delta_H$  3.80 and the carbon at  $\delta_C$  145.7 (C-3). The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.12) of **103** were in good agreement with those published for scopoletin (Darmawan *et al.*, 2012). Thus, compound **103** was characterised as scopoletin, a coumarin with a range of biological activities including antidiabetic, antimicrobial, anti-oxidant and antitumor (Zhao *et al.*, 2015; Napiroon *et al.*, 2018).

Position	$\delta_{\rm H} m (J \text{ in Hz})$	δ <sub>C</sub>	Position	$\delta_{\rm H} m (J \text{ in Hz})$	δ <sub>C</sub>
2	-	161.5	7	-	151.5
3	6.20 d (9.4)	112.0	8	7.19 s	109.9
4	7.90 d (9.4)	145.1	9	-	111.2
5	6.77	103.1	10	-	145.7
6	-	149.8	OCH <sub>3</sub>	3.80 s	56.4

Table 3.12 <sup>1</sup>H (600 MHz, DMSO-d<sub>6</sub>) and <sup>13</sup>C NMR (150 MHz, DMSO-d<sub>6</sub>) data of **103** 

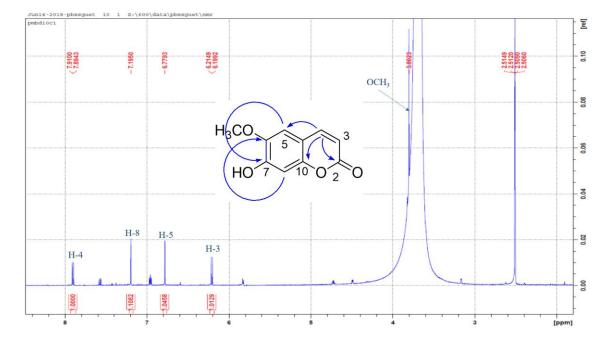


Figure 3.54 <sup>1</sup>H NMR (600 MHz, DMSO-d<sub>6</sub>) and key HMBC correlations of **103** 

## 3.3.3.2 Structure elucidation of isovetexin (172)

Compound **172** was isolated as a dark yellow amorphous powder. Its molecular formula  $C_{21}H_{20}O_{10}$  was confirmed by the peak at m/z 455 [M+Na]<sup>+</sup>, in its ESI-MS spectrum obtained in positive ion mode. The <sup>1</sup>H NMR spectrum (Figure 3.55, Table 3.13) showed

12 peaks belonging to a glycosylated molecule bearing one sugar moiety. The aglycone part of the compound was identified as apigenin by the presence of an  $A_2B_2$  system resonating at  $\delta_{\rm H}$  7.87 (2H, d, J = 8.8 Hz) and 6.95 (2H, d, J = 8.8 Hz) characteristic of the 1,4-disubstituted B ring; a singlet at  $\delta_{\rm H}$  6.63 attributable to the H-3 proton of the C ring and another singlet at  $\delta_{\rm H}$  6.54 ppm attributable to the H-6 or H-8 proton of the A ring suggesting an additional substituent on that ring at one of those positions. The sugar moiety was characterised by the presence of the methylene protons at  $\delta_{\rm H}$  3.75 (dd, J =5.3, 12.1 Hz) and 3.90 (dd, J = 2.3, 12.1 Hz) and four methines including the anomeric proton at  $\delta_{\rm H}$  4.92 (d, J = 9.9 Hz), which on the HSQC spectrum (Figure 3.56) correlated with the carbon at  $\delta_{\rm C}$  75.3 suggesting the direct C-C attachment of the sugar moiety to the aglycone ring. The observed <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.13) associated with the sugar moiety were in agreement with those published for  $\beta$ -D-glucose (Agrawal, 1992). In the HMBC spectrum (Figure 3.57), the singlet at  $\delta_{\rm H}$  6.54 showed long-range correlations with the carbons at  $\delta_{\rm C}$  105.2 (C-10), 158.7 (C-9) and 164.9 (C-7) and was therefore assigned to the proton at C-8 of the A ring. A  ${}^{3}J$  long-range correlations were observed between the anomeric proton of the glucosyl moiety ( $\delta_{\rm H}$  4.92, H-1'') and the carbon at  $\delta_{\rm C}$  109.3 (C-6), 162.0 (C-5) and 164.9 (C-7) confirming the direct attachment of the sugar carbon C-1" at C-6 by a C-C linkage. All the observed NMR data (Table 3.13) were consistent with those published for apigenin-6-C- $\beta$ -D-glucoside or isovitexin (Krafczyk *et al.*, 2008; Ganbaatar et al., 2015). Thus, compound 172 was identified as isovitexin. Isovitexin has been identified and isolated from several genus of Anarcadiaceae and possesses a range of biological properties including anti-inflammatory, antispasmodic, antimicrobial and antinociceptive (Picerno et al., 2006; Correa et al., 2012; Abu-Reidah et al., 2015).

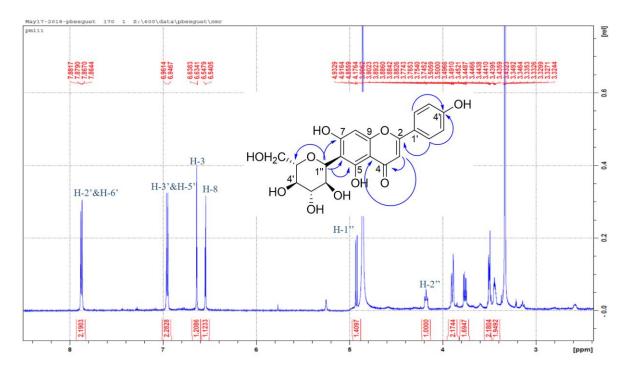


Figure 3.55  $^{1}$ H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **172** 

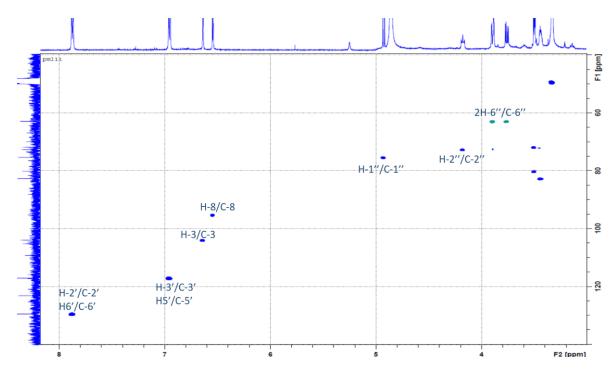


Figure 3.56 HSQC spectrum of 172

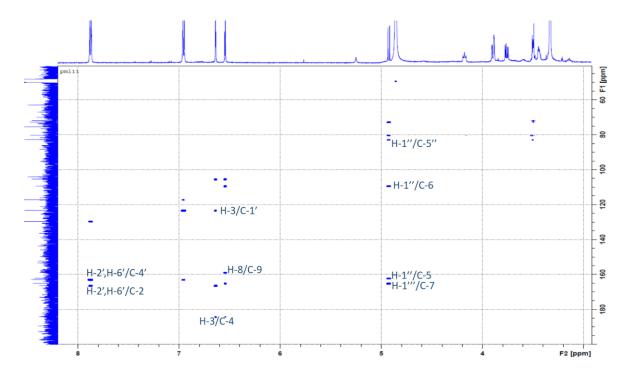


Figure 3.57 HMBC (600 MHz, CD<sub>3</sub>OD) of 172

### 3.3.3.3 Structure elucidation of apigenin 7-O- $\beta$ -D-neohesperidoside (173)

Compound **173** was isolated as a dark brown amorphous powder. Its molecular formula  $C_{27}H_{30}O_{14}$  was determined from the peak at m/z 601 [M+Na]<sup>+</sup>, in its ESI-MS spectrum in positive mode ion. Its <sup>1</sup>H NMR spectrum (Figure 3.58, Table 3.13) suggested a glycosylated apigenin bearing two sugar moieties payable to the presence of two anomeric protons at  $\delta_H$  5.30 (d, J = 1.7 Hz) and 5.22 (d, J = 7.6 Hz) and the proton signals at  $\delta_H$  7.91 (2H, d, J = 8.8 Hz), 6.95 (2H, d, J = 8.8 Hz), 6.82 (1H, d, J = 2.2 Hz), 6.68 (1H, s) and 6.48 (1H, d, J = 2.2 Hz) attributable to the apigenin aglycone. The two sugar moieties were identified as  $\alpha$ -L-rhamnose and  $\beta$ -D-glucose by comparison of the observed data with those published for the respective molecules (Osterdahl, 1979; Beier and Mundy, 1980). In the HMBC spectrum (Figure 3.58), a correlation observed between the anomeric proton of the glucosyl unit ( $\delta_H$  5.22, H-1'') and the carbon at  $\delta_C$  162.5 (C-7) confirmed its direct attachment to the aglycone. While, the correlation of the rhamnosyl

anomeric proton ( $\delta_{\rm H}$  5.30, H-1''') with the carbon C-2'' of the glycosyl at  $\delta_{\rm C}$  77.6 led to the fixation of the rhamnosyl unit on the carbon C-2'' of the glycoside. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.13) of **173** and the key HMBC correlations (Figure 3.58) were in good agreement with those published for apigenin 7-O- $\beta$ -D-neohesperidoside (Osterdahl, 1979; Zhang *et al.*, 2016). Thus, compounds **173** was identified as apigenin 7-O- $\beta$ -Dneohesperidoside, a flavone glycoside with antioxidant properties (Zhang *et al.*, 2016).

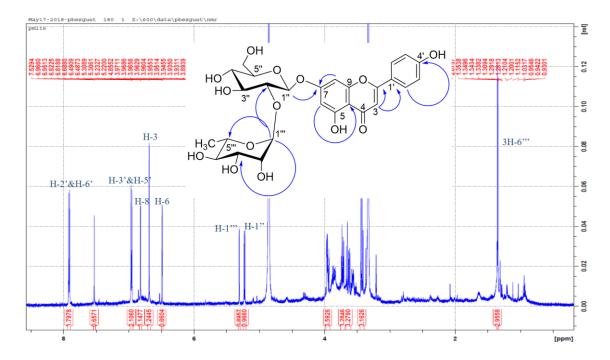


Figure 3.58 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **173** 

## 3.3.3.4 Structure elucidation of pithecellobiumol B (174)

Compound **174** was isolated as a brown amorphous powder. Its molecular formula  $C_{10}H_{10}O_4$  was determined from the *pseudo*molecular peak at m/z 217 [M+Na]<sup>+</sup>, in its ESI-MS spectrum. Its <sup>1</sup>H NMR spectrum (Figure 3.59, Table 3.14) depicted seven peaks which could be attributed to five methines including three aromatic methine at  $\delta_H$  7.58 (dd, J = 7.3, 8.3 Hz), 6.97 (dd, J = 0.9, 8.3 Hz) and 6.96 (dd, J = 0.9, 7.3 Hz) and two

oxymethines at  $\delta_{\rm H}$  4.75 (dq, J = 2.1, 6.6 Hz) and 4.50 (dd, J = 2.1, 6.3 Hz); a doublet methyl at  $\delta_{\rm H}$  1.40 (J = 6.6 Hz) and a doublet proton at  $\delta_{\rm H}$  5.71 (d, J = 6.3 Hz) which was attributed to the proton of a hydroxyl group as no cross peak correlation was observed for this signal in the HSQC-DEPT spectrum. In the HMBC (Figure 3.59), long-range correlations were observed between the methine at  $\delta_{\rm H}$  4.50 (H-3) and the carbon at  $\delta_{\rm C}$ 107.7 (C-10) and 142.8 (C-9) and between the methine at  $\delta_{\rm H}$  7.58 (H-7) and the carbons at  $\delta_{\rm C}$  142.8 (C-8a) and 160.6 (C-5). The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.14) of compound **174** were similar to those published for pithecellobiumol B (Wang *et al.*, 2017). In addition, pithecellobiumol B and **174** were found to have quite identical <sup>1</sup>H NMR spectra when ran in DMSO. Thus, **174** was recognised as pithecellobiumol B, a chromanone previously isolated from the leaves and twigs of *Pithecellobium clypearia* (Fabaceae) (Wang *et al.*, 2017).

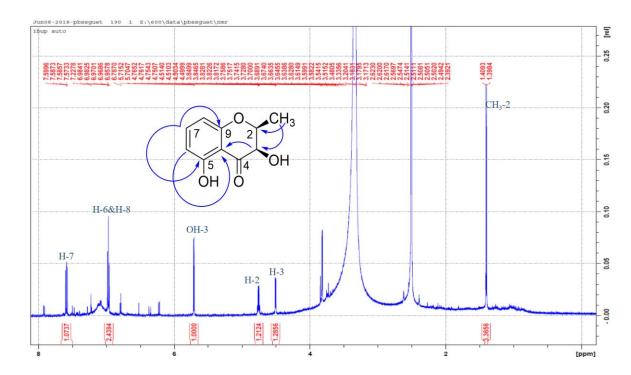


Figure 3.59 <sup>1</sup>H NMR (600 MHz, DMSO-d<sub>6</sub>) and key HMBC correlations of **174** 

Position	$\delta_{\mathrm{H}} m$ (	(J in Hz)		δ <sub>C</sub>
	172	173	172	173
1	-	-	-	-
2	-	-	166.3	164.8
3	6.63 s	6.68 s	103.9	102.7
4	-	-	183.9	182.1
5	-	-	162.0	nd
6	-	6.48 d (2.2)	109.3	99.5
7	-	-	164.9	162.5
8	6.54 s	6.82 d (2.2)	95.4	94.7
9	-	-	158.7	157.0
10	-	-	105.2	105.5
1'	-	-	123.1	121.4
2'	7.87 d (8.8)	7.91d (8.8)	129.4	128.6
3'	6.95 d (8.8)	6.95 d (8.8)	117.0	115.9
4'	-	-	162.8	161.3
5'	6.95 d (8.8)	6.95 d (8.8)	117.0	115.9
6'	7.87 d (8.8)	7.91d (8.8)	129.4	128.6
1"	4.92 d (9.9)	5.22 d (7.6)	75.3	98.4
2"	4.17 brt (9.3)	3.71 ov	72.6	77.6
3"	3.49 ov	3.65 ov	80.1	77.6
4"	3.47 ov	3.42 m	71.8	72.6
5"	3.44 m	3.55 m	82.8	76.9
6"	3.75 dd (5.3, 12.1)	3.73 m	62.8	61.1
	3.90 dd (2.3,12.1)	3.94 m		
1""		5.30 d (1.7)		101.3
2""		3.97 m		70.7
3'''		3.62 m		70.7
4""		3.43 m		70.1
5'''		3.96 m		68.4
6'''		1.34 d (6.2)		16.8

Table 3.13  $^{1}$ H (600 MHz, CD<sub>3</sub>OD) and  $^{13}$ C NMR (75 MHz, CD<sub>3</sub>OD) data of **172** and **173** 

nd not determined

Position	$\delta_{\rm H}  {\rm m}  (J  {\rm in}  {\rm Hz})$	$\delta_{\mathrm{C}}$	Position	$\delta_{\rm H} { m m}  (J { m in} { m Hz})$	$\delta_{C}$
1	-	-	8	6.97 dd (0.9, 8.3)	117.6
2	4.75 dq (1.9, 6.6)	79.2	9	-	142.8
3	4.50 dd (2.1, 6.3)	65.8	10	-	107.7
4	-	nd	2-CH3	1.40 d (6.6)	16.1
5	-	160.6	3-ОН	5.71 d (6.3)	-
6	6.96 dd (0.9, 7.3)	119.6	5-OH	10.9 brs	-
7	7.58 dd (7.3, 8.3)	137.3			

Table 3.14 <sup>1</sup>H (600 MHz, DMSO-d<sub>6</sub>) and <sup>13</sup>C NMR (150 MHz, DMSO-d<sub>6</sub>) data of **174** 

nd not determined

#### 3.3.4 Phytochemistry of Zanthoxylum leprieurii

Silica gel CC and RP-HPLC analyses of the *n*-hexane and DCM extracts of the fruits of *Z. leprieurii* afforded nine compounds including five kaurane diterpenes i.e., kaurenoic acid (**175**), xylopic acid (**176**), *ent*-kauran-16 $\beta$ -ol-19-oic acid (**177**), *ent*-kauran-16 $\beta$ -ol-19-al (**178**) and *ent*-kauran-16 $\beta$ -ol (**179**); two flavanes i.e., dulcisflavane (**180**) and epicatechin (**181**); caffeic acid (**114**) and icariside D2 (**182**) (Figure 3.60) along with a mixture of sterols. All the above listed compounds are here reported from the first time from *Z. leprieurii*. Compounds **114** and **180-182** have previously been reported from other *Zanthoxylum* species including *Z. piperitum*, *Z. schinifolium* (Kusuda *et al.*, 2006; Fang *et al.*, 2010). This is the first report on the occurrence kaurane diterpenoids (**175-179**) in the genus *Zanthoxylum*. Kaurane diterpenes are mainly found in the Asteraceae, Annonaceae and Euphorbiaceae families among others (Garcia *et al.*, 2007). First occurrence of kaurane diterpenes in the Rutaceae was the isolation of kaurenoic acid and its 15 $\beta$ -hydroxylated derivative from the genus *Fortunella* (El-Shafae and Ibrahim, 2003).

Diterpenes are rather rare in the Rutaceae and only a few have been reported from the genus *Citrus*, *Evodia*, *Glycosmis* and *Pamburus* (Dreyer and Park, 1975; Seger *et al.*, 1998; Garcia *et al.*, 2007).

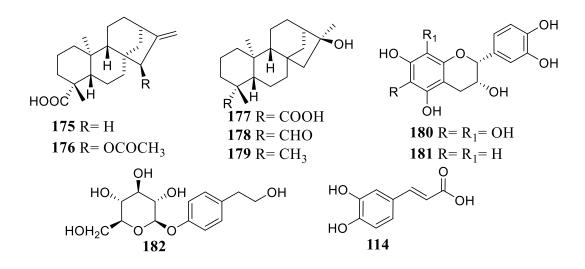


Figure 3.60 Isolated compounds from the fruits of Z. leprieurii

## 3.3.4.1 Structure elucidation of kaurenoic acid (175)

Compound **175** was isolated as white crystalline powder from the hexane extract of *Z*. *leprieurii*. Its molecular formula C<sub>20</sub>H<sub>30</sub>O<sub>2</sub> was determined from its HR ESI-MS spectrum (Figure 3.61) obtained in negative ion mode where the peak at m/z 301.2170 [M-H]<sup>-</sup> calculated for C<sub>20</sub>H<sub>29</sub>O<sub>2</sub>, 301.2173 was observed. The <sup>1</sup>H NMR spectrum (Figure 3.62, Table 3.15) of **175** showed characteristic signals of two tertiary methyls at  $\delta_{\rm H}$  0.92 (s, H-20) and 1.25 (s, H-18), two singlets at  $\delta_{\rm H}$  4.67 and 4.72 (1H each) typical of the methylene protons (H-17) of an exocyclic double bond and one methine proton at  $\delta_{\rm H}$  2.56 (brs, H-13). The <sup>13</sup>C NMR spectrum (Figure 3.62, Table 3.15) depicted 20 peaks which could be attributed to two methyls at  $\delta_{\rm C}$  15.6 (C-20) and 28.9 (C-18); ten methylenes at  $\delta$  103.0 (C-17), 48.9 (C-15), 41.3 (C-7), 40.7 (C-1), 39.7 (C-14), 37.8 (C-3), 33.1 (C-12), 21.8 (C-6), 19.1 (C-2) and 18.4 (C-11); three methines at  $\delta_{\rm C}$  57.0 (C-5), 55.1 (C-9) and 43.7

(C-13) and five quaternary carbons including a carboxylic acid function at  $\delta_{\rm C}$  184.6 (C-19) and an olefinic carbon at  $\delta_{\rm C}$  155.9 (C-16) suggesting **175** was a diterpene (Pacheco *et al.*, 2009). In the HMBC spectrum (Figure 3.64), long-range correlations could be observed between the olefinic protons H-17 and the carbons C-13, C-15 and C-16 and between the methine proton H-13 and the carbons C-8, C-12, C-15, C-16 and C-17. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.15) of **175** and the key correlations observed (Figure 3.64) were in good agreement with those published for *ent*-kaurenoic acid (Mitscher *et al.*, 1983; Viera *et al.*, 2002). Thus, compound **175** was identified as *ent*-kaurenoic acid, a kaurene diterpene with a range of biological activities including analgesic, antidiabetic, anti-inflammatory, antioxidant and neurological activities (Villa-Ruano *et al.*, 2016).

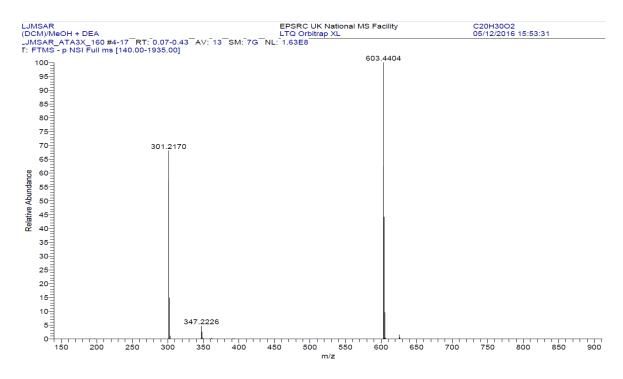


Figure 3.61 ESI-MS spectrum of 175

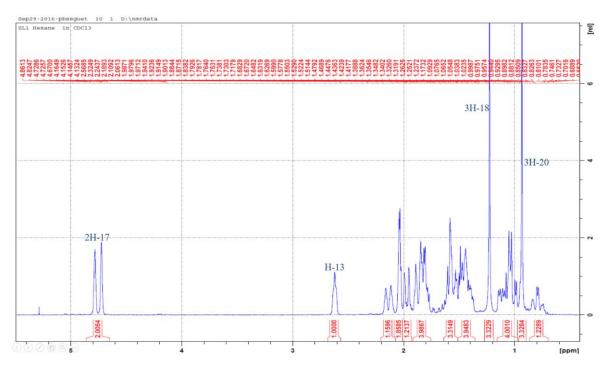


Figure 3.62  $^1\text{H}$  NMR (300 MHz, CDCl<sub>3</sub>) of 175

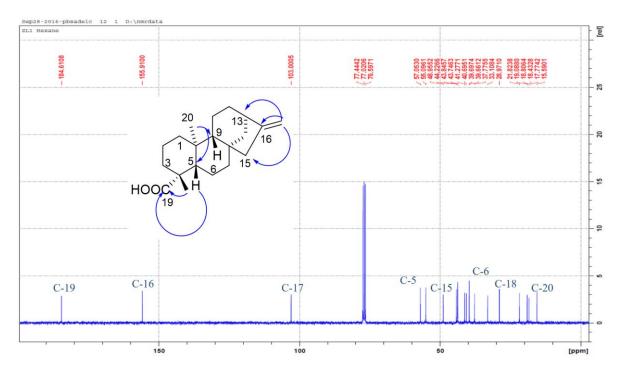


Figure 3.63 <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **175** 

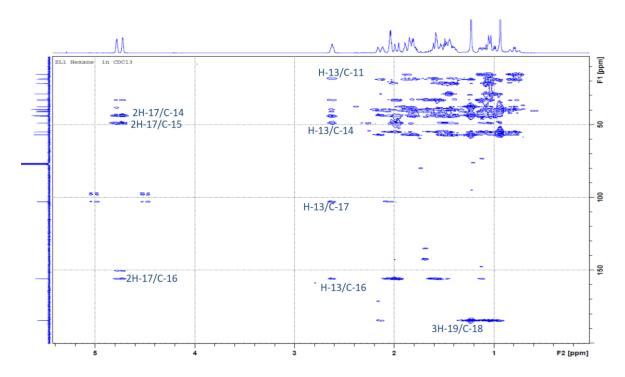


Figure 3.64 HMBC spectrum of 175

## 3.3.4.2 Structure elucidation of xylopic acid (176)

Compound **176** was also isolated as white crystalline powder. Its molecular formula  $C_{22}H_{32}O_4$  was determined from its HR ESI-MS spectrum by the peak at m/z 361.2375 [M+H]<sup>+</sup> calculated for  $C_{22}H_{33}O_4$ , 361.2379. Its <sup>1</sup>H NMR spectrum (Figure 3.65, Table 3.15) was similar to that of **175**. The only difference was the presence in the <sup>1</sup>H NMR spectrum of **176** of an additional methyl singlet at  $\delta_H$  2.26 suggesting the presence of an acetoxy function in the molecule, and an oxymethine at  $\delta_H$  5.16 (t, J = 2.5, 4.9 Hz). This was supported in the <sup>13</sup>C NMR spectrum (Figure 3.66, Table 3.15) by the peaks at  $\delta_C$  171.4 and 21.3 corresponding to the carbonyl and methyl of the acetoxy respectively and the oxymethine carbon at  $\delta_C$  81.6. In the HMBC spectrum (Figure 3.66), correlations observed between the olefinic methylene protons (H-17) and the carbons at  $\delta_C$  40.6 (C-13), 81.6 (C-15) and 153.7 (C-16) and between the oxymethine H-15 and the carbons at  $\delta_C$  45.9 (C-8), 106.1 (C-17), 153.7 (C-16) and 171.4 led to the fixation of the acetoxy 162

function on the carbon C-15 at  $\delta_{\rm C}$  81.6. The orientation of the acetoxy function on C-15 was determined to be  $\beta$ -oriented using the NOESY experiment. All the <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.15) of **176** supported by the key HMBC correlations observed were in agreement with those published for *ent*-15 $\beta$ -acetoxy-kaur-16-en-19-oic acid or xylopic acid (Takahashi *et al.*, 2001). This was further confirmed in the MS spectrum of **176** by the peak at *m*/*z* 301.2245 which correspond to the loss of the acetoxy function [M-OAc+H]<sup>+</sup>. Thus, compound **176** was identified as xylopic acid, a kaurene diterpene widespread in the *Xylopia* genus of the Annonaceae family (Takahashi *et al.*, 2001; Silva *et al.*, 2015).

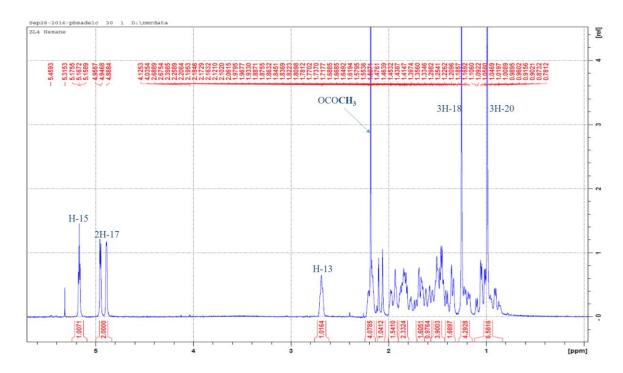


Figure 3.65 <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>) of **176** 

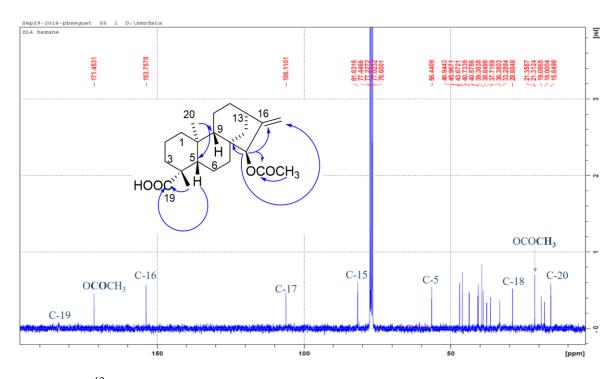


Figure 3.66<sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **176** 

# 3.3.4.3 Structure elucidation of *ent*-kauran-16 $\beta$ -ol-19-oic acid (177)

Compound **177** was isolated as a white granular powder. Its molecular formula  $C_{20}H_{32}O_3$  was confirmed from its HR ESIMS spectrum obtained in negative ion mode by the peak at m/z 319.2273 [M-H]<sup>-</sup> calculated for  $C_{20}H_{31}O_3$ , 319.2279. Its <sup>1</sup>H NMR spectrum (Figure 3.67, Table 3.15) was similar to that of **175**. The only difference was the absence in the <sup>1</sup>H NMR spectrum of **177** of the olefinic methylene protons suggesting an oxidation, and the presence of an additional tertiary methyl singlet at  $\delta_H$  1.46. This was supported in its <sup>13</sup>C NMR spectrum (Figure 3.68, Table 3.15) by the absence of the olefinic carbons and the presence of a quaternary oxygenated carbon at  $\delta_C$  78.5 and a methyl at  $\delta_C$  23.2 which show in the HSQC spectrum cross peak correlations could be observed between the methyl signal at  $\delta_H$  1.46 (H-17) and the carbons at  $\delta_C$  48.8 (C-13), 58.0 (C-15) and 78.5 (C-16) and between the methine proton at  $\delta_H$  1.76 (H-13) and the carbons  $\delta_C$  78.5 (C-16),

58.0 (C-15), 45.4 (C-8), 26.2 (C-12) and 23.2 (C-17). The orientation of the hydroxy function on C-16 was determined to be  $\beta$ -oriented using the NOESY experiment. All the <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.15) of **177** supported by the key HMBC correlations observed were in good agreement with those published for *ent*-16 $\beta$ -hydroxykauran-19-oic acid (Takahashi *et al.*, 1995). Thus, compound **177** was identified as *ent*-16 $\beta$ -hydroxykauran-19-oic acid, previously isolated from *Xylopia frutescens* (Takahashi *et al.*, 1995).

## 3.3.4.4 Structure elucidation of *ent*-kauran-16 $\beta$ -ol (**179**)

Compound 179 was isolated as white powder. Its molecular formula  $C_{20}H_{35}O$  was determined from its HR ESI-MS spectrum obtained in positive ion mode by the peak at m/z 290.2843 [M-H<sub>2</sub>O+NH<sub>4</sub>]<sup>+</sup> calculated for C<sub>20</sub>H<sub>32</sub>NH<sub>4</sub>, 290.2848. Its <sup>1</sup>H NMR spectrum (Figure 3.69, Table 3.15) was similar to that of **177**. The only difference was the presence of an additional tertiary methyl singlet at  $\delta_{\rm H}$  0.78 in the <sup>1</sup>H NMR spectrum of **179**. In its <sup>13</sup>C NMR spectrum (Figure 3.70, Table 3.15), the presence of an additional methyl signal at  $\delta_{\rm C}$  21.5 and the absence of the carboxylic acid carbon were clearly visible. In the HMBC spectrum (Figure 3.70), long-range correlations could be observed between the methyl signal at  $\delta_{\rm H}$  0.78 (H-19) and the carbons at  $\delta_{\rm C}$  18.6 (C-2), 33.2 (C-4), 33.6 (C-18), 42.0 (C-3) and 56.2 (C-5). All the <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.15) of **179**, supported by its key HMBC correlations, were in agreement with those published for ent-kauran- $16\beta$ -ol (Morris *et al.*, 2005). This was further confirmed in the MS spectrum of **179** by the peak at m/z 273.2578 [M-H<sub>2</sub>O+H]<sup>+</sup> arising from the loss of a water molecule due to the dehydration of the tertiary alcohol function. Thus, compound 179 was identified as ent-kauran-16 $\beta$ -ol, a kaurane diterpene previously isolated from sunflowers (Morris et al., 2005).

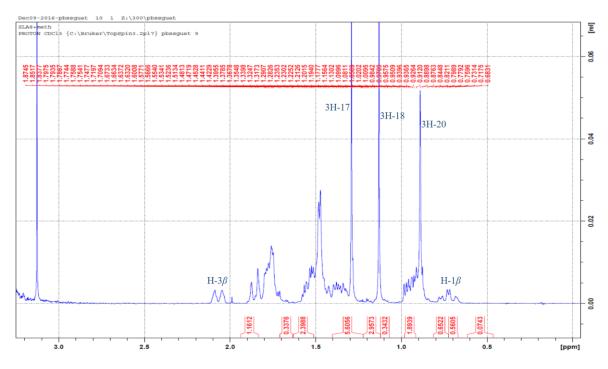


Figure 3.67  $^{1}$ H NMR (300 MHz, CDCl<sub>3</sub>+ drops MeOH) of **177** 

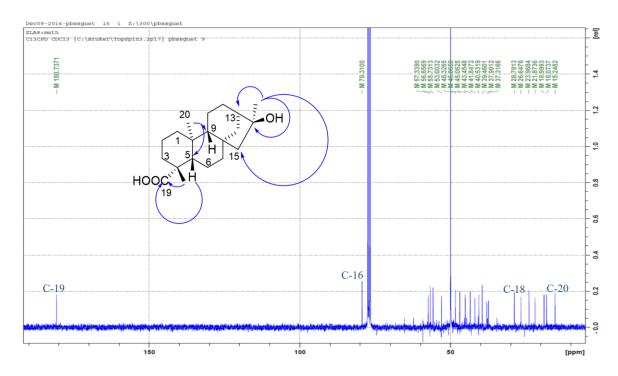


Figure 3.68 <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>+ drops MeOH) and key HMBC correlations of

177

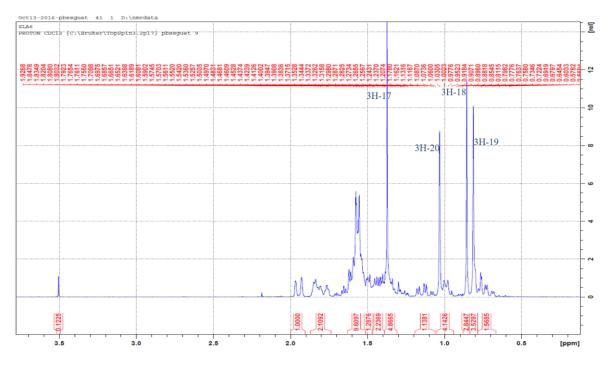


Figure 3.69  $^{1}$ H NMR (300 MHz, CDCl<sub>3</sub>) of **179** 

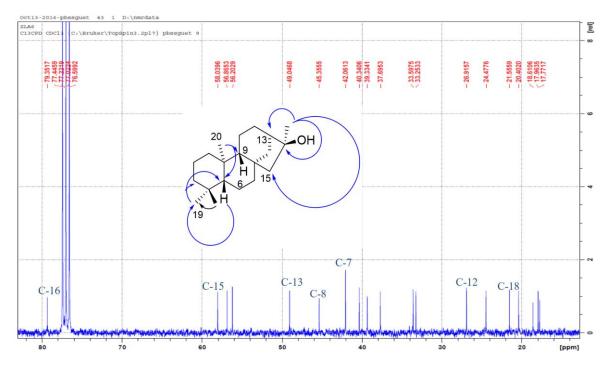


Figure 3.70 <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **179** 

Position	$\delta_{\rm H} { m m}  (J { m in} { m Hz})$									δ <sub>C</sub>							
	175		176			177 <sup>a</sup>			178 <sup>b</sup>			179	175	176	177 <sup>a</sup>	178 <sup>b</sup>	179
1	0.75 ddd 12.9, 5.5)	(13.0,	0.91 8.8, 6		(13.1,	0.69 <i>12.0</i> ,	ddd 3.0)	(15.0,	0.84 <i>13.6</i> ,		(17.0,	0.67 m	40.7	40.7	40.5	39.7	40.3
	1.76 m		1.97	m		1.76 r	n		1.86	m		1.76 m					
2	1.32 m		1.49	m		1.35 m 1		1.48	m		1.40 m	19.1	19.0	18.9	18.3	18.6	
	1.79 m	1.79 m 1.93 m 1.83 m		1.66 m 1.63 m			1.63 m										
3	0.94 m	0.94 m 1.08 m		0.96 m		1.05 m 1.08 m			37.8	38.8	37.3	34.2	42.0				
	2.06 dm		2.18	m		2.08 dm		2.16 dm		1.46 m							
4													43.8	43.6	48.3	48.4	33.2
5	1.03 m		1.04	m		0.96 r	0.96 m		1.17 dd (2.5, 2.1)		5, 2.1)	0.97 m	57.0	56.4	56.6	56.6	56.2
6	1.71 m		1.87	m		1.76 m 1.72 m 1.28 m			1.28 m	21.8	21.3	21.8	20.0	20.4			
			2.26	m					1.91	m		1.59 m					
7	1.40 m 1.50 m			1.39 m		1.52 d ( <i>3.8</i> )		1.31 m	41.3	39.9	43.4	41.9	42.0				
						1.46 r	n		1.72	m		1.65 m					
8													44.2	45.9	45.0	45.1	45.3
9	0.97 m		1.35	m		0.93 r	n		1.05	m		0.75 m	55.1	46.9	55.7	55.4	56.8
10													39.6	36.3	39.4	39.3	39.3
11	1.43 m		1.55	m		1.46 r	n		1.60	m		1.57 m	18.4	18.0	18.0	18.0	17.9
			1.66	m													

Table 3.15  $^{1}$ H NMR (CDCl<sub>3</sub>, 300 MHz) and  $^{13}$ C NMR (CDCl<sub>3</sub>, 75 MHz) data of **175-179** 

Position	1		$\delta_{\rm H} { m m}  (J { m in} { m Hz})$					$\delta_{C}$		
	175	176	1 <b>77</b> <sup>a</sup>	178 <sup>b</sup>	179	175	176	177 <sup>a</sup>	178 <sup>b</sup>	179
12	1.38 m	1.61 m	1.46 m	1.53 m	1.61 m	33.1	33.2	26.6	26.6	26.9
	1.47 m	1.73 m		1.62 m						
13	2.56 bs	2.68 bm	1.76 m	1.88 m	1.84 m	43.7	40.6	48.3	48.8	49.0
14	1.09 m	1.29 m	1.46 m	1.64 m	1.61 m	39.7	37.7	37.9	37.7	37.6
	1.90 d ( <i>3</i> .7)	2.16 d (3.5)	1.83 m	1.94 m	1.93 m					
15	1.98 bs	5.16 t (2.5, 4.9)	1.46 bs	1.60 bs	1.55 bs	48.9	81.6	57.3	57.6	58.0
16						155.9	153.7	79.3	79.3	79.3
17	4.67 s	4.88 bs	1.12 s	1.39 s	1.38 s	103.0	106.1	23.9	24.5	24.4
	4.72 s	4.94 d (2.6)								
18	1.25 s	1.33 s	0.93 s	1.05 s	0.83 s	28.9	28.8	28.7	24.2	33.6
19	-	-	-	9.75 d (1.4)	0.78 s	184.6	184.2	180.7	205.9	21.5
20	0.93 s	1.01 s	0.69 s	0.89 s	1.03 s	15.6	15.8	15.2	16.4	17.7
CH <sub>3</sub> CO		2.26 s					21.3			
CH <sub>3</sub> CO							171.4			

Table 3.15 *continued* 

<sup>a</sup> recorded in CDCl<sub>3</sub>+drops of MeOH; <sup>b</sup> recorded at 600 MHz for <sup>1</sup>H and 150 MHz for <sup>13</sup>C

Compound 180 was isolated as a brown amorphous powder. Its molecular formula  $C_{15}H_{14}O_8$  was determined from its ESI-MS spectrum obtained in positive ion mode, where the sodiated ion peak was observed t m/z 345 [M+Na]<sup>+</sup>. Its <sup>1</sup>H NMR spectrum (Figure 3.71, Table 3.16) revealed the presence of three aromatic protons at  $\delta_{\rm H}$  6.77 (d, J = 8.1 Hz), 6.81 (dd, J = 1.6, 8.1 Hz) and 6.99 (d, J = 1.6 Hz) members of a 1,3,4trisubstituted benzene as well as two oxygenated methines at  $\delta_{\rm H}$  4.19 and 4.83 and a set of methylene protons at  $\delta_{\rm H}$  2.75 (dd, J = 2.8, 16.5 Hz) and 2.88 (dd, J = 4.5, 16.5 Hz). The <sup>13</sup>C NMR spectrum (Figure 3.72, Table 3.16) of **180** depicted thirteen peaks which could be assigned based on the correlation observed in the HSQC and HMBC spectra (Figures 3.73-3.74) to one methylene, five methines and nine quaternary carbons. In the HMBC spectrum (Figure 3.72), correlations could be observed between the methylene protons (H-4) and the carbons at  $\delta_{\rm C}$  67.5 (C-3), 79.8 (C-2), 100.2 (C-10) and 157.6 (C-5) and between the oxymethine at  $\delta_{\rm H}$  4.83 (H-2) and the carbons at  $\delta_{\rm C}$  29.3 (C-4), 67.5 (C-3), 115.3 (C-2'), 119.5 (C-6'), 132.5 (C-1') and 157.3 (C-9). The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 16) of 180 were in good agreement with those published for dulcisflavan (Deachathai et al., 2005). Thus, compound 180 was identified as dulcisflavan, a flavan with antioxidant activity previously isolated from the fruit of Garcinia dulcis (Deachathai et al., 2005).

#### 3.3.4.6 Structure elucidation of icariside D2 (182)

Compound **182** was isolated as a dark brown amorphous powder. Its molecular formula  $C_{15}H_{14}O_8$  was determined from its pseudomolecular ion peak at m/z 345 [M+Na]<sup>+</sup> in the ESI-MS spectrum obtained in positive ion mode Its <sup>1</sup>H NMR spectrum (Figure 3.75,

Table 3.17) suggested a glycosylated compound bearing a sugar moiety by the presence of an anomeric proton at  $\delta_{\rm H}$  4.90 (d, J = 7.5 Hz). Four aromatic protons forming an A<sub>2</sub>B<sub>2</sub> system at  $\delta_{\rm H}$  7.18 (d, J = 8.7 Hz) and 7.06 (d, J = 8.7 Hz) suggesting a 1,4-disubstituted benzene, five oxygenated methines as well as three methylenes including two oxygenated ones at  $\delta_{\rm H}$  4.69 (t, J = 7.0 Hz) and 3.90 (dd, J = 2.2, 12.1 Hz) and 3.71 (dd, J = 5.5, 12.1 Hz) were also observed. In the COSY spectrum, a chain of correlation from the anomeric proton at  $\delta_{\rm H}$  4.90 to the methylenes at  $\delta_{\rm H}$  3.90 and 3.71 led to the identification of the sugar moiety as  $\beta$ -D-glucoside. Their corresponding carbons were determined by analysis of the <sup>13</sup>C (Figure 3.76, Table 3.17) and HSQC-DEPT spectra. The obtained NMR data of the sugar moiety (Table 3.17) were consistent with those published for  $\beta$ -Dglucopyranoside (Agrawal, 1992). In the HMBC spectrum (Figure 3.76), correlations could be observed between the anomeric proton at  $\delta_{\rm H}$  4.90 (H-1') and the carbon at  $\delta_{\rm C}$ 158.3 (C-4), 78.1 (C-5'), 78.0 (C-3') and 71.4 (C-10) and between the methylene at  $\delta_{\rm H}$ 3.24 (t, J=7.0 Hz) and the carbons at  $\delta_{\rm C}$  131.7 (C-1), 130.7 (C-2, C-6) and 77.5 (C- $\beta$ ). All the <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.17) of **182** supported by the observed key HMBC correlations, were comparable with those published for 4-(2-hydroxyethyl)phenyl- $\beta$ -Dglucopyranoside or icariside D2 (Miyase et al., 1989). Thus, compound 182 was identified as icariside D2, a lignan glycoside with cytotoxic properties first isolated from Epimedium diphyllum (Berberidaceae) (Miyase et al., 1989; Hien et al., 2015).

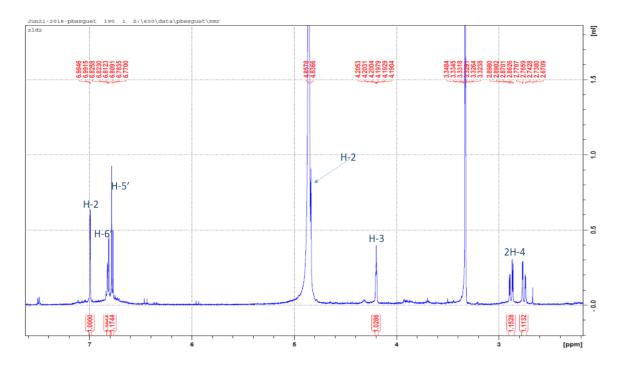


Figure 3.71 <sup>1</sup>H NMR (300 MHz, CD<sub>3</sub>OD) of **180** 

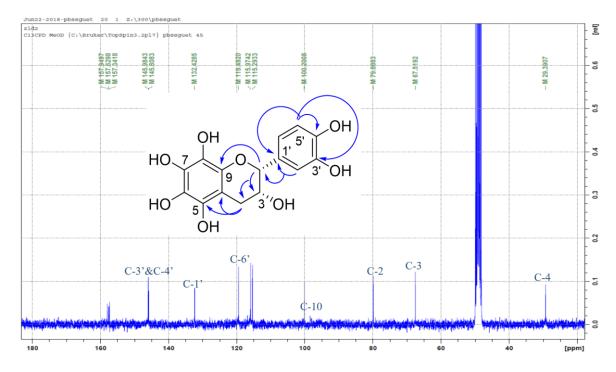


Figure 3.72 <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **180** 

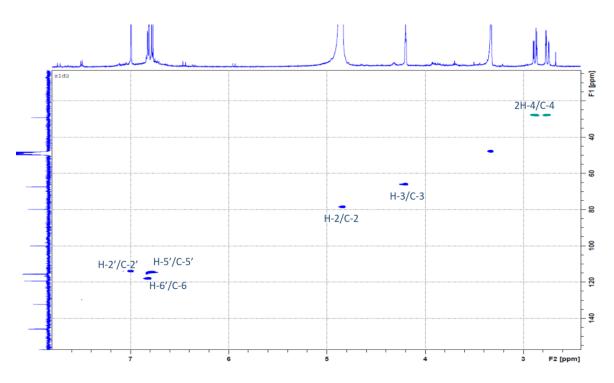


Figure 3.73 HSQC spectrum of 180

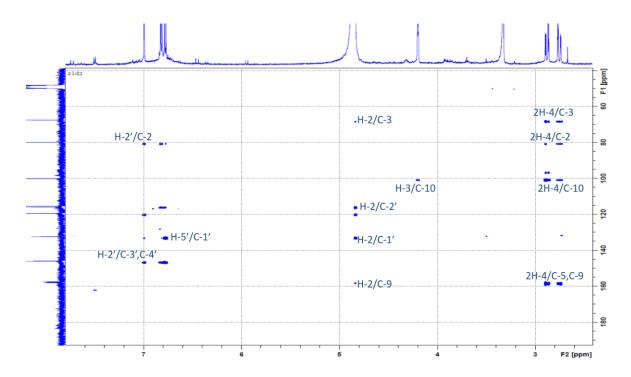


Figure 3.74 HMBC spectrum of 180

Position	$\delta_{\rm H} \ {\rm m} \ (J \ {\rm in} \ {\rm Hz})$	$\delta_{C}$	Position	$\delta_{\rm H}$ m ( <i>J</i> in Hz)	$\delta_{\mathrm{C}}$
1	-	-	9	-	157.3
2	4.86 ov	79.8	10	-	100.2
3	4.19 m	67.5	1'	-	132.4
4	2.75 dd (2.8, 16.5)	29.3	2'	6.99 d ( <i>1.6</i> )	115.3
	2.88 dd (4.5, 16.5)		3'	-	145.8
5	-	157.3	4'		145.9
6	-	132.4	5'	6.77 d (8.1)	115.9
7	-	157.6	6'	6.81 dd ( <i>1.6</i> , <i>8.1</i> )	119.5
8	-	157.9			

Table 3.16  $^{1}$ H (600 MHz, CD<sub>3</sub>OD) and  $^{13}$ C NMR (75 MHz, CD<sub>3</sub>OD) data of **180** 

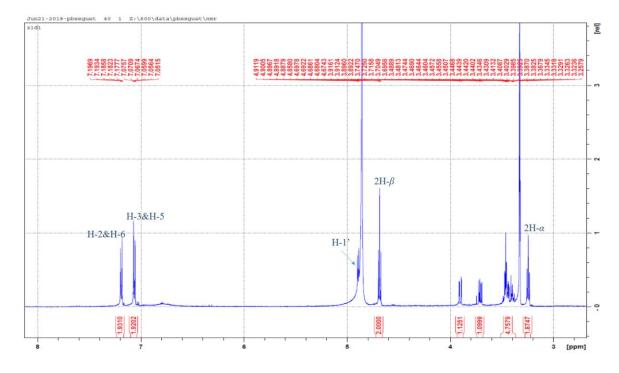


Figure 3.75  $^{1}$ H NMR (600 MHz, CD<sub>3</sub>OD) of **182** 

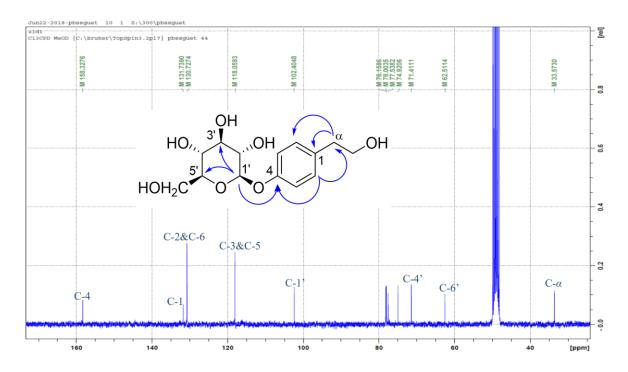


Figure 3.76<sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **182** 

Position	$\delta_{\rm H}$ m ( <i>J</i> in Hz)	δ <sub>C</sub>	Position	$\delta_{\rm H}$ m ( <i>J</i> in Hz)	δ <sub>C</sub>
1		131.7	3'	3.47 ov	78.0
2	7.18 d (8.7)	130.7	4'	3.40 ov	71.4
3	7.06 d (8.7)	118.0	5'	3.44 ov	78.1
4		158.3	6'	3.71 dd (2.2, 12.1)	62.5
5	7.06 d (8.7)	118.0		3.90 dd (5.5, 12.1)	
6	7.18 d (8.7)	130.7	C-α	3.24 t (7.0)	33.5
1'	4.90 d (7.5)	102.4	С-β	4.69 t (7.0)	77.5
2'	3.46 ov	74.9			

Table 3.17  $^{1}$ H (600 MHz, CD<sub>3</sub>OD) and  $^{13}$ C NMR (75 MHz, CD<sub>3</sub>OD) data of **182** 

### 3.3.5 Phytochemistry of Zanthoxylum zanthoxyloides

Preparative RP-HPLC and TLC analyses of the DCM and MeOH extracts of the fruits *Z*. *xanthoxyloides* afforded fourteen compounds including eight quinoline alkaloids, skimmianine (55), atanine (128), *N*-methylplatydesminium cation (183), isoplatydesmine (184), myrtopsine (185), ribalinine (186) and *N*-methylatanine (187); five alkamides, *trans*-fagaramide (79), zanthoamides G-I (188-190) and; a lignan, sesamin (54); and two flavanones, hesperidin (59) and hesperetin (191) (Figure 3.77). Compounds 188-190 are new alkamides, whilst compounds 183-187 are here reported for the first time from *Z*. *zanthoxyloides*.

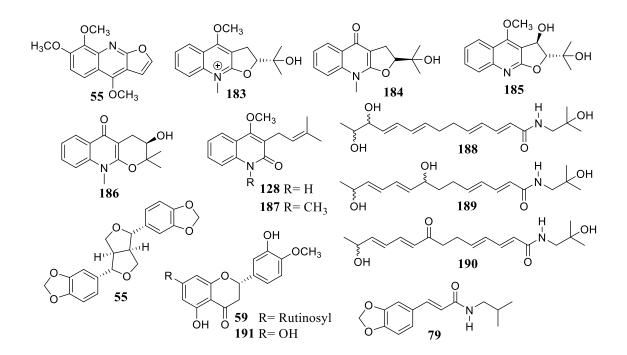


Figure 3.77 Structures of isolated compounds from the fruits of Z. zanthoxyloides

#### 3.3.5.1 Structure elucidation of skimmianine (55)

Compound 55 was isolated as white needles. Its molecular formula  $C_{14}H_{13}NO_4$  was determined from its HR ESI-MS spectrum (Figure 3.78) obtained in positive ion mode,

where the *pseudo*molecular ion peak was observed at *m/z* 260.0919 [M+H]<sup>+</sup> calculated for C<sub>14</sub>H<sub>13</sub>NO<sub>4</sub>H, 260.0917. Its <sup>1</sup>H NMR spectrum (Figure 3.79, Table 3.18) showed characteristic signals of three methoxy groups at  $\delta_{\rm H}$  3.89 (OCH<sub>3</sub>-8), 4.02 (OCH<sub>3</sub>-7) and 4.49 (OCH<sub>3</sub>-4) and four aromatic methines at  $\delta_{\rm H}$  7.28 (d, *J* = 2.6 Hz, H-1<sup>'</sup>), 7.36 (d, *J* = 9.4 Hz, H-6), 7.75 (d, *J* = 2.6 Hz, H-2<sup>'</sup>) and 8.05 (d, *J* = 9.4 Hz, H-5). Its <sup>13</sup>C NMR (Figure 3.80, Table 3.18) revealed 14 signals attributable to three methoxy, four methines and seven quaternary carbons including the deshielded signal at  $\delta_{\rm C}$  164.6 suggesting the presence of an amide in the molecule. Analysis of the different correlations present in the HMBC spectrum (Figure 3.80) revealed the core structure of the molecule to be quinoline alkaloid. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.18) of **55** were in good agreement with those published for skimmianine (Chakravarty *et al.*, 1999). Thus, compound **55** was identified as skimmianine, a common quinoline alkaloids in the Rutaceae family (Adamska-Szewczyk *et al.*, 2016).

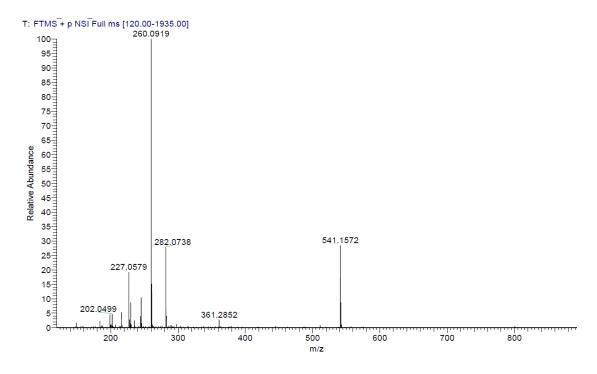


Figure 3.78 ESI-MS spectrum of 55

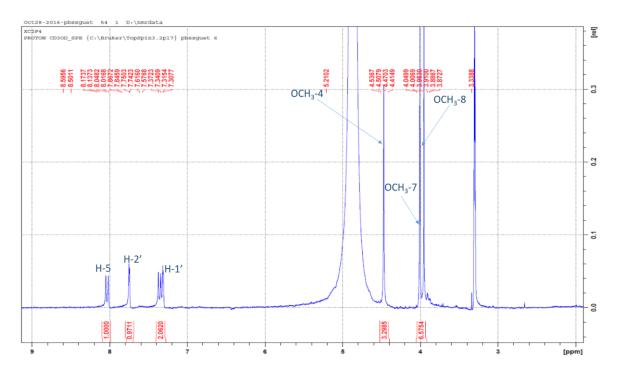


Figure 3.79  $^{1}$ H NMR (300 MHz, CD<sub>3</sub>OD) of **55** 

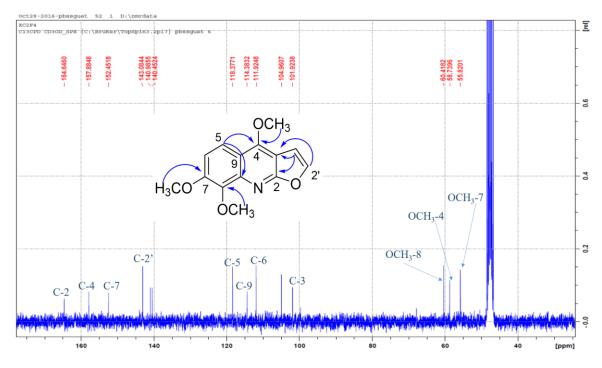


Figure 3.80 <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **55** 

Compound 183 was isolated as a dark yellow amorphous powder. Its molecular formula  $C_{16}H_{20}NO_3$  was determined from its HR ESI-MS spectrum by the peak at m/z 274.1440 [M]<sup>+</sup> calculated for C<sub>16</sub>H<sub>20</sub>NO<sub>3</sub>, 274.1438. Its <sup>1</sup>H NMR spectrum (Figure 3.81, Table 3.18) showed characteristic signals of an *ortho*-disubstituted benzene ring at  $\delta_{\rm H}$  8.32 (d, J = 9.0 Hz), 8.03 (ov), 7.96 (ov) and 7.71 (td, J = 6.0, 9.0 Hz). Four methyl singlets including two deshielded methyls at  $\delta_{\rm H}$  4.11 and 4.53; a methylene at  $\delta_{\rm H}$  3.97 (dd, J = 6.0, 15.0 Hz) and an oxymethine at  $\delta_{\rm H}$  (dd, J = 6.0, 15.0 Hz) were also observed. Analysis of its <sup>13</sup>C NMR (Figure 3.82, Table 3.18) led to the identification of 16 signals which could be attributed to four methyls including an N-methyl at  $\delta_C$  33.0 ( $\delta_H$  4.11); a methylene at  $\delta_{\rm C}$  28.6; five methines including an oxymethine at  $\delta_{\rm C}$  94.3 and six quaternary carbons including the deshielded signal at  $\delta_{\rm C}$  166.1 suggesting the presence of an amide in the molecule. In the HMBC spectrum, correlations observed between the methyl signals at  $\delta_{\rm H}$  1.25 (3H-4') and 1.42 (H-5') with the carbons at  $\delta_{\rm C}$  23.4 or 24.6 (C-4' or C-5'), 70.5 (C-3') and 94.3 (C-2') suggested they were in geminal position on the carbon C-3' and the entire group fixed on C-2'. Correlations observed between the methyl at  $\delta_{\rm H}$  4.11 (N-CH<sub>3</sub>) and the carbons at  $\delta_{\rm C}$  136.7 (C-10) and 166.1 (C-2) confirmed its position on the nitrogen atom. The proton H-2'was determined to be  $\beta$ -oriented due to the observed coupling constants (J = 6.0, 15.0 Hz) with the adjacent methylene 2H-1'. All the <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.18) of **183** were in good agreement with those published for *N*methylplatydesminium cation (Boyd and Grundon, 1967; Boyd et al., 2007). Thus, compound 183 was identified as N-methylplatydesminium cation, a quaternary quinoline alkaloid first isolated from Skimmia Japonica (Rutaceae) (Boyd and Grundon, 1967).

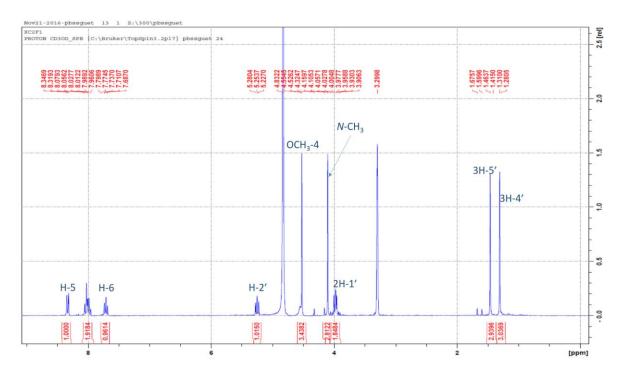


Figure 3.81  $^{1}$ H NMR (300 MHz, CD<sub>3</sub>OD) of **183** 

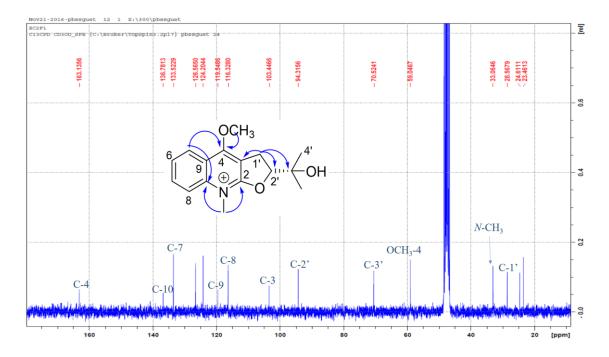


Figure 3.82 <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **183** 

### 3.3.5.3 Structure elucidation of isoplatydesmine (184)

Compound **184** was isolated as yellow amorphous powder. Its molecular formula  $C_{15}H_{17}NO_3$  was determined from its HR ESI-MS spectrum obtained in positive ion mode by the peak at m/z 260.1284 [M+H]<sup>+</sup> calculated for  $C_{15}H_{17}NO_3H$ , 260.1281. Its 1D and 2D NMR spectra were similar to those of **183**. The only difference was the absence in the <sup>1</sup>H NMR spectrum (Figure 3.83, Table 3.18) of **184** of the methoxy proton signal present in that of **183** suggesting a demethylation on position C-4. This was further confirmed by the different <sup>13</sup>C chemical shifts in the <sup>13</sup>C NMR (Figure 3.84) especially the carbon C-4 which appeared at  $\delta_C$  172.6 as well as the key correlations observed in its HMBC spectra (Figure 3.84). Therefore, compound **184** was identified as isoplatydesmine and all its <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.18) were in good agreement with the published data (Boyd and Grundon, 1967; Boyd *et al.*, 2007).

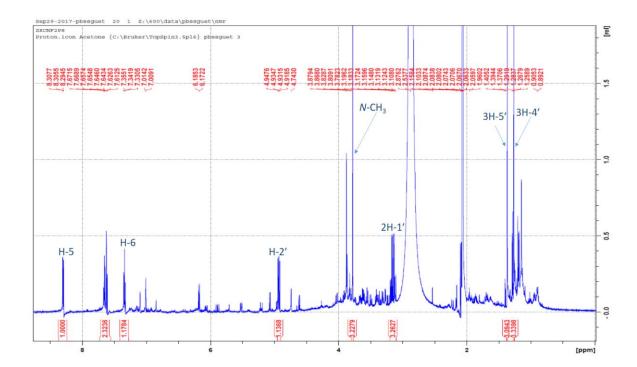


Figure 3.83 <sup>1</sup>H NMR (600 MHz, acetone- $d_6$ ) of **184** 

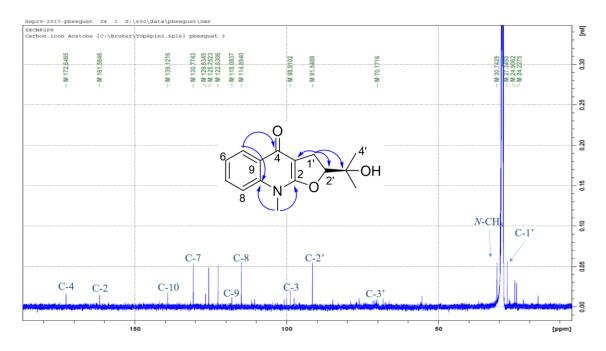


Figure 3.84<sup>13</sup>C NMR (150 MHz, acetone-d<sub>6</sub>) and key HMBC correlations of **184** 

## 3.3.5.4 Structure elucidation of ribalinine (186)

Compound **186** was isolated as yellow amorphous powder. Its molecular formula  $C_{15}H_{17}NO_3$  was determined from its HR ESI-MS spectrum obtained in positive ion mode by the peak at m/z 260.1284 [M+H]<sup>+</sup> calculated for  $C_{15}H_{17}NO_3H$ , 260.1281. Its 1D and 2D NMR spectra were very similar to those of **184**. The only difference was the chemical shifts observed for the C-3 side chain. In the <sup>1</sup>H NMR spectrum (Figure 3.85, Table 3.18) of **186**, the H-2' proton ( $\delta_H$  3.91, dd, J = 5.0, 5.6 Hz) was found to be shielded compared to the corresponding proton in **184**. In the <sup>13</sup>C NMR spectrum (Figure 3.86, Table 3.18), the signal of carbon C-3' ( $\delta_C$  84.3) bearing the two *gem*-methyls was deshielded while that of C-2' ( $\delta_C$  69.0) was shielded compared to the values observed for the same carbons in **184**. This observation suggested a six membered side ring. The 2'-OH was determined to be  $\beta$ -oriented due to the small coupling constants (J = 5.0, 5.6 Hz) for its coupling with the 2H-1' protons. Therefore, compound **186** was identified as ribalinine (Corral and

Orazi, 1967). Its <sup>1</sup>H and <sup>13</sup>C NMR data (Tables 3.21-3.22) were in good agreement with the published data (Corral and Orazi, 1967; Boyd *et al.*, 2007).

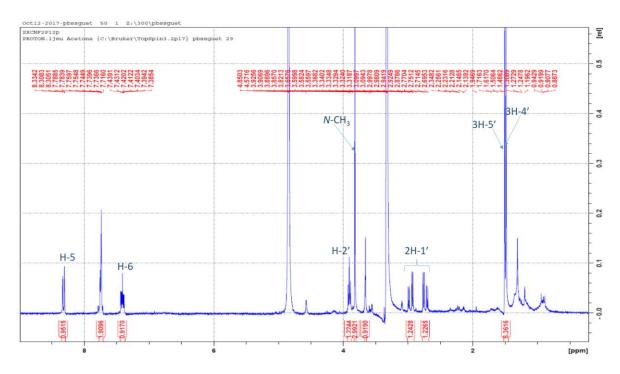


Figure 3.85  $^{1}$ H NMR (300 MHz, CD<sub>3</sub>OD) of **186** 

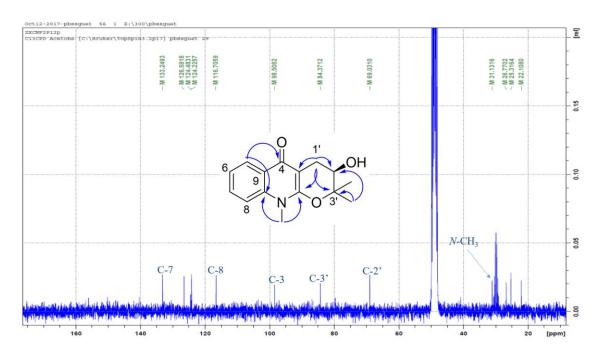


Figure 3.86  $^{13}$ C NMR (75 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **186** 

Compound **128** was isolated as yellow amorphous powder. Its molecular formula  $C_{15}H_{17}NO_2$  was determined from its HR ESI-MS spectrum obtained in positive ion mode by the peak at m/z 244.1337 [M+H]<sup>+</sup> calculated for  $C_{15}H_{17}NO_2H$ , 244.1338. Its 1D and 2D NMR spectra similar to those of **183** also suggested a quinoline skeleton. The chemical shifts observed for the C-3 side chain in **128** suggested an open cycle compared to the previous isolated quinolones. In addition, the absence of the *N*-methyl signal in the <sup>1</sup>H NMR spectrum (Figure 3.87, Table 3.18) of **128** was clearly visible. The methine signal appearing at  $\delta_H 5.25$  showing cross peak correlation in the HSQC-DEPT spectrum with the carbon at  $\delta_C 122.7$  was attributed to H-2'. In the HMBC spectrum (Figure 3.87), correlations observed between the two *gem*-methyls ( $\delta_C 25.8$ , C-4'; 18.0, C-5';  $\delta_H 1.71$ , H-4'; 1.82, H-5') with the carbons at  $\delta_C 18.0$  or 25.8 (C-5' or C-4'), 122.7 (C-2') and 133.3 (C-3') suggested the C-3 side chain to be a dimethylallyl moiety. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.18) of **128** was identified as atanine, a quinolone alkaloid first isolated from *Z. zanthoxyloides* by Eshiett and Taylor (1968).

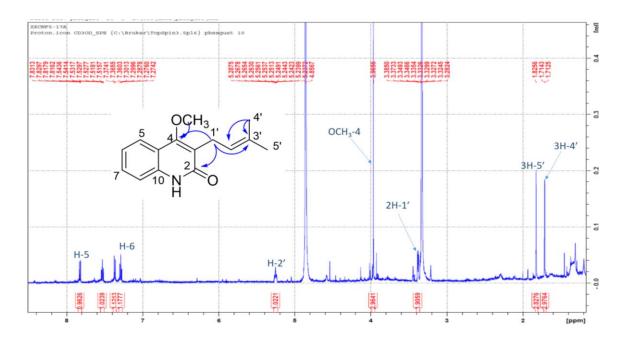


Figure 3.87 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) and key HMBC correlations of **128** 

184

### 3.3.5.6 Structure elucidation of *N*-methylatanine (187)

Compound **187** was isolated as yellow amorphous powder. Its molecular formula  $C_{16}H_{19}NO_2$  was determined from its HR ESI-MS spectrum obtained in positive ion mode by the peak at m/z 258.1494 [M+H]<sup>+</sup> calculated for  $C_{16}H_{19}NO_2H$ , 258.1494. Its 1D and 2D NMR spectra were very similar to those of **128**. The only difference was the presence in the <sup>1</sup>H NMR spectrum (Figure 3.88, Table 3.18) of **187** of a singlet methyl at  $\delta_H$  3.74 attributable to *N*-methyl. This attribution was confirmed in the HMBC spectrum (Figure 3.88) by the correlation between the *N*-methyl protons and the carbons at  $\delta_C$  139.0 (C-10) and 164.5 (C-2). Thus, **187** was identified as *N*-methylatanine and all its <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.18) were in good agreement with the published data (Brown, 1980).

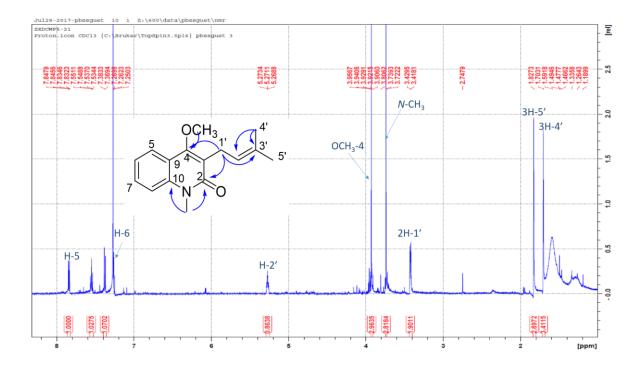


Figure 3.88 <sup>1</sup>H NMR (600 MHz, CDCl<sub>3</sub>) and key HMBC correlations of **187** 

Position						δι	C						
	55		128	183	184	186	187	55	128	183	184	186	187
1								-	-	-	-	-	-
2								164.6	164.3	166.1*	161.2	156.8	164.5
3								101.9	122.7	103.4	98.0	98.4	122.6
4								157.8	162.1	163.1	172.6	nd	159.8
5	8.05 (9. <i>4</i> )	d	7.82 dd (1.0, 8.1)	8.32 d (9.0)	8.31 dd ( <i>1.4</i> , 7.9)	8.32 dd ( <i>1.4</i> , 7.9)	7.84 dd (1.4, 7.9)	118.3	124.0	124.2	125.8	124.1	123.4
6	7.36 (9.4)	d	7.29 ddd (1.2, 7.1, 8.1)	7.71 td (6.0, 9.0)	7.34 ddd (1.0, 7.0, 7.9)	,	7.27 m	111.9	123.5	126.5	122.0	126.5	121.8
7			7.53 ddd (1.3, 7.1, 8.3)	7.96 ov	7.65 ddd (1.5, 6.4, 8.4)	7.75 ov	7.56 td (1.4, 7.2, 8.4)	152.4	131.4	133.5	130.7	133.2	130.0
8			7.36 d (8.2)	8.03 ov	7.63 brd (8.4)	7.75 ov	7.39 d (8.4)	104.9	116.6	116.3	114.8	116.6	114.1
9								114.6	117.0	119.5	118.0	124.4	117.8
10								104.4	142.1	136.7	139.1	140.7	139.0
1'	7.28 (2.6)	d	3.37 ov	3.97 dd (6.0, 15.0)	3.13 dd (7.4, 16.6)	2.74 dd (5.6, 16.5)	3.43 d (6.8)	104.9	24.2	28.6	27.3	26.7	24.3
					3.17 dd (9.6, 16.6)	2.96 dd (5.0, 16.5)							
2'	7.75 (2.6)	d	5.25 m	5.25 dd (6.0, 15.0)	4.98 dd (9.6, 7.4)	3.91 dd (5.0, 5.6)	5.28 m	143.9	122.7	94.3	91.5	69.0	121.5

Table 3.18 <sup>1</sup>H and <sup>13</sup>C NMR data of **55**, **128**, **183-184** and **186-187** 

Position		δ		δ <sub>C</sub>							
55	128	183	184	186	187	55	128	183	184	186	187
3'	-	-	-	-	-	-	133.3	70.5	70.1	84.3	132.5
4'	1.71	1.25 s	1.29 s	1.48 s	1.71 s	-	25.8	23.4	24.2	25.3	25.7
5'	1.82 s	1.42 s	1.38 s	1.50 s	1.83 s	-	18.0	24.6	24.9	22.1	18.0
N-CH <sub>3</sub>	-	4.11 s	3.78 s	3.82 s	3.74 s	-	-	33.0	30.7	31.1	29.7
4-OCH <sub>3</sub> 4.49 s	3.96 s	4.53 s	-			58.7	62.4	59.0	-	-	61.7
7-OCH <sub>3</sub> 4.02 s	-	-	-			55.8					
8-OCH <sub>3</sub> 3.89 s	-	-	-			60.4					

Table 3.18 continued

\*Values determined from the HMBC spectrum

Compound 188 was obtained as white viscous liquid. Its molecular formula was determined as  $C_{18}H_{29}NO_4$  from its HR ESI-MS spectrum (Figure 3.89) by the pseudomolecular ion at m/z 346.1984 calculated 346,1989 for its sodium adduct [M + Na]<sup>+</sup> (C<sub>18</sub>H<sub>29</sub>NO<sub>4</sub>Na). The <sup>1</sup>H NMR spectrum of **188** (Figure 3.90, Table 3.19) exhibited 14 signals corresponding to 24 protons including the signals of the proton of a secondary amine at  $\delta_{\rm H}$  8.56 (1H, brs), six olefinic protons at  $\delta_{\rm H}$  7.12 (1H, dd, J = 10.4, 15.1 Hz),  $6.60 (1H, dd, J = 11.1, 15.1 Hz), 6.27 (1H, dd, J = 10.8, 15.1 Hz), \delta 5.99 (1H, d, J = 15.0 Hz)$ Hz), 5.70 (1H, dd, J = 6.4, 15.0 Hz) and 5.44 (1H, m, J = 7.4 Hz), signals at  $\delta_{\rm H}$  1.18 (6H, brs) and 1.11 (3H, d, J = 6.3 Hz) for three methyls, two oxygenated methines at  $\delta_{\rm H}$  3.58 (1H, m, J = 6.3 Hz) and 3.91 (1H, m, J = 6.4 Hz) and three methylene signals including a broad singlet at  $\delta_{\rm H}$  3.27 and two double doublets at  $\delta_{\rm H}$  2.29 (J = 7.0, 13.2 Hz) and 2.35 (J = 7.3, 14.8 Hz). Overlapping signals occurring between  $\delta_{\text{H}} 6.0-6.15$  ppm resulting from two olefinic protons resonances were also observed. The <sup>13</sup>C NMR spectrum (Figure 3.91, Table 3.19) revealed signals for 18 carbons attributable to three methyls ( $\delta_C$  27.2 x 2,  $\delta_C$ 18.8), three methylenes with one appearing relatively downfield at  $\delta_{\rm C}$  51.1, ten methines including two oxymethines at  $\delta_C$  77.8 and 71.7, eight olefinic methines at  $\delta_C$  143.0, 142.3, 134.0, 131.7, 130.2, 129.9, 128.2 and 123.2, and two quaternary carbons including an amide at  $\delta_{\rm C}$  169.4 and an oxygenated quaternary carbon at  $\delta_{\rm C}$  71.6. The assignment of the carbon signals was consistent with the resonances observed in the <sup>1</sup>H NMR experiment as well as the <sup>1</sup>J <sup>1</sup>H-<sup>13</sup>C correlations observed in the HSQC-DEPTQ spectrum (Figure 3.92). In the <sup>1</sup>H-<sup>1</sup>H COSY spectrum (Figure 3.93), a chain of vicinal correlations observed from H-2 to H-14, helped to construct one part of the compound structure as CH<sub>3</sub>-CH=CH-CH=CH-CH<sub>2</sub>-CH<sub>2</sub>-CH=CH-CH=CH-. In the HMBC spectrum (Figure 3.94), correlations from  $\delta_C$  1.18 (H-3' and H-4') to carbon signals at  $\delta_C$  51.1 (C-1'), 71.6 (C-2') and 27.2 (C3' or C-4'), and from  $\delta_C$  3.27 (s, H-1') to C-2', C-3' and C-4' identified the 188 other part of the molecule as 2-hydroxy-isobutyl moiety. Further correlations in the HMBC spectrum from H-1' ( $\delta_{H}$  3.27), H-2 ( $\delta_{H}$  5.99) and H-3 ( $\delta_{H}$  7.12) to C-1 ( $\delta_{C}$  169.4), in addition to the appearance of the methylene of the isobutyl moiety at  $\delta_{C}$  55.1, established that the 2-hydroxy-isobutyl moiety was linked to the nitrogen of the amide group and the aliphatic moiety to the carbonyl of the amide. The geometry of the double bond C<sub>2</sub>-C<sub>3</sub> was deduced as *trans* ( $J_{H2/H3} = 15.0 \text{ Hz}$ ) like those of C<sub>4</sub>-C<sub>5</sub>, C<sub>8</sub>-C<sub>9</sub> and C<sub>10</sub>-C<sub>11</sub>. Compound **188** did not show any optical activity suggesting that it was obtained as a racemic mixture. Thus compound **188** was identified as (12*RS*, 13*RS*)-(2*E*, 4*E*, 8*E*, 10*E*)-*N*-(2-hydroxy-2-methylpropyl)-12,13-dihydroxy-2,4,8,10-tetradecatetraenamide, a new alkamide from natural source and given the trivial name zanthoamide G. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.19) were similar to those published for zanthoamide C isolated from *Zanthoxylum bungeanum* (Wang *et al.*, 2016) with the only difference being the additional olefinic bond between C3 and C6 present in **188**.

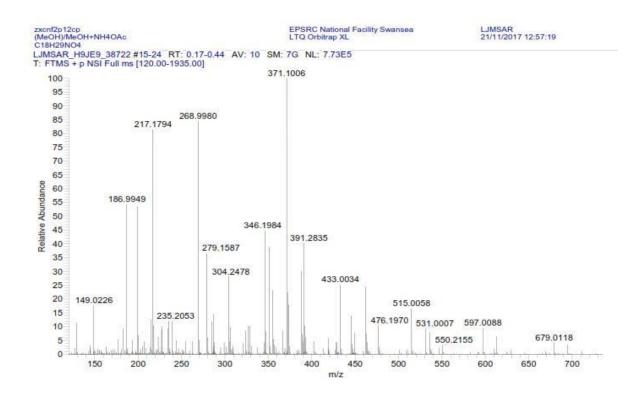


Figure 3.89 ESI-MS spectrum of 188

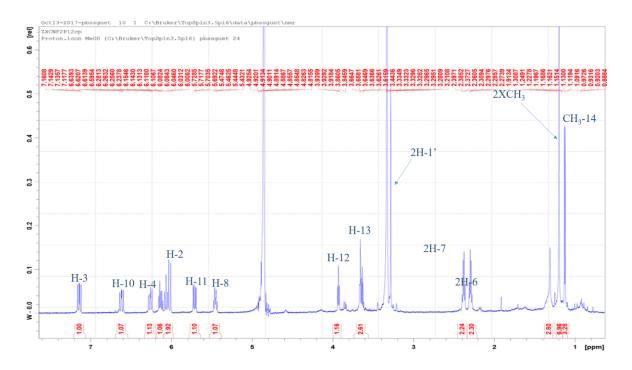


Figure 3.90  $^{1}$ H NMR (600 MHz, CD<sub>3</sub>OD) of **188** 

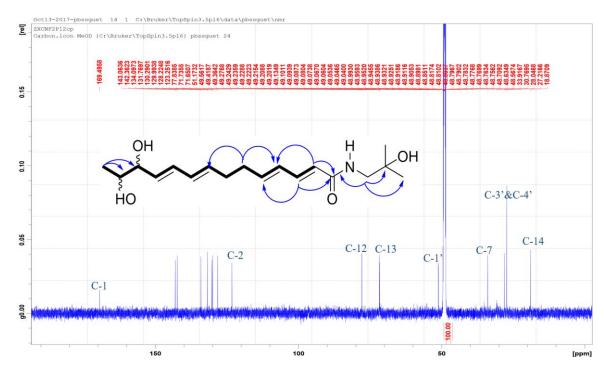


Figure 3.91 <sup>13</sup>C NMR (150 MHz, CD<sub>3</sub>OD), COSY (–) and Key HMBC ( $\rightarrow$ ) correlation of **188** 

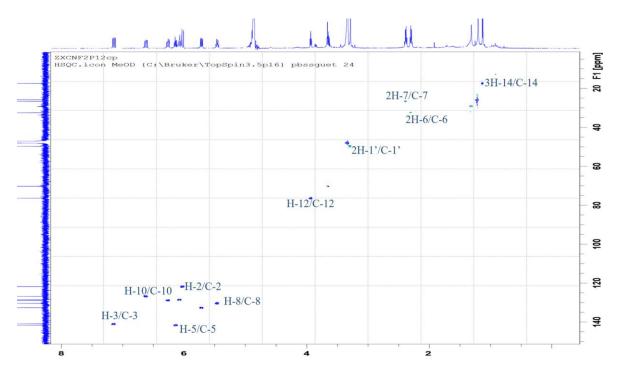


Figure 3.92 HSQC spectrum of 188

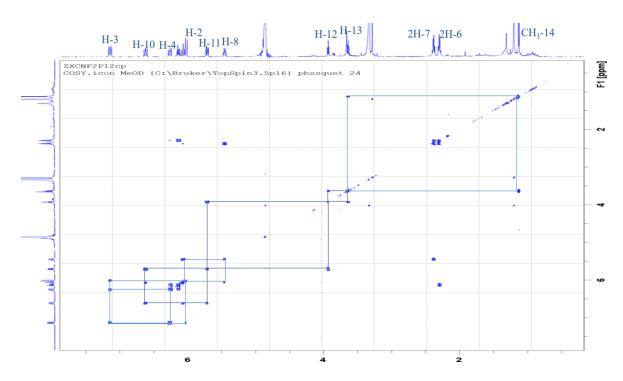


Figure 3.93 <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **188** 

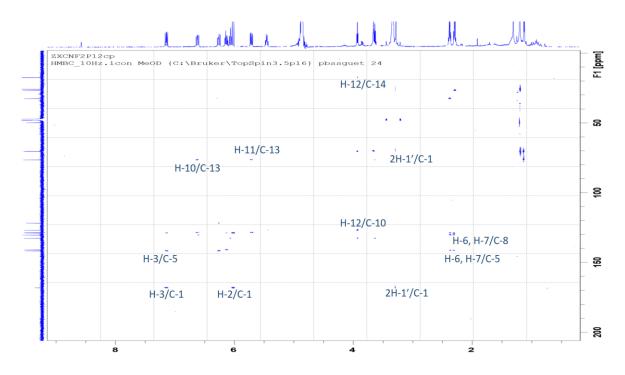


Figure 3.94 HMBC spectrum of 188

## 3.3.5.8 Structure elucidation of zanthoamides H (189)

Compound **189** was also obtained as white viscous liquid. Its molecular formula was determined as  $C_{18}H_{29}NO_4$  from its HR ESI-MS spectrum, where a *pseudo*molecular ion was observed as at *m*/z 346.1992 [M + Na]<sup>+</sup> (calculated 346.1989 for  $C_{18}H_{29}NO_4Na$ ). The <sup>1</sup>H and <sup>13</sup>C NMR spectra (Figures 3.95-3.96, Table 3.19) of **189** established it as an aliphatic amide, similar to **188**, with the only difference being the placement of the two oxymethines, which were not adjacent to each other like in **188**. The COSY spectrum (Figure 3.96) revealed scaler couplings between oxymethine at  $\delta_H$  4.28 (H-13) and the methyl at  $\delta_H$  1.20 (H-14) and with the olefinic proton at  $\delta_H$  5.73 (H-12), and the correlations from the other oxymethine at  $\delta_H$  4.10 (H-8) to the methylene at  $\delta_H$  1.60 (H-7) and the olefinic proton at  $\delta_H$  5.68 (H-9). These correlations from H-8 to C-6 ( $\delta_C$  29.9), C-7 ( $\delta_C$  37.4) and C-9 ( $\delta_C$  137.0), and from H-13 to C-12 ( $\delta_C$  138.6) and C-14 ( $\delta_C$  23.5)

were observed. The geometry of the olefinic bonds were confirmed as *trans* from relevant coupling constants ( $J_{H2/H3} = 15.1$ ,  $J_{H4/H5} = 15.2$ ,  $J_{H9/H10} = 14.6$  and  $J_{H11/H12} = 14.5$ ). Compound **189** was optically inactive suggesting that it was a racemic mixture. Therefore, compound **189** was identified as (8*RS*, 13*RS*)-(2*E*, 4*E*, 9*E*, 11*E*)-*N*-(2-hydroxy-2-methylpropyl)-8,13-dihydroxy-2,4,9,11-tetradecatetraenamide, and given the trivial name, zanthoamide H.

## 3.3.5.9 Structure elucidation of zanthoamides I (190)

Compound 190 had a molecular formula  $C_{18}H_{27}NO_4$  determined at m/z 322.2017 by HR ESI-MS  $[M+H]^+$  (calculated for  $[C_{18}H_{27}NO_4+H]^+$ , 322.2013) corresponding to six double bonds and rings equivalent. The comparison of <sup>1</sup>H NMR spectrum (Figure 3.97, Table 3.19) of **190** to those of **189** clearly disclosed that **190** was also an aliphatic amide. The obvious difference between the two compounds was that the hydroxyl group at C-8 on 189 was replaced by a ketone group in 190, which was confirmed by the signal at  $\delta_{\rm C}$  202.0 in the <sup>13</sup>C NMR spectrum (Figure 3.98, Table 3.19) of **190**. The position of the ketone group was confirmed using HMBC spectrum (Figure 3.98) by the correlation observed between this later and the protons at  $\delta_{\rm H}$  2.48 (H-6), 6.21 (H-9), 2.79 (H-7) and 7.29 (H-10). This was also supported by the downfield shifts of C-10 and C-12 from  $\delta_C$  129.8 and 138.6 in **189** to  $\delta_{\rm C}$  144.5 and 149.3 respectively in **190** confirming the presence of a ketone group nearby. The geometry of the double bonds  $C_2$ - $C_3$ ,  $C_4$ - $C_5$ ,  $C_9$ - $C_{10}$  and  $C_{11}$ - $C_{12}$  was determined as trans-configured (J = 15.0, 15.0, 15.6 and 15.1 Hz, respectively). Compound **190** was found to be optically active,  $[\alpha]_D^{25} = -25.6$ . However, the absolute configuration could not be determined because of paucity of this sample. Thus, compound 190 was identified as  $(13^*)$ -(2E, 4E, 9E, 11E)-N-(2-hydroxy-2-methylpropyl)-13hydroxy-2,4,9,11-tetradecatetraenamide and given the trivial name, zanthoamide I.

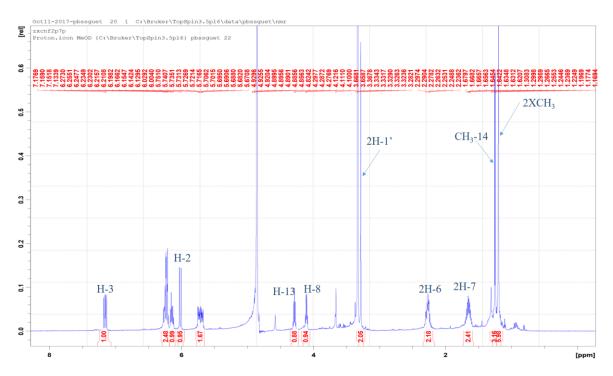


Figure 3.95  $^{1}$ H NMR (600 MHz, CD<sub>3</sub>OD) of **189** 

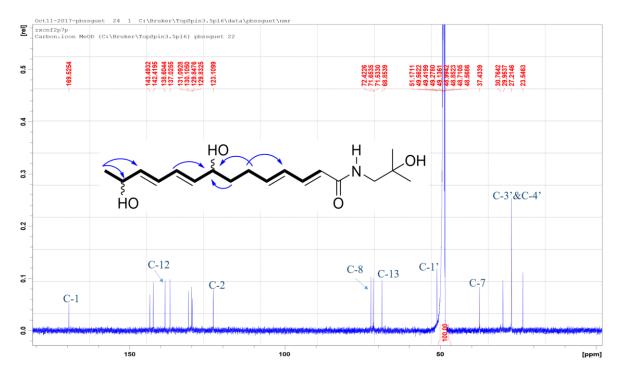


Figure 3.96 <sup>13</sup>C NMR (150 MHz, CD<sub>3</sub>OD), COSY (–) and Key HMBC ( $\rightarrow$ ) correlation of **189** 

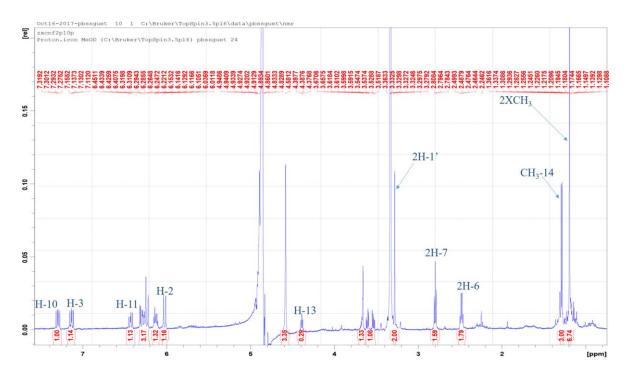


Figure 3.97 <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) of **190** 

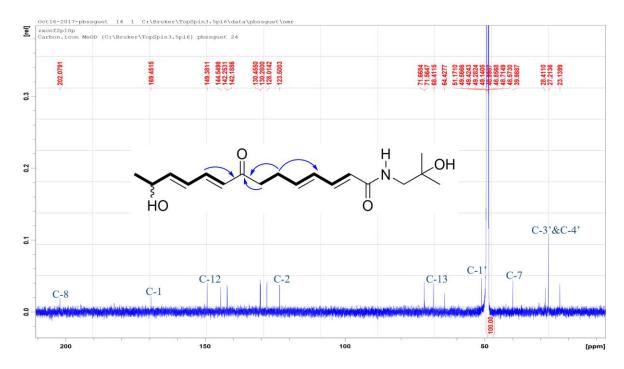


Figure 3.98 <sup>13</sup>C NMR (150 MHz, CD<sub>3</sub>OD), COSY (–) and Key HMBC ( $\rightarrow$ ) correlation of **190** 

1		δ <sub>C</sub>		
	188	189	190	188 189 190
1	-	-	-	169.4 169.5 169.4
2	5.99 d ( <i>15.0</i> )	6.01 d ( <i>15.1</i> )	6.01 d ( <i>15.0</i> )	123.2 123.1 123.5
3	7.12 dd ( <i>10.4</i> ,	7.15 dd ( <i>10.7</i> ,	7.13 dd ( <i>10.7</i> ,	142.3 142.4 142.1
	15.1)	15.0)	15.1)	
4	6.27 dd (10.8,	6.25 dd (10.7,	6.26 m	130.2 130.1 130.4
	15.1)	15.2)		
5	6.11 m	6.14 m	6.13 dt (7.0,	143.0 143.4 142.2
			15.0)	
6	2.29 dd (7.0,	2.20 m	2.48 dd (7.2,	33.9 29.9 28.4
	13.2)		14.1)	
7	2.35 dd (7.3,	1.60 m	2.79 dd (7.2,	28.0 37.4 39.9
	14.8)		14.5)	
8	5.44 dd (7.2,	4.10 q ( <i>6.5</i> , <i>13.0</i> )	-	131.7 72.4 202.0
	15.1)			
9	6.07 m	5.68 dd (6.7,	6.21 d ( <i>15.6</i> )	129.9 137.0 130.2
		14.6)		
10	6.60 dd (11.1,	6.20 m	7.29 dd (10.8,	128.2 129.8 144.5
	15.1)		15.6)	
11	5.70 dd (6.4,	6.20 m	6.42 dd (10.3,	134.0 131.0 128.0
	15.0)		15.1)	
12	3.91 dd (6.4,	5.73 m	6.31 m	77.8 138.6 149.3
	12.9)			
13	3.58 m	4.28 q ( <i>6.3</i> , <i>12.5</i> )	4.38 m	71.7 68.8 68.4
14	1.11 d (6.3)	1.20 d (6.4)	1.22 d (6.5)	18.8 23.5 23.1
1'	3.27 s	3.28 s	3.27 s	51.1 51.1 51.1
2'	-	-	-	71.6 71.6 71.5
3'	1.18 s	1.19 s	1.18 s	27.2 27.2 27.2
4'	1.18 s	1.19 s	1.18 s	27.2 27.2 27.2
NH	8.56 br s	8.56 br s	8.56 br s	

Table 3.19  $^{1}$ H (600 MHz) and  $^{13}$ C (150 MHz) NMR Data of zanthoamide G-I (188-190)

Compound 191 was isolated as yellow amorphous powder. Its molecular formula  $C_{20}H_{30}O_2$  was determined from its HR ESI-MS spectrum (Figure 3.99) obtained in negative ion mode by the peak at m/z 301.2170 [M-H]<sup>-</sup> calculated for C<sub>20</sub>H<sub>29</sub>O<sub>2</sub>, 301.2173. Its <sup>1</sup>H NMR spectrum (Figure 3.100, Table 3.20) revealed the presence of a methoxy at  $\delta_{\rm H}$  3.88 (s); a 1,3,4-trisubstituted benzene ring formed by the protons at  $\delta_{\rm H}$  7.06 (d, J = 1.7 Hz), 7.00 and 6.99; two meta coupling protons at  $\delta_{\rm H}$  5.97 (dd, J = 3.0, 16.8 Hz) and an oxymethine at  $\delta_{\rm H}$  5.45 (dd, J = 3.0, 12.6 Hz) which were found to couple in the COSY spectrum with the methylene protons appearing at  $\delta_{\rm H}$  2.76 (dd, J = 3.0, 16.8 Hz) and 3.16 (dd, J = 12.6, 16.8 Hz). The <sup>13</sup>C NMR spectrum (Figure 3.101, Table 3.20) of **191** depicted sixteen peaks which could be assigned based on the correlation observed in the HSQC spectrum to one methoxy at  $\delta_C$  55.2; one methylene at  $\delta_C$  42.9, six methines including an oxymethine at  $\delta_{\rm C}$  79.0 and eight quaternary carbons including the highly deshielded carbon at  $\delta_{\rm C}$  196.2 suggesting a conjugated carbonyl. The <sup>1</sup>H and <sup>13</sup>C NMR data (Table 3.20) of 191 and its key HMBC correlations (Figure 301) were in good agreement with those described for hesperetin (EI-Shafae et al., 2002). The C-2 proton at  $\delta_{\rm H}$  5.97 (d, J = 3.0, 16.8 Hz) was determined to be  $\beta$ -oriented due to the coupling constant observed from its coupling with the C-3 methylene. Thus, compound 191 was identified as hesperetin, a flavone commonly isolated from Citrus species (Rutaceae) (EI-Shafae et al., 2002; Gattuso et al., 2007).

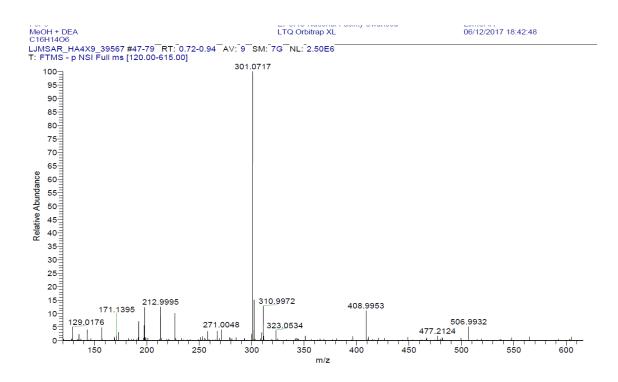


Figure 3.99 ESI-MS spectrum of 191

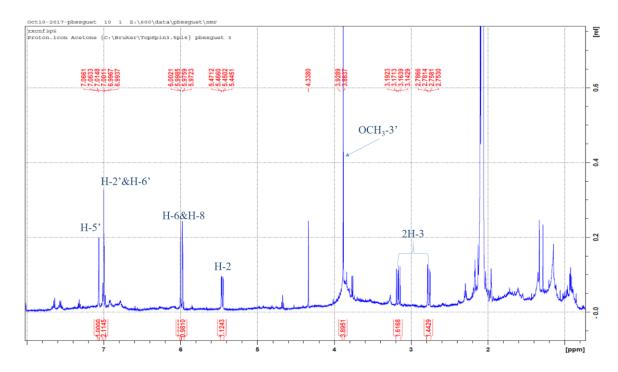


Figure 3.100  $^{1}$ H NMR (600 MHz, acetone-d<sub>6</sub>) of **191** 

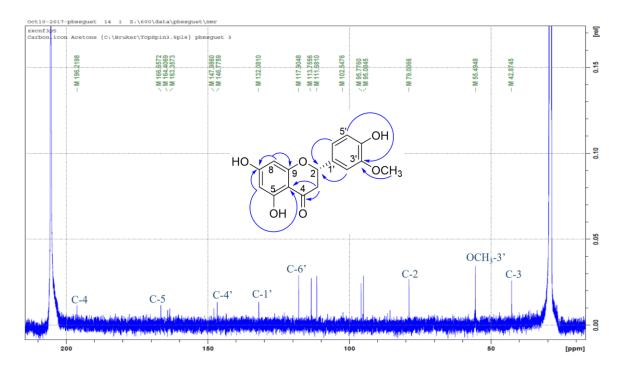


Figure 3.101  $^{13}$ C NMR (150 MHz, acetone-d<sub>6</sub>) of compound **191** 

Table 3.20 $^{1}$ H (600 MHz, acetone-d <sub>6</sub> ) at	d <sup>13</sup> C NMR data (150 MHz, acetone-d <sub>6</sub> ) of <b>191</b>

Position	$\delta_{\rm H} { m m}  (J { m in} { m Hz})$	$\delta_c$	Position	$\delta_{\rm H} { m m}  (J { m in} { m Hz})$	$\delta_c$
1	-	-	9	-	163.3
2	5.45 dd (3.0, 12.6)	79.0	10	-	102.5
3	2.76 dd (3.0, 16.8)	42.8	1'	-	132.0
	3.16 dd ( <i>12.6</i> , <i>16.8</i> )		2'	7.00 ov	111.6
4	-	196.2	3'	-	147.9
5	-	166.6	4'	-	146.7
6	5.97 d (2.1)	95.7	5'	7.06 d ( <i>1.7</i> )	113.7
7	-	164.4	6'	6.99 ov	117.9
8	6.00 d (2.1)	95.0	OCH <sub>3</sub>	3.88 s	55.4

#### 3.4 Biogenesis and chemotaxonomic significance of isolated compounds

Three main classes of secondary metabolites including alkaloids, diterpenes and flavonoids were isolated in the course of this study.

## 3.4.1 Alkaloids

Quinoline alkaloids (**55**, **128**, **183-187**) and alkamides (**188-190**) were isolated from *Z*. *zanthoxyloides* (Rutaceae). *Zanthoxyllum* is a rich source of alkaloids including acridone, aliphatic and aromatic amides, aporphine and benzophenantridine (Adesina, 2005; Patino *et al.*, 2012).

This is the first report on the occurrence of compounds **183-187**, in addition to three new alkamides (**189-191**), in *Z. zanthoxyloides*. Several alkamides with structures similar to zanthoamides G-I (**189-191**) were previously reported in several *Zanthoxylum* species including *Z. zanthoxyloides*, *Z. bungeanum*, *Z. syncarpum*, *Z. piperitum*, *Z. ailanthoides*, *Z. integrifoliolum*, *Z. schinifolium*, *Z. armatum*, *Z. tessmannii*, *Z. achtoum*, and *Z. heitzii* (Ross *et al.*, 2005; Wang *et al.*, 2016; Chruma *et al.*, 2018). *N*-methylplatydesminium catium (**183**) was previously isolated from *Z. usambarense* and *Z. chalybeum* (Kato *et al.*, 1996), isoplatydesmine (**184**) from *Z. nididum* (Ishikawa *et al.*, 1995), myrtopsine (**185**) from *Z. integrifoliolum* (Ishii *et al.*, 1982), ribalinine (**186**) from *Z. mayu* (Torres and Cassels, 1978), and *N*-methylatanine (**187**) from *Z. beecheyanum* (Cheng *et al.*, 2004) and *Z. rigidum* (Moccelini *et al.*, 2009).

Quinoline alkaloids are biosynthesized from anthranilic acid, which is formed from chorismate and L-glutamine, while alkamides derived from the condensation of amino acid and/or cinnamic acid unit (Waterman, 1975). For quinoline synthesis, anthranilic acid is acetylated followed by cyclisation to form the basic 2,4-dihydroxyquinoline ring,

which may then undergo a nucleophilic addition of an isoprenyl unit on the highly reactive C-3 carbon to give rise to 3-(3',3'-dimethylallyl)-4-hydroxy-2-quinolone (Waterman, 1975; Grundon, 1988). Subsequent modifications on the obtained quinolone including alkylation, cyclisation (e.g. C-2 and C-2' or C-3') and oxidation may lead to the formation of the known quinoline alkaloids such as those isolated from *Z. zanthoxyloides* in the course of this study (Figure 3.102) (Fish and Waterman, 1973; Waterman, 1978; Boyd *et al.*, 2000).

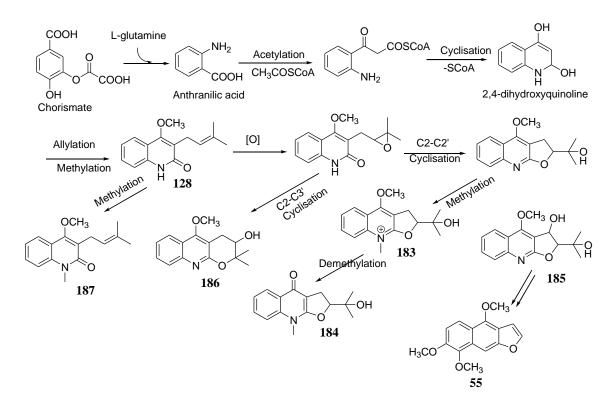


Figure 3.102 Possible biosynthesis pattern of the quinolines isolated from Z. *zanthoxyloides* 

### **3.4.2 Diterpenes**

Abietane and clerodane were isolated from *C. oligandrus*, and kaurane from *Z. leprieurii*, respectively.

The genus *Croton* is a great natural source of diterpenes. Its phytochemical studies have led to the isolation of several diterpenes including clerodanes and abietanes, which represent about 27% and 4%, respectively, of all the diterpenes isolated (Xu *et al.*, 2018). A previous study of *C. oligandrus* has led to the isolation and identification of clerodane, trachylobane and labdane type diterpenes (Abega *et al.*, 2014).

Regarding Z. leprieurii, this is the first report on the occurrence of kaurane diterpenes in Z. leprieurii, and in the genus Zanthoxylum in general. Diterpenes are rare in the Rutaceae, and only a few including abietane, clerodane, kaurane and labdane have been reported from the genera *Citrus, Evodia, Fortunella, Glycosmis, Pamburus* and *Phebalium* (Cannon *et al.,* 1966; Dreyer and Park, 1975; Billet *et al.,* 1976; Seger *et al.,* 1998; Luis *et al.,* 2000; El-Shafae and Ibrahim, 2003). In addition to alkaloids, *Zanthoxylum* is known to be rich in coumarins and lignans, and contains few flavonoids (Adesina, 2005; Patino *et al.,* 2012; Li *et al.,* 2014). Alkaloids including acridone, alkamide, aporphine and benzophenanthridine, coumarins and lignans have been reported from *Z. leprieurii* (Adesina, 2005; Tchinda *et al.,* 2009). In the present study, in addition to kauranes, two flavonoids and a lignan previously reported from other *Zanthoxylum* species, have also been identified.

The biogenesis of diterpenes have extensively been studied. Diterpenes are now recognized to derive from geranylgeranyl diphosphate (GGPP) (Peters, 2010), though some few have been found to arise from carotene degradation (Seger *et al.*, 1998). Acid catalyzed cyclisation of GGPP can give rise to a fused decalin bicylic intermediate, copalyl diphosphate (CPP), from which subsequent modifications including internal additions, rearrangements and elimination may lead to bicyclic, tricyclic and tetracyclic diterpenes e.g. clerodane, abietane and kaurane, respectively (Figure 3.103) (Wilson *et al.*, 1976; Gershenzon and Croteau, 1993; Brunetton, 1999; Peters, 2010). Secondary modifications including oxidization and addition of different oxygen containing groups

may give rise to the range of diverse structural diterpenes occurring in nature (Gershenzon and Croteau, 1993) such as those isolated in the present study.

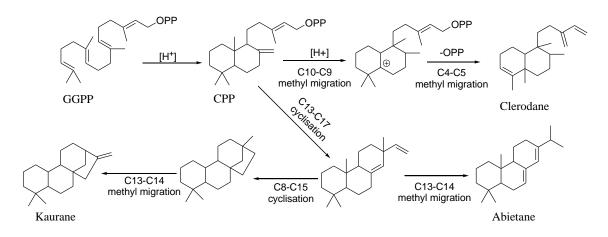


Figure 3.103 Summarised biosynthetic pathway toward abietane, clerodane and kaurane diterpenoids basic skeletons

#### 3.4.3 Flavonoids

Flavonoids were isolated from *J. hypocrateriformis* and *P. microcarpa*. This is the first phytochemical study of *P. microcarpa* and the third of *J. hypocrateriformis*.

Previous phytochemistry of *J. hypocrateriformis* have reavealed the presence of pyrrolidine alkaloids (Roessler *et al.*, 1978; Neukomm *et al.*, 1983). In the current study, in addition to pyrrolidine alkaloids, we have also isolated flavones. A similitude in terms of phytochemical constituents were found between *J. hypocrateriformis* and two other Acanthaceae species, *Justicia secunda* and *Graphtophyllum grandulosum* as all the compounds isolated, excluding the two new reported flavonoids **166** and **167**, have also been reported earlier in those two species.

Regarding *P. microcarpa*, two of the flavonoids isolated, **164** and **166**, have also been identified in *J. hypocrateriformis*. The genus *Pseudospondias* is constituted of two species, *P. microcarpa* and *P. longifolia*, and this study constitutes the first phytochemical

study of the genus. Species known today within *Pseudospondias* genus, were once members of the *Spondias* genus (Burkill, 1985). *Spondias* is a rich source of phenolic compounds including, tannins, flavonoids and phenolic acid derivatives (Sameh *et al.*, 2018). Flavonoids have been reported from *Spondias* as well as other Anarcadiaceae genus e.g. *Rhus* from which isovitexin and several glycosylated flavones and flavonols have been identified and isolated (Shrestha *et al.*, 2012; Abu-Reidah *et al.*, 2015).

Flavonoids are a large group of natural products. They usually occur in plants as glycosides (Ikan, 1991). Flavonoid subclasses are all related by a common biosynthetic pathway, which incorporates precursors from both the shikimate and acetate-malonate pathways. Condensation between *p*-coumarylCoA and three acetylCoA units results in the formation of a chalcone, the biogenetic intermediate to other structural subclasses of flavonoids (Figure 3.104) (Markham, 1982; Ikan, 1991; Schijlen *et al.*, 2004). Further modification of the flavonoids skeleton including addition (or reduction) of hydroxyl groups, arylation, dimerization and glycosylation may occur at various stages resulting in the diverse structural known flavonoids (Markham, 1982).

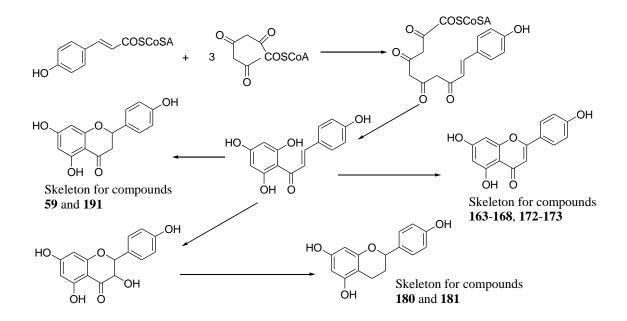


Figure 3.104 Summarised biosynthetic pathway toward basic flavonoid skeletons isolated

## 3.5 Bio-activity of the isolated compounds

## **3.5.1** Cytotoxicity of isolated compounds

The cytotoxicity of the isolated compounds (except **114**, **155**, **169**, **181** and sucrose) were evaluated against A549 (adenocarcinomic human alveolar basal epithelial), MCF7 (human breast adenocarcinoma), PC3 (human prostate cancer) and PNT2 (human normal prostate epithelium) cells using the MTT assay. The obtained  $LC_{50}$  values are shown in Table 3.21. Doxorubicin was used as positive control.

Table 3.21 Cell viability effect (LC<sub>50</sub> in  $\mu$ M) of isolated compounds against cancer and non-cancer (PNT2) cells

Plant	Cpds	LC <sub>50</sub> in µM					
COB		A549	MCF-7	PC-3	PNT2		
	7	136.8 ± 18.9	>200	172.3 ± 39.7	$167.5 \pm 25.3$		
	8	>200	>200	135.6 ± 21.1	>200		
	147	>200	>200	>200	>200		
	148	>200	>200	>200	$106.6 \pm 6.0$		
	149/150	>200	>200	160.9±36.2	>200		
	151	$106.6 \pm 27.2$	>200	>200	>200		
	152	>200	>200	131.1 ± 10.8	>200		
	153	63.8 ± 13.8	$136.2 \pm 22.7$	>200	>200		
	154	138.6 ± 22.1	$171.3 \pm 51.4$	>200	>200		
	156/157	$128.6\pm31.0$	>200	$111.2 \pm 2.9$	>200		
	158/159	>200	>200	>200	>200		
	160	>200	>200	153.0 ± 9.1	$97.6 \pm 8.9$		
	161	>200	>200	$68.9\pm6.6$	>200		
	162	>200	>200	>200	197.7 ± 14.9		
JHL	163	>200	>200	>200	>200		
	164	>200	>200	$146.6 \pm 20.4$	>200		
	165	>200	>200	128.1 ± 8.5	>200		
	166	>200	>200	87.1 ± 10.6	>200		

Table 3.21 *continued* 

Plant	Cpds		LC <sub>50</sub> in µM				
JHC		A549	MCF-7	PC-3	PNT2		
	167	>200	>200	117.4 ± 12.3	>200		
	168	>200	>200	>200	>200		
	170/171	>200	>200	>200	>200		
PMB an	103	>200	>200	125.7 ± 11.8	>200		
PML	172	>200	>200	>200	>200		
	173	>200	>200	112.5 ± 9.4	>200		
	174	>200	>200	180.5 ± 23.4	>200		
ZLF	175	>200	132.1 ± 14.9	33.3 ± 9.1	>200		
	176	>200	>200	138.9 ± 16.0	>200		
	177	82.0 ± 7.2	62.0 ± 1.8	$94.4 \pm 6.9$	10.6 ± 3.6		
	178	>200	>200	$147.9 \pm 14.1$	>200		
	179	116.8 ± 13.7	$167.9 \pm 10.1$	>200	>200		
	180	>200	>200	$112.7 \pm 7.5$	>200		
	182	>200	>200	149.5±21.9	>200		
ZZF	54	$108.5 \pm 22.1$	>200	33.4 ± 9.8	>200		
	55	113.4±0.15.8	53.7±09.5	164.7±0.21.3	$104.4 \pm 16.2$		
	59	29.5 ± 7.5	$74.2 \pm 17.8$	51.7 ± 8.7	$129.0 \pm 20.3$		
	79	>200	>200	>200	>200		
	128	$112.0 \pm 17.4$	>200	195.3 ± 22.6	>200		
	183	>200	$152.2 \pm 33.6$	>200	>200		
	184	>200	$172.2 \pm 31.4$	>200	>200		
	185	>200	>200	>200	>200		
	186	>200	>200	>200	>200		
	187	114.7 ± 18.3	$142.5 \pm 17.0$	>200	>200		
	188	>200	$153.6 \pm 32.7$	>200	>200		
	189	>200	>200	>200	181.6 ± 35.7		
	190	>200	>200	>200	>200		
	191	151.4 ± 25.4	>200	159.7 ± 28.5	>200		
	Doxy	1.3 ± 0.3	0.7 ± 0.1	16.4 ± 2.9	1.5 ± 0.3		

Data are represented as mean  $\pm$  SEM (n = 3); LC<sub>50</sub> = sample concentration that caused

50% of cells death

The LC<sub>50</sub> values of the screened compounds ranged from 29.5 to 138.6  $\mu$ M (A549), 53.7 to 172.2  $\mu$ M towards MCF-7 cells, 33.3 to 195.3  $\mu$ M (PC-3), and 10.6 to 197.7  $\mu$ M (PNT2). The isolated flavonoids from JHL, PMB and PML were found to have selective effect on PC-3 cells. None of the isolated compounds has shown a significant cytotoxic effect. They have showed moderate or low cytotoxic effect at the tested concentrations. A pure compound to be consider as promising candidate for drug development should have its LC<sub>50</sub> value below 25  $\mu$ M (Choudhary and Thomsen, 2003). Among the tested compounds, kaurenoic acid (**175**) and sesamin (**54**) were found to exert reasonable cytotoxicity against PC3 with IC<sub>50</sub> both about 33  $\mu$ M. Hesperidin (**59**) and *ent*-kauran-16 $\beta$ -ol-19-oic acid (**177**) were the only compounds with wide active spectrum, having effect against all the cell lines tested even on the normal prostate cells PNT2 with IC<sub>50</sub> 129.0 ± 20.3 and 10.6 ± 3.6  $\mu$ M, respectively. Hesperidin was the most active compound with IC<sub>50</sub> 29.5 ± 7.5, 74.2 ± 17.8 and 51.7 ± 8.7  $\mu$ M against A549, MCF-7 and PC3, respectively.

#### **3.5.2** Chemopreventive activity of isolated compounds

3.5.2.1 Cytotoxicity of the compounds against AREc32 cells

Compounds isolated from plants extracts, which have shown potent chemopreventive properties, were screened to assess their chemopreventive property by evaluating their effect on the activation of the level of luciferase in AREc32 cells. This include compounds isolated from *C. oligandrus* and *Z. zanthoxyloides*. The toxicity of the compounds on the viability of AREc32 cells was first determined by the MTT assay to find a suitable concentration at which to conduct the following luciferase assay which would not cause significant cell death (concentrations causing more than 10% cell death, relative to control were not used, Table 3.22).

C. oligandrus					Z. zanthoxyloides			
Cpds	LC <sub>10</sub>	Cpds	LC <sub>10</sub>	Cpds	$LC_{10}$	Cpds	LC <sub>10</sub>	
	(µM)		(µM)		(µM)		(µM)	
7	6.0	153	12.5	54	6.0	185	12.5	
8	12.5	154	12.5	55	6.0	186	12.5	
147	25.0	156/157	6.0	59	6.0	187	6.0	
148	25.0	158/159	6.0	79	25.0	188	6.0	
149/150	25.0	160	25	128	25.0	189	6.0	
151	6.0	161	6.0	183	12.5	190	25.0	
152	6.0	162	6.0	184	12.5	191	12.5	

Table 3.22 Least toxic concentration (no more than 10% cells death) of isolated compounds against AREc32 cells

## 3.5.2.2 Luciferase activity

The luciferase activity was measured using the Steady-Glo luciferase kit provided by Promega Corp. After the exposure of AREc32 cells for 24 h to *t*BHQ 6 and 12  $\mu$ M, the activity of luciferase increased by 2.51 and 7.37-fold compared to the control (Figure 3.105). Among the tested compounds, the mixtures of epimeric clerodanes **156/157** and **158/159** (crotonolins A-D), skimmianine (**55**), hesperidin (**79**) and myrtopsine (**185**) were found to produce 2.7, 2.4, 2.4, 1.8 and 2.0-fold induction, respectively. Other tested compounds had slight or negligible effects on the luciferase activity at the dosage tested. The results suggested the above active compounds might be potential activators of the NrF2 pathway, meaning potent chemopreventive compounds at non cytotoxic dosages. The diterpenes **156-159** were the most activitors of the Nrf2 activity. Their activities were comparable to that *t*BHQ when tested at the same concentration. Several diterpenes including clerodanes have been reported to possess cytoprotective and cytotoxic properties (Thoppil and Bishayee, 2011; Islam, 2017). For example, the clerodane columbin has been revealed to have chemoproventive activity against colorectal cancer when administered as a diet to male rats in the early phase of azoxymethane-induced colon carcinogenesis (Kohno *et al.*, 2002). Therefore, clerodanes, represent an interesting group of compounds to explore for the discovery of cancer chemopreventive agents.

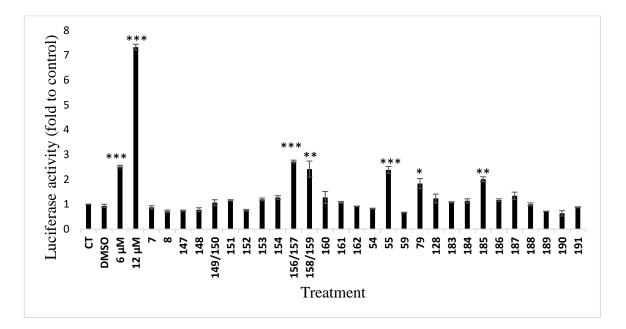


Figure 3.105 Luciferase activity of isolated compounds

AREc32 cells were seeded in 96-well plates at 1.2 x  $10^4$  cells/well. After 24 h, *t*BHQ (6 and 12  $\mu$ M) and the compounds were added to the medium. The cells were then incubated for another 24 h and assayed for luciferase activity as detailed in Chapter 2, 2.4.4. The value of luciferase activity of untreated cells (CT) was set at 1. Values shown are mean  $\pm$  SEM of three experiments. \*p < 0.01; \*\*p < 0.001, \*\*\*p < 0.0001, significantly increased versus control. DMSO 0.1%: (cells treated with 0.1% DMSO medium).

## 3.5.3 Haem polymerisation assay of the isolated compounds

The isolated compounds from extracts, which have shown antimalarial activity, were screened to evaluate their antimalarial potential using haem polymerisation assay. Quinoline antimalarials in use today, such as chloroquine, mefloquine and quinine, have been reported to act in the erythrocytic stage of the parasite life cycle by blocking the polymerisation of free haem generated during haemoglobin digestion to its polymer haemozoin, a step which is essential for the parasite survival (Foley and Tilley, 1998). None of the isolated compounds have demonstrated an effect at the tested concentration (0-250  $\mu$ M). These compounds included the diterpenes isolated from *C. oligandrus* and *Z. leprieurii*, and flavonoids from *J. hypocrateriformis* and *P. microcarpa*. These results were not consistent with our early finding regarding the activity associated with the extracts from which the compounds have been isolated. However, as previously noted, the activity of an extract is the result of different interactions occurring between the chemical constituents within the extract. In addition, the isolated compounds might not be iron chelators making it difficult to bind with haem via the ion present at its center (Sarker *et al.*, 2016), hence their inactivity.

# Chapter 4 Conclusion and Future Prospects

# 4. Conclusion and future prospects

Bioassay-guided isolation of active compounds was carried out with five Cameroonian medicinal plants, including *Croton oligandrus* Pierre ex Hutch (Euphorbiaceae), *Justicia hypocrateriformis* (Vahl) Milne-Redh (Acanthanceae), *Pseudospondias microcarpa* (A. Rich.) Engl. (Anacardiaceae), *Zanthoxylum lepreurii* Guill. and Perr. and *Zanthoxylum zanthoxyloides* (Lam.) Zepern. and Timler (Rutaceae).

The *n*-hexane and the DCM extracts of *Croton oligandrus* bark, and the DCM and MeOH extracts of *Z. zanthoxyloides* fruits were active in the luciferase assay causing 18, 21, 34 and 36-fold induction of the level of luciferase in AREc32 cells, respectively. The diterpenes crotonolins A-D, isolated from the DCM extract of *C. oligandrus* were identified as the most active principle of the plant with 2-fold or greater. From *Z. zanthoxyloides*, skimmianine isolated from both the DCM and MeOH extracts, and hesperidin and myrtopsine isolated from the MeOH extract, increased the level of luciferase by 2.4, 1.8 and 2.0-fold, respectively.

The antimalarial activity of the selected plants was evaluated using the haem polymerisation assay. *Pseudospondias microcarpa* was the most active plant with IC<sub>50</sub>  $73.9 \pm 25.8$ ,  $2.5 \pm 1.5$  and  $4.0 \pm 1.7 \mu$ M for the stem bark *n*-hexane, DCM and MeOH extracts, respectively, and  $13.0 \pm 9.0 \mu$ M for the leaves MeOH extract. No inhibition of haem polymerisation was observed for the isolated compounds at the assayed concentrations.

Three major classes of phytochemicals including alkaloids, diterpenes and flavonoids were isolated from the five plants studied. Some of the compounds, 12-*epi*-megalocarpoloide D (**154**), crotonolins A-F (**155-160**), justicialosides A and B (**166-167**) and zanthoamides G-I (**188-190**) were being reported for the first time from natural sources. Kaurane diterpenes were reported for the first time from the genus *Zanthoxylum* 

and the quinoline alkaloids *N*-methylplatydesminium cation (**183**), isoplatydesmine (**184**), myrtopsine (**185**), ribalinine (**186**) and *N*-methylatanine (**187**) from the species *Z*. *zanthoxyloides* for the first time. Other isolated compounds included two triterpenes, acetyl aleuritolic acid (**7**) and lupeol (**8**); two lignans, sesamine (**54**) and icariside D2 (**183**); ferulic acid derivatives (**114**, **147-150**); a quindoline, 10*H*-quindoline (**169**); a coumarin, scopoletin (**103**); and a chromanone, pithecellobiumol B (**174**). The cytotoxicity of all the isolated compounds were evaluated against three cancer cell lines (A549, MCF7 and PC3) and a non cancer cell line (PNT2). Hesperidin (**59**) was the most cytotoxic compound with LC<sub>50</sub> 29.5  $\pm$  7.5, 74.2  $\pm$  17.8, 51.7  $\pm$  8.7 and 129.0  $\pm$  20.3  $\mu$ M against A549, MCF7, PC3 and PNT2, respectively.

This study generated the first, second and third phytochemical report of *P. microcarpa*, *C. oligandrus* and *J. hypocrateriformis*, respectively. The chemotaxonomy of the isolated compounds has also been discussed. *J. hypocrateriformis* and *J. secunda* were observed to have similar compounds and should be investigated further to establish whether the two names refer to the same species, and kaurane diterpenes were identified as new markers of *Z. leprieurii*.

This study also supports the use of *C. oligandrus* and *Z. zanthoxylum* as alternative and complementary medicine for the treatment and prevention of cancer, and *P. microcarpa* for the treatment of malaria. However, further studies need to be carried out to define standardised dosage as those plants were found to contain moderate cytotoxic compounds against the human normal prostate epithelium cells (PNT2).

Future studies may involve:

 chemical modification of active compounds as well as the novel compounds and evaluation of the activity of obtained analogues;

- investigation of the mechanism of action of chemopreventive extracts and compounds identified;
- investigation of the antimalarial activity of isolated compounds using a different assay e.g. *in vitro* against parasites;
- investigation of synergism, antagonism and additive interactions as a contributor to activity of crude plant extracts specially those active in the antimalarial assay.

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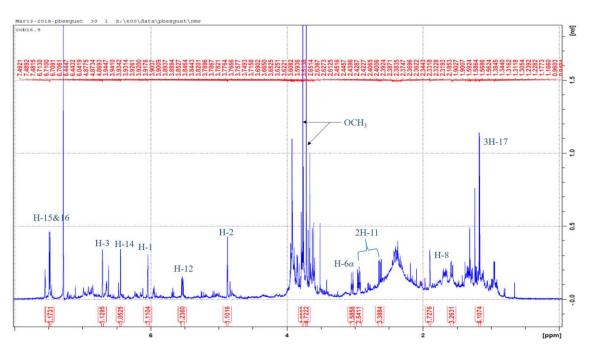
Zhao, P., Dou, Y., Chen, L., Li, L., Wei, Z., Yu, J., Wu, X., Dai, Y. and Xia, Y. (2015). SC-III3, a novel scopoletin derivative, induces autophagy of human hepatoma HepG2 cells through AMPK/mTOR signaling pathway by acting on mitochondria, *Fitoterapia*, 104, 31-41.

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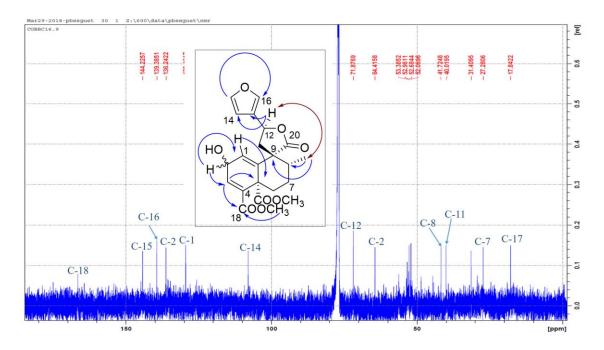
### Appendix I MS, <sup>1</sup>H and <sup>13</sup>C NMR spectra of crotonolin E (155)

White amorphous powder,  $[\alpha]_D^{25}$  -70.0 (c 0.00005, MeOH).

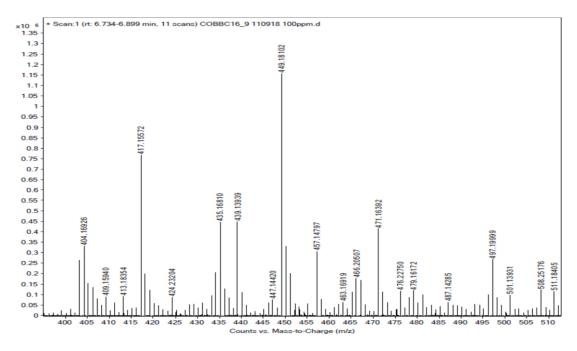
Molecular formula  $C_{22}H_{24}O_8$  determined from the peak at m/z 417.1557 calculated for  $C_{22}H_{25}O_8$  [M+H]<sup>+</sup>, 417.1543



<sup>1</sup>H NMR spectrum (CDCl<sub>3</sub>, 600 MHz) of **155** 

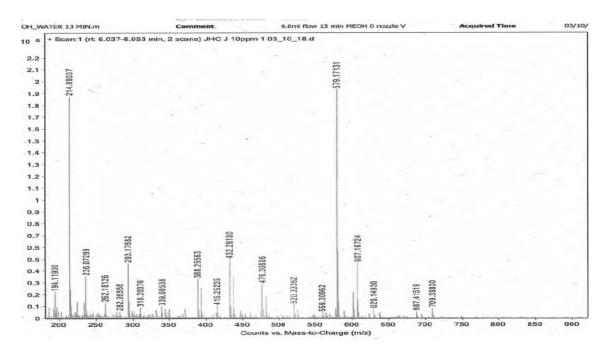


<sup>13</sup>C NMR spectrum (150 MHz, CDCl<sub>3</sub>) and Keys <sup>1</sup>H-<sup>1</sup>H NOESY ( $\leftrightarrow$ ) and HMBC ( $\rightarrow$ ) correlations of **155** 

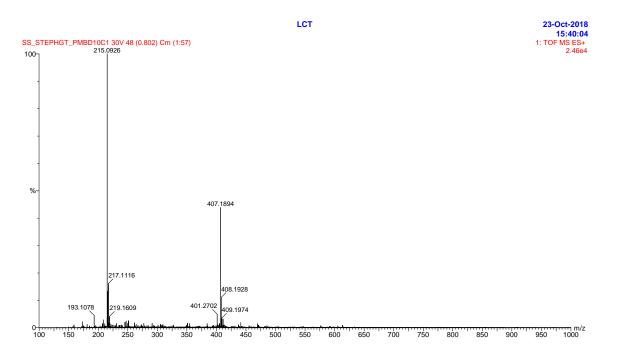


ESI-MS spectrum of 155

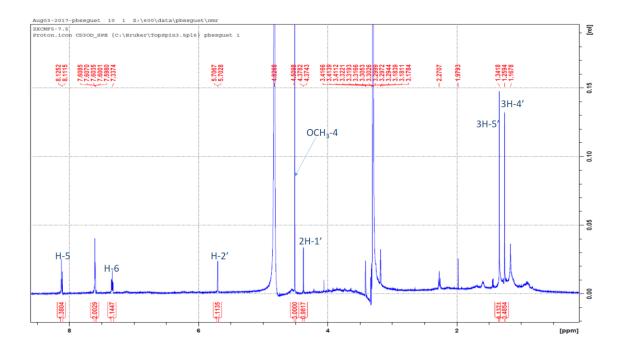




## **Appendix III ESI-MS spectrum of scopoletin (103)**



# Appendix IV <sup>1</sup>H NMR (600 MHz, CD<sub>3</sub>OD) spectrum of myrtopsine (185)



#### Appendix V List of publications and presentations

#### **Publications**

**Guetchueng, S.T.**, Nahar, L., Ritchie, K.J., Ismail, F.M., Wansi, J.D., Evans, A. and Sarker, S.D. (2018). Zanthoamides G-I: Three new alkamides from *Zanthoxylum zanthoxyloides*. *Phytochemistry Letters*, 26, 125-129.

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**Guetchueng, S. T.**, Nahar, L., Ritchie, K. J. and Sarker, S. D. (2018). Assessment of potential antimalarial activity of selected Cameroonian medicinal plants. Oral presentation- PSE-YSM 2018, Liverpool, UK, 02-05 July 2018.

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**Guetchueng, S. T.**, Nahar, L., Ritchie, K. J. and Sarker, S. D. (2017). Screening of extracts of six selected medicinal plants as activators of the keap1-Nrf2 pathway to assess potential cancer chemopreventive activity. Oral presentation- The 9<sup>th</sup> International Conference of the Kenya Chemical Society (KCS), Kenya, 9-12 May 2017.