CHEMICAL CONSTITUENTS FROM THE STEMS OF Garcinia cylindrocarpa AND THE STEM BARK OF Garcinia picrorhiza AND Garcinia tetrandra



A Dissertation Submitted in Partial Fulfillment of the Requirements for the Degree of Doctor of Philosophy in Chemistry Department of Chemistry FACULTY OF SCIENCE Chulalongkorn University Academic Year 2019 Copyright of Chulalongkorn University

องค์ประกอบทางเคมีจากลำต้น Garcinia cylindrocarpa เปลือกลำต้น Garcinia picrorhiza และ Garcinia tetrandra



วิทยานิพนธ์นี้เป็นส่วนหนึ่งของการศึกษาตามหลักสูตรปริญญาวิทยาศาสตรดุษฎีบัณฑิต สาขาวิชาเคมี คณะวิทยาศาสตร์ จุฬาลงกรณ์มหาวิทยาลัย ปีการศึกษา 2562 ลิขสิทธิ์ของจุฬาลงกรณ์มหาวิทยาลัย

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เอ็ดวิน ริสกี้ ซูกานดา : องค์ประกอบทางเคมีจากลำต้น Garcinia cylindrocarpa เปลือกลำต้น Garcinia picrorhiza และ Garcinia tetrandra. (CHEMICAL CONSTITUENTS FROM THE STEMS OF Garcinia cylindrocarpa AND THE STEM BARK OF Garcinia picrorhiza AND Garcinia tetrandra) อ.ที่ปรึกษาหลัก : รศ. ดร.สันติ ทิพยางค์

เพื่อการค้นหาสารออกฤทธิ์ทางชีวภาพชนิดใหม่ จากพืชสกุล Garcinia จึงได้เลือกทำการศึกษาองค์ประกอบทางเคมี และฤทธิ์ทางชีวภาพจากพืชสกุล Garcinia 3 ชนิด และทำการเก็บตัวอย่างพืชเหล่านี้จากประเทศอินโดนีเซีย สำหรับการแยกองค์ประกอบทางเคมีจากพืชสกุล Garcinia ให้ บริสุทธิ์จะ ต้องอาศัยเทคนิคทางโครมาโทกราฟี และในการพิสูจน์ทราบเอกลักษณ์ทางโครงสร้างของสารบริสุทธิ์ ต้องอาศัยการวิเคราะห์ข้อมูลทาง สเปกโทรสโกปี จากนั้นสารบริสุทธิ์ทั้งหมดที่แยกได้จะถูกนำไปทดสอบความเป็นพิษต่อเซลล์มะเร็ง 5 ชนิด (HeLa S3, Hep G2, HT-29, MCF-7 และ KB) ด้วยวิธี MTT assay สาร xanthones ชนิดใหม่ 4 ชนิด ชื่อ cylindroxanthones D-G และสาร biphenyls ชนิดใหม่ 2 ชนิด ชื่อ cylindrobiphenyls A และ B รวมทั้งสารที่เคยมีการรายงานมาแล้วอีก 28 ชนิด ถูกแยกได้จากลำต้น ของ G. cylindrocarpa สาร xanthones ชนิดใหม่ 9 ชนิด ชื่อ tetrandraxanthones A-I และสารที่เคยมีการรายงานมาแล้วอีก 22 ชนิด ถูกแยกได้จากเปลือกลำต้น ของ G. tetrandra สาร polyprenylated benzoylphloroglucinols (PPBPs) ชนิดใหม่ 8 ชนิด ชื่อ picrorhizones A-H และสารที่เคยมีการรายงานมาแล้วอีก 4 ชนิด ถูกแยกได้จากเปลือกลำต้น ของ G. *picrorhiza* ผลจากการทอสอบพบว่า สาร 9-hydroxycalabaxanthone มีความเป็นพิษต่อเซลล์มะเร็งทั้ง 5 ชนิด โดยมีค่า IC₅₀ ในช่วง 1.6–3.4 µM ในขณะที่สาร 2deprenylrheediaxanthone B และ picrorhizone F มีความเป็นพิษต่อเซลล์มะเร็งทั้ง 4 ชนิด ยกเว้นเซลล์มะเร็ง HT-29 โดยมีค่า IC₅₀ ในช่วง 2.2 ถึง 9.4 μM

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Aiming to discover novel bioactive secondary metabolites from genus Garcinia, we performed phytochemical and biological investigations of three Garcinia species collected from Indonesia. A series of chromatographic techniques and spectroscopic data analysis were used to isolate and characterize the Garcinia phytochemicals. The cytotoxicity of the isolated compounds was evaluated against five human cancer cell lines (HeLa S3, Hep G2, HT-29, MCF-7, and KB) using MTT assay. Four new xanthones, cylindroxanthones D-G, and two new biphenyls, cylindrobiphenyls A and B, together with 28 known compounds were successfully isolated from G. cylindrocarpa stems. From the stem bark of G. tetrandra, nine new xanthones, tetrandraxanthones A-I, were identified along with 22 previously described analogues. Eight new polyprenylated benzoylphloroglucinols (PPBPs), picrorhizones A-H, and four known analogues were reported from the stem bark of G. picrorhiza. Among the tested compounds, 9-hydroxycalabaxanthone showed significant cytotoxicity against five human cancer cell lines with IC₅₀ values in the range of 1.6–3.4 μ M, while 2deprenylrheediaxanthone B and picrorhizone F were active against four cancer cells, except HT-29, with IC₅₀ values ranging from 2.2 to 9.4 μ M.

Field of Study:	Chemistry	Student's Signature
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LIST OF ABBREVIATIONS

- HR-ESI-MS High Resolution Electrospray Ionization Mass Spectroscopy
- ECD Electronic Circular Dichroism
- ¹H NMR Proton Nuclear Magnetic Resonance
- ¹³C NMR Carbon-13 Nuclear Magnetic Resonance
- 1D-NMR One-Dimensional Nuclear Magnetic Resonance
- 2D-NMR Two-Dimensional Nuclear Magnetic Resonance
- COSY ¹H⁻¹H Correlation Spectroscopy
- HSQC Heteronuclear Single Quantum Correlation
- HMBC Heteronuclear Multiple Bond Correlation
- NOESY Nuclear Overhauser Effect Spectroscopy
- *m/z* mass per charge number of ions (Mass Spectroscopy)
- $\delta_{
 m H}$ chemical shift of proton (NMR)
- $\delta_{
 m C}$ chemical shift of carbon (NMR)
- J coupling constant (NMR) MODULA
- s singlet (NMR)
- d doublet (NMR)
- t tirplet (NMR)
- m multiplet (NMR)
- brs broad singlet (NMR)
- dd doublet of doublets (NMR)
- tt triplet of triplets (NMR)
- DMSO-d₆ deuterated dimethyl sulfoxide

- CDCl₃ deuterated chloroform
- Acetone-d₆ deuterated acetone
- MeOH-d₆ deuterated methanol
- calcd. calculated
- m.p melting point
- TLC Thin Layer Chromatography
- MTPA α -methoxy- α -trifluoromethylphenylacetic acid in Mosher's method
- PPBP polyprenylated benzoylphloroglucinol
- *i*NOS inducible nitric oxide synthase
- COX cyclooxygenase enzyme
- MTT 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide, a reagent in cytotoxic assay
- EC₅₀ the molar concentration of an agonist that produces 50% of the maximal possible effect of that agonist
- IC₅₀ the molar concentration of an antagonist that reduces the response to an agonist by 50%
- CC₅₀ the concentration of a compound required for the reduction of cell viability by 50%

Chapter I Introduction

Cancer remains one of serious health problems across the world and caused over nine million deaths in 2018. In general, cancer arise from alteration of genes which control cell functions, especially in cell growth and division [1]. As the result, cells with certain genetic changes will divide out of control. These abnormal and malignant cells become invasive by spreading into surrounding tissues and inducing normal cells to form blood vessels for their oxygen and nutrients supply. In many cases, metastasis also occurs where cancer cells are able to travel through blood or lymph vessels to other parts of the body and form other cancerous tumors [2, 3]. Several effective cancer treatments have been developed, including chemotherapy, in which drugs having anticancer properties are used to inhibit growth of cancer cells and its metastasis, restrict cancer invasion, and increase cancer cells death [4]. Recent advance of chemotherapy has been developed by targeting specific enzymes and growth factor receptors involved in carcinogenesis as many previous chemotherapeutic drugs had broad spectrum of cells destruction causing numerous side effects [5].

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Natural products play a pivotal role in an attempt to discover potential chemotherapeutic agents and over 60% of current anticancer drugs are bioactive substances originally produced by terrestrial plants, marine invertebrates (e.g. mollusks, sponges, echinoderms, cnidarians, and tunicates), and microorganisms (e.g. bacterial and fungal cultures) [6]. The secondary metabolites are produced in a limited number and used by specific living organisms as response to protect and adapt in various environmental conditions. Their roles in nature provide an insight to explore those bioactive compounds for human health benefit with the major use in cancer therapy [7]. Due to relatively ease to access in sample collection and

preparation, medicinal plants or herbs are frequently targeted to find potential anticancer candidates and a million of phytochemicals has been hitherto isolated and pharmacologically screened [8].



Figure 1.1. Natural product-derived compounds as anticancer agents.

Vinblastine (1.1) and vincristine (1.2), two vinca alkaloids originally isolated from *Catharanthus roseus* G. Don. (common name: Madagascar periwinkle), were the first introduced anticancer drugs prescribed to combat childhood leukemia, Hodgkin's lymphoma, and testicular teratoma. An extensive research revealed that both compounds were capable to destabilize microtubules and obstruct cancer cells during metaphase stage causing the cells apoptosis [9, 10]. Later, the discovery of paclitaxel (Taxol[®]) (1.3), a bioactive constituent isolated from the bark of *Taxus brevifolia* Nutt. (common name: Pacific yew), became one of great breakthroughs in cancer research area. This clinically effective chemotherapeutic drug was found to be active against a variety of human cancers, including ovarian, non-small cell lung,

breast, and Kaposi's sarcoma by promoting tubulin heterodimers polymerization and suppressing dynamic changes in microtubules leading to mitotic obstruction [9]. Several structural modifications of the above natural product-based anticancer agents were also carried out to overcome clinical trials limitation, such as low solubility which affect drug delivery process, high toxicity causing several side effects, and drug resistance associated with the current drugs. Etoposide (1.4), a synthetic analog of podophyllotoxin isolated *Podophyllum peltatum* L. (common name: American mandrake/mayapple), was approved for ovarian, lymphoma, testicular teratoma, small-cell lung cancers treatment. The latter anticancer agent was found to have different mode of action to kill cancer cells via topoisomerase II inhibition and give higher efficacy than its precursor [7, 9].

The above-mentioned data illustrate a potency of natural products as chemotherapeutic agents and exploration of bioactive secondary metabolites, especially from medicinal plants, is highly recommended to be done. This research will focus on the phytochemical and biological investigation *Garcinia* plants which have been reported previously to contain many pharmacologically active substances.

ุหาลงกรณ์มหาวิทยาลัย

1.1. Genus Garcinia: botanical aspect, distribution, and beneficial uses

The genus *Garcinia*, one of the largest genera to the family Clusiaceae, is widely distributed in tropical rain forests of Southeast Asia, South America, and West and Central Africa. This genus is pantropical and comprise high level of species diversity with more than 250 species of evergreen, lactiferous, dioecious, and small shrubs to medium-sized trees [11, 12]. There are approximately 77 species of this genus distributed throughout Indonesia, with the main population in Sumatera, Kalimantan, and Papua Islands [13]. In its natural habitat, *Garcinia* species can be found along rivers, swamps, and streams as wild plants or cultivated in garden and widely grow on sandy, clay, chalk, silt, peat, and loam soils [14, 15].

Some *Garcinia* plants have been consumed traditionally by local people as Indonesian dishes and herbal medicines to prevent and treat several health problems. The fruits of *G. mangostana* (local name: manggis), well known as mangosteen, have mixed sweet and sour taste and become one of potential commodities in market. The pericarps of mangosteen are extracted for additive in herbal soap and food [12, 15]. *G. atroviridis* fruits (local name: Gelugur) are edible with sour taste and used for food spices after drying and grinding process and its leaves are prepared for food salad. In medical aspect, the infusion of the leaves of *G. atroviridis* and *G. parvifolia* (local name: Kandis) is consumed for stomach pain remedy associated with pregnancy. The latex of *G. mangostana* is topically applied for oral ulcer treatment, while the decoction of its inner bark is consumed to treat dysentery [15, 16]. The pericarps of *G. mangostana* are prepared as a gargling solution for oral hygiene protection [17]. Moreover, the pericarps of *G. mangostana* and *G. cambogia* are formulated as a herbal supplement for weight loss and natural antioxidant [18].

1.2. Biosynthesis of Garcinia secondary metabolites

Phytochemical investigation of the genus *Garcinia* resulted in the isolation of **CHULALONGKORN ONIVERSITY** structurally diverse secondary metabolites, especially phenolic compounds, such as xanthones, biphenyls, polyprenylated benzoylphloroglucinols, isocoumarins, depsidones, and biflavonoids. Biosynthetically, amino acids L-phenylalanine or L-tyrosine, derived from shikimate pathway, act as a C_6C_3 building block and precursor for the phenolic metabolites. Plant employs phenylalanine ammonia lyase (PAL) enzyme to eliminate ammonia of the amino acids and produce cinnamic acid (Figure 1.2). The hydration of cinnamic acid via coenzyme A ester substrate (cinnamoyl-CoA) and oxidation of the hydrated intermediate generate a β -ketoester and subsequently lose acetyl-CoA through reverse Claisen rearrangement to introduce a CoA-ester of benzoic acid. The chain elongation of benzoyl-CoA with three malonyl-CoA derived from acetate pathway generates a shikimate-acetate intermediate I, which is involved later in an intramolecular cyclization to yield several Garcinia phenolics [19]. Claisen condensation of intermediate I catalyzed by benzophenone synthase (BPS), a type III polyketide synthase, afford a benzophenone, 2,4,6-trihydroxybenzophenone. This metabolite is oxidized by benzophenone 3'-hydroxylase (B3'H) to give 2,4,6,3'tetrahydroxybenzophenone and converted to xanthones by 1,3,5-/1,3,7trihydroxyxanthone synthases (TXSs) via intramolecular oxidative C-O coupling reaction [20]. Further oxidation of xanthones via Baeyer-Villiger rearrangement by specific enzymes gives a depsidone, which is rarely found in higher plants. In general, depsidones are well known as lichen secondary metabolites derived polyketide biosynthesis pathway [21]. A biphenyl-type compound, 3,5-dihydroxybiphenyl, is constructed through aldol condensation and enolization of the intermediate I by biphenyl synthase (BIS) [22].

As depicted in Figure 1.3, a simple benzophenone can be converted to more complex structure, so-called polyprenylated benzophenone (PPB)/benzoylphloroglucinol (PPBP), via polyprenylation which utilize plant enzymes and DMAPP (dimethylallyl pyrophosphate), a C₅-building block from mevalonate pathway [23]. Multiple prenylation of benzophenone generate two type-D PPB intermediates and different attacking position at phloroglucinol A-ring results various type of PPBs. A C-5 enol attack to the double bond of prenyl unit at C-1 generates type-C PPB, while C-1 and C-3 attacking position to a prenyl moiety at C-5 introduce type-A and —B PPBs, respectively. Such enzyme-catalyzed modifications afford various PPBs with a high degree of complexity [24].







Figure 1.3. Biosynthesis of polyprenylated benzophenone in plant.

The biosynthesis of flavonoid and isocoumarin lie to the precursor CoA-ester of cinnamic acid with three malonyl-CoA which give intermediate II with chain extension, as described in Figure 1.4. Later, this intermediate is cyclized via Claisen condensation, followed by enolization to form naringenin, a flavonoid-type secondary metabolite. Intermolecular oxidative C-C and C-O coupling reaction initiated by specific plant enzymes (e.g. peroxidase) produce biflavonoid structures [19]. The aldol condensation and enolization of intermediate II give intermediate III which is subsequently hydrated and lactonized to generate an isocoumarin [25].



Figure 1.4. Biosynthesis of isocoumarin, flavonoid, and biflavonoid in plant.

1.3. Diverse structures of Garcinia secondary metabolites

In this section, several secondary metabolites which have been reported previously are briefly reviewed based on the core structures and substituent modifications. The biological and pharmacological results of the isolated compounds are also described and restricted to those having potent activities, particularly in anticancer study. This is due to the fact that *Garcinia* phytochemicals, particularly xanthone and benzophenone derivatives, are well recognized to exhibit significant effects on cancer growth inhibition, such as α -mangostin (1.8), gambogic acid (1.32),

and garcinol (**1.76**), wherein detailed mechanism of action of those compounds against cancer cells are described below.

1.3.1. Xanthones



Figure 1.5. Simple oxygenated and prenylated xanthones (1.5–1.19) from *Garcinia* plants.

Xanthone is an oxygenated heterocyclic compound bearing dibenzo- γ -pyrone skeleton and derived biosynthetically from shikimate-acetate pathway, as described

in Figure 1.2. Xanthone derivatives can be found in higher plants, microorganisms, marine sources, and lichens, while they are isolated as a chemotaxonomic marker and major component of the genus *Garcinia* with high chemical diversity [26].

Most common xanthones found in Garcinia plants bear oxygenated functional groups such as hydroxy and methoxy substituents, as well as prenyl and geranyl side chains (Figure 1.5). For example, three tri- and tetra-oxygenated xanthones, 1,3,6,7-tetrahydroxyxanthone (1.5), 1,6-dihydroxy-7-methoxyxanthone (1.6), and 1,6-dihydroxy-3,7-dimethoxyxanthone (1.7), were obtained from the leaves and twigs of G. multiflora [27]. Another phytochemical investigation led to the isolation of prenylated xanthone α -mangostin (1.8) and its analogues, garcinone D (1.9) and fuscaxanthone D (1.10), with hydrated and hydroxylated prenyl side chains from the roots of G. cowa and all compounds showed moderate cytotoxicity against KB and HeLa cancer cells with IC₅₀ values ranging from 11.7 to 33.5 μ M. Moreover, α magostin, originally isolated from the pericarps of mangosteen (G. mangostana), had a cytostatic effect on colon cancer cells by inducing G1 cell cycle arrest and promote the suppression of breast tumor growth in vivo [28]. Xanthones with modified prenyl moieties were found in the leaves of G. bracteata and G. polyantha, such as allaxanthone A (1.11) and bracteaxanthone VIII (1.12) bearing a dimethylallyl unit and O-prenylated polyanxanthone A (1.13) [29, 30].

Three furanoxanthones were obtained from the roots of *G. fusca* and the stem bark of *G. dulcis*, including garbogiol (1.14), garciniaxanthone D (1.15), and subelliptenone C (1.16), whereas three pyranoxanthones, ananixanthone (1.17) and nigrolineaxanthones Q and X (1.18 and 1.19), were isolated from the leaves of *G. nigrolineata* [31-33]. The five- and six-membered ring moieties of the six xanthones are formed from cyclization of prenyl unit and its adjacent hydroxy group (Figure 1.5). Compound 1.14 was found to inhibit α -glucosidase enzyme with IC₅₀ value of 21.2

 μ M, while **1.18** exhibited antibacterial activity against *Micrococcus luteus* with MIC value of 8 μ g/mL.



Figure 1.6. Xanthones with extended prenyl units (1.20–1.25) from Garcinia plants.

The extension of C₅-prenyl side chain on a xanthone skeleton was found in the extracts of *Garcinia* species, such as cowanol (1.20) with a C₁₀-geranyl substituent, garciniacowone H (1.21) bearing a C₁₅-farnesyl unit, and garciniacowone I (1.22) substituted by a C₂₀-geranylgeranyl side chain (Figure 1.6) [34]. The three phytochemicals were obtained from fractionation of the acetone extract of *G. cowa* leaves and they demonstrated NO (nitric oxide) production inhibition in LPS-induced RAW 263.7 macrophage cells and α -glucosidase inhibition with IC₅₀ values in the range of 5.8–13.4 μ M and 15.4–28.7 μ M, respectively. Modification of a geranyl functionality was observed in kaennacowanol B (1.23) and fuscaxanthone J (1.24) isolated from *G. cowa* and *G. fusca* roots via intramolecular cyclization to give furan and pyran rings [31, 35], while a metabolite with an uncommon 7-oxo-[2.2.1]-bicyclic system, parvixanthone I (**1.25**), was obtained from the bark of *G. parvifolia* (Figure 1.6) [36].



Figure 1.7. Rearranged polyprenylated and caged xanthones (**1.26–1.34**) from *Garcinia* plants.

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Extensive phytochemical studies on the leaves and twigs of G. oligantha resulted in discovery of unique xanthones, including dihydro and tetrahydroxanthones garoliganthin I (1.26) and garoliganthin H (1.27), as well as (-)garoliganthin F (1.28), (-)-garoliganthin G (1.29), (+)-oliganthin T (1.30), and (-)garoliganthin B (1.31) featuring structurally rearranged xanthone cores (Figure 1.7). The formation of such unprecedented scaffolds in the ring A of 1.28-1.31 was biosynthetically proposed by pinacol rearrangement to form 1.28 and 1.29 and an oxidative ring cleavage followed by a series of enzyme-catalyzed oxidation and hydration reactions and intramolecular cyclization to achieve 1.30 and 1.31 [37, 38]. Gambogic acid (1.32), gaudichaudiic acid (1.33), and epi-gambogic acid C (1.34), originally exist in *G. hanburyi* and *G. gaudichaudii*, represent another class of complex modified xanthones with 4-oxo-tricyclo[$4.3.1.0^{3,7}$]dec-8-en-2-one ring system containing a highly substituted tetrahydrofuran moiety, so-called caged xanthones [28, 39, 40]. Additionally, compound 1.34 also bears a bridged tricyclic ring moiety which is derived from an unusual cyclization of geranyl functionality at the xanthone skeleton (Figure 1.7). Cytotoxic evaluation showed that 1.33 was active against P388/DOC and Messa cell lines with IC₅₀ values of 3.4 and 3.8 μ g/mL. Moreover, pharmacological study revealed that 1.32 was capable to induce apoptosis, reverse multidrug resistance of cancer cells, inhibit cell proliferation, and anti-angiogenic activities and it has been completed a phase IIa clinical trial in China for patients with lung, colon, and renal cancers.

xanthone possessing an unusual α,β -unsaturated- γ -lactone unit. А garcinexanthone F (1.35), was obtained from G. xanthochymus bark with its scavenging potential against DPPH and OH radical at IC_{50} values of 22.3 and 1.2 $\mu {\rm M}$ [41]. Garciduol A (1.36), isolated from the stem bark of G. dulcis, was found to be a trioxygenated xanthone connected with 2,4,6-trihydroxybenzophenone [33]. The isolation of schomburgkixanthone (1.37) from G. schomburgkiana twigs and garmoxanthone (1.38) and garcinoxanthone B (1.39) from the bark and pericarps of G. mangostana revealed the existence of bixanthone structures, the coupled xanthones via C-C or C-O bond formation (Figure 1.8) [42-44]. Compound 1.38 significantly inhibited the growth of MRSA ATCC 43300 and CGMCC 1.12409 strains with MIC values of 3.9 μ g/mL and was moderately active against *Vibrio* strains with MIC values in the range of $15.6-31.2 \,\mu$ g/mL, whereas **1.39** showed NO production inhibition at IC₅₀ value of 11.3 μ M and suppressed *i*NOS expression in a concentration-dependent curve.



Figure 1.8. Unusual substituted xanthones (1.35–1.39) from Garcinia plants.

1.3.2. Depsidones

This secondary metabolite class contains two benzene rings connected by ester and ether bridges to generate 11*H*-dibenzo[b,e][1,4]-dioxepin-11-one moiety. The compound is found in lichens as the main constituents with common structures containing methyl groups at C-1, C-9, and/or C-6 and oxygenated substituents at C-3 and C-8, based on its biosynthetic occurrence via polyketide pathway. However, a limited number of depsidones are existed in higher plants which are putatively generated from hydroperoxylation reaction of a xanthone carbonyl to form an ester, as described in Figure 1.2 [21, 45]. In *Garcinia* plant, not over than 40 depsidones have been discovered with a typical 1,3,6- or 1,3,8-trioxygenated form, together with other functionalities, such as hydroxy, methoxy, prenyl, and geranyl moieties. Brevipsidone (1.40) is the simple depsidone structure obtained from the stem bark of *G. brevipedicellata* with only hydroxy and methoxy groups attached to the skeleton [46]. A depsidone with unusual methyl groups, polyanthadepsidone (1.41), was

isolated from *G. polyantha* leaves, as methylated depsidones were commonly found in lichen materials (Figure 1.9) [29].



Figure 1.9. Depsidones and its derivatives (1.40-1.51) from Garcinia plants.

Six prenylated depsidones including those with modified prenyl units, garcinisidone (1.42), paucinervin Q (1.43), oliveridepsidones A–C (1.44–1.46), and garcinisidone D (1.47), were reported from four *Garcinia* species [47-50]. From the twigs of *G. parvifolia*, geranylated depsidones parvifolidones A and B (1.48 and 1.49) were well characterized (Figure 1.9) [51]. Biological activity studies demonstrated that

1.43 was cytotoxic against HL-60 and Caco-2 cancer cells with IC₅₀ values of 3.1 and 6.8 μ M, while **1.47** was found to inhibit Eipstein—Barr virus early antigen (EPV-EA) activation induced by TPA with IC₅₀ value of 360 mol ratio/TPA without significant cytotoxicity against Raji cells, which was better than positive control β -carotene.

Pucinervin B (1.50), isolated from the stems of *G. paucinervis*, was identified as a pseudodepsione with two aromatic rings connected by ether linkage [50]. In addition, depside 2-O-(3,4-dihydroxybenzoyl)-2,4,6-trihydroxypehnylacetic acid (1.51) was isolated from the pericarps of *G. mangostana* (Figure 1.9) [52]. This type of compound is considered the precursor of depsidones, characterized by two or more phenolic rings linked by an ester bond. Unfortunately, these two substances were reported to have no biological activities.

1.3.3. Biphenyls

Among secondary metabolites structurally elucidated from the genus *Garcinia*, biphenyl is one of those having small structures with two aromatic rings linked by C-C bond and substituted by the same functional groups as xanthone and depsidone, including hydroxy, methoxy, and prenylated derivatives. Compared to depsidone, this class of compound is widespread to various *Garcinia* plants with more than 50 structures. Phytochemical investigation on the stems of *G. speciosa* resulted the isolation of two simple biphenyls garcibiphenyl C (**1.52**) and garciosine A (**1.53**), together with garciosine B (**1.54**) bearing methylenedioxy unit formed by an oxidative ring closure of hydroxyl phenol group and adjacent methoxy substituent (Figure 1.10) [53]. A biphenyl with an ester unit, 3-methoxy-5-methoxycarbonyl-4-hydroxy-biphenyl (**1.55**), was identified from *G. oligantha* [54].



A series of prenylated biphenyls were obtained from the twigs of *G. bracteata* and the stems of *G. lancilimba*, including bractebiphenyls B (**1.56**) and A (**1.57**) and garcilancibiphenyls A (**1.58**) [55, 56]. Biphenyls featuring furan and pyran units, including oblongifoliagarcinines C (**1.59**) and D (**1.60**), multibiphenyls B (**1.61**) and C (**1.62**), bractebiphenyl C (**1.63**), 2,2-dimethyl-3,5-dihydro-7-(4-hydroxypehnyl)chromane (**1.64**), garciosine C (**1.65**), and 2-isopropenyl-6-methoxy-7-hydroxy-(4-hydroxyphenyl)-dihydrobenzofuran (**1.66**), have been reported from five *Garcinia* species [53, 56-59]. Dibenzofuran paucinervin C (**1.67**) was obtained from the

leaves of *G. paucinervis* [60]. This compound is naturally formed via an intramolecular cyclization of hydroxyl group at either C-2 or C-2' and aromatic carbon at opposite position [61].

Bioactivities study revealed that compounds **1.52–1.54** and **1.65** had anti-HIV-1 effect against $d^{Tat/Rev}$ MC99 virus in 11A2 cells with EC₅₀ values of 14.2–56.3 μ M, without significant cytotoxicity on the cells. Compound **1.55** showed potent cytotoxicity against SHSY5Y, A549, and MCF-7 cancer cells at IC₅₀ values of 4.8–7.1 μ M, while **1.56** and **1.57** had cytotoxic effects against SHSY5Y and MCF-7 cells with IC₅₀ values lower than 10 μ M. Biphenyls **1.63** was cytotoxic against five cancer cells, including NB4, A549, SHSY5Y, PC3, and MCF-7, whereas **1.64** showed cytotoxic effects on NB4, SHSY5Y, and MCF-7 with IC₅₀ values less than 10 μ M. Compounds **1.58**, **1.61**, **1.62**, and **1.66** displayed anti-rotavirus activity on rotavirus infected to MA104 cells with EC₅₀ values of 10.9–17.6 μ M without any cytotoxicity against the cells (CC₅₀>125 μ M).

1.3.4. Benzophenones

Biosynthetically, benzophenone is categorized as a xanthone intermediate composed by two aromatic rings connected via carbonyl linkage, so-called diarylketone. Benzophenones appear as one of major *Garcinia* phytochemicals in two forms, which are basic benzophenone with unmodified aromatic rings (Figure 1.11) and polyprenylated benzophenone (PPB)/benzoylphloroglucinol (PPBP) possessing highly rearranged prenyl or geranyl side chains at the ring A of phloroglucinol backbone (Figure 1.12). The latter form commonly features a bicyclo[3.3.1]nonane-2,4,9-trione moiety or more complex cyclized derivatives attached to the 13,14-dihydroxylated benzoyl unit (B-ring), resulting diverse chemical architectures and many of which showed pharmacological potentials, especially as anticancer agent [24].



Figure 1.11. Simple benzophenones (1.68–1.73) from Garcinia plants.

Maclurin (1.68) and 2,3',4,5'-tetrahydroxy-6-methoxybenzophenone (1.69) are two simple oxygenated benzophenones and garcihombrianone (1.70) is a benzophenone bearing benzoyl unit which is the first report of its appearance in the genus *Garcinia*. These compounds were reported from the leaves and twigs of *G. multiflora* and the roots of *G. hombroniana* [27, 62]. Two prenylated and geranylated benzophenones, garciosone B (1.71) and gakolanone (1.72), were successfully isolated from *G. speciosa* and *G. kola* [53, 63], while garcimangosone D (1.73), a benzophenone glycoside, was identified from the fruit hulls of *G. mangostana* (Figure 1.11) [64].

Four PPBs-class compounds were characterized from *G. multiflora* and *G. indica*, including garciniagifolone A (1.74), garcinielliptone GC (1.75), garcinol (1.76), and symphonone H (1.77) (Figure 1.12). Compound 1.75 is classified to be a type-A PPB derivative with its benzoyl unit at C-1 of the phloroglucinol skeleton and 1.76 is identified as a type-B PPB with benzoyl ring linked to the C-3. Further cyclization of type-A yields a PPB having an adamantyl skeleton as compound 1.74, while oxidative coupling between hydroxyl group at C-2 and aromatic carbon at C-16 in type-B constructs a γ -pyrone ring as compound 1.77 [27, 28].



Figure 1.12. Polyprenylated benzophenones (1.74–1.82) from Garcinia plants.

Two unusual PPBs containing unprecedented imine functionality at C-10, garciyunnanimines A and B (1.78 and 1.79), were identified from the chemical investigation of the whole parts of G. yunnanensis and were the first occurrence of natural PPBs class with such moiety [65]. PPBs with more complex polycyclic system were observed in garcimulin В (1.80)featuring caged а tetracyclo[5.4.1.1^{1,5}.0^{9,13}]tridecane on the phloroglucinol skeleton [66], as well as (-)garmultin A (1.81) composed by two coupled 2,11-dioxatricyclo[4.4.1.0^{3,9}]undecane and tricyclo[$4.3.1.0^{3,7}$]decane cage structures and its precursor, (–)-garmultin C (**1.82**). The isolated compounds were characterized from the leaves and twigs of G. multiflora [67].

All PPBs compounds described above showed anticancer activities, including compounds **1.74**, **1.75**, **1.77**, and **1.80–1.82** which displayed cytotoxic properties against five cancer cells (SMMC-7721, A-549, HL-60, MCF-7, and SW480) with IC₅₀ values ranging from 1.8 to 17.4 μ M, while **1.78** and **1.79** were cytotoxic against Hep G2, A-549, and RPMI-8226 cancer cells with IC₅₀ values of 1.7–10.1 μ M. Compound **1.82** also demonstrated moderate NO production inhibitory effect in LPS activated RAW 264.7 macrophage cells at IC₅₀ value of 15.1 μ M and was capable of inducing apoptosis in human erythroleukemia (HEL) cells and inhibiting the expression of Fli-1 protein, an oncogene which plays an important role in carcinogenesis. Moreover, **1.76** was found to induce apoptosis and cell cycle arrest in leukemia, pancreatic, and lung cancer cells, as well as modulate gene expression leading to inhibition of pancreatic cells growth.

1.3.5. Flavonoids and biflavonoids

Flavonoid, a group of polyphenolic compounds, is widespread in most of fruits and vegetables and has potent antioxidant and anti-inflammatory capacities. As its natural co-occurrence shares the similar building block and biosynthetic pathway with xanthone, this phytonutrient also appears in *Garcinia* plants with several flavonoid substructures (Figure 1.13) [68]. Pinocembrin (**1.83**), a flavanone-class compound, along with its glycoside and 3-hydroxy forms, naringenin-7-rhamnoglucoside (**1.84**) and aromadendrin-8-C- β -D-glucopyranoside (**1.85**), were identified from three *Garcinia* species [64, 69, 70]. The flavone apigenin (**1.86**) and its prenylated derivative 5,7,4'-trihydroxy-6-[3"-methylbut-3"-enyl]-flavone (**1.87**), isolated from *G. mckeaniana* and *G. xanthochymus*, are the elimination products of a flavanone at C-2 and C-3 [71, 72], while hydroxylation of a flavone at C-3 gives a flavonol class, such as kaempferol (**1.88**) and quercetin (**1.89**) [32].


Figure 1.13. Flavonoids (1.83–1.95) from Garcinia plants.

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The flavanol (–)-epicatechin (1.90), an addition form of a flavonol, was reported in the pericarps of *G. mangostana* [73]. A 1,2-aryl shift of flavone initiated by radical rearrangement yields an isoflavone, such as pauciisoflavone A (1.91) and dulcisisoflavone (1.92) which were originally derived from daidzein and genistein and isolated from the stems of *G. paucinervis* and the green fruits of *G. dulcis* [74, 75]. Compound 1.91 was significantly cytotoxic against NB4, SH-SY5Y, and MCF-7 with IC₅₀ values lower than 10 μ M. A chalcone, 4,4'-dihydroxy-2-methoxychalcone (1.93), and a dihydrochalcone, loureirin A (1.94), were reported from the seeds of *G. dulcis* [76]. Chalcone is a key intermediate to construct various flavonoid derivatives, including

pauciaurone A (**1.95**), an aurone molecule with five-membered ring in the C-ring of flavonoid skeleton [75].

It is worth noting that pinocembrin, apigenin, kaempferol, quercetin, epicatechin, daidzein, and genistein are common flavonoids that are exist in many dietary sources, including propolis, honey, soybean, green and white tea, grape and apple fruits, berries, spinach, and broccoli and they are considered as a basic framework to yield various flavonoid structures in genus *Garcinia*. They were also reported to have broad range of therapeutic potentials for cancer therapy, inflammatory treatment, immune booster, and other degenerative diseases treatment, including diabetes and Alzheimer's disease [77].

Enzymatic coupling reaction of two or more flavonoids lead to the formation of a polymer of flavonoid, such as biflavonoid or flavonoid dimer, trimer, and tetramer, wherein the classification depends on the number of flavonoid molecules connected via C-C and/or C-O-C bonds (Figure 1.14). A previous study showed that a biflavonoid fraction containing morelloflavone (1.96), volkensiflavone (1.97), and amentoflavone (1.98) from *G. madruno* extract exhibited potent atheroprotective activity for cardiovascular diseases treatment by lowering the circulation level of cholesterol and lipid peroxidation product [78]. These three flavonoid dimers are commonly isolated from the polar part of many *Garcinia* species in large amounts. Paucinervin K (1.99) is a flavanone-chromone dimer isolated from the leaves of *G. paucinervis* [79]. Dulcisbiflavonoid B (1.100), isolated from *G. dulcis* green branches, is a prenylated biflavonoid which is rarely found in nature, while the other two prenylated analogs which are not described in this section, dulcisbiflavonoids A and C, were also isolated from the same species [80].



Figure 1.14. Flavonoid dimers and trimers (1.96–1.104) from *Garcinia* plants.

A flavonol dimer and trimer, proanthocyanidin A2 (1.101) and procyanidin C1 (1.102), were reported from the water-soluble fractions of pericarps of *G. mangostana*. The two molecules are obtained from the polymerization of monomer epicatechin and classified to be a group of proanthocyanidin or condensed tannin. Compound 1.101 showed antioxidant and α -glucosidase inhibitory activities with IC₅₀ values of 11.6 and 3.5 μ M, respectively [52, 81]. A biflavonol and flavanone trimer

coupling with a trihydroxyxanthone, garcineflavonol A (**1.103**) and garcineflavanone A (**1.104**), were successfully characterized from the stem bark of *G. atroviridis* and both compounds were linked via ether C-O-C bonds. Compound **1.103** displayed AChE and BChE inhibitions with IC₅₀ values of 14.0 and 14.5 μ g/mL, while **1.04** selectively inhibited AChE enzyme with IC₅₀ value of 28.5 μ g/mL [82].

1.3.6. Terpenoid quinones

Most of terpenoid quinones isolated from *Garcinia* plants are categorized as a group of fat-soluble vitamin E molecules, including tocotrienol (with unsaturated isoprene unit) and tocopherol (with saturated isoprene unit) derivatives. These constituents are naturally occurred from the building block homogentisic acid, a derivative of 4-hydroxypehyl-pyruvic acid, which is subsequently involved in alkylation reaction with phytyl diphosphate containing four isoprene unit and followed by decarboxylation and cyclization to form a δ -tocotrienol, while reduction isoprene double bonds gives a δ -tocopherol. Methylation on both compounds by SAM produces the other three derivatives which are defined based on the number and position of methyl units attached to the aromatic part, including α - (methylated at C-5, C-7, and C-8), β - (methylated at C-5 and C-8), and γ - (methylated at C-7 and C-8) forms [19].

Phytochemical investigation of the stem bark of *G. virgata* led to the purification of δ -tocotrienol (**1.105**), β -tocotrienol 2,8-dimethyl-2-[(3*E*,7*E*)-4,8,12-trimethyldeca-3,7,11-trienyl]-5-formylchroman-6-ol (**1.106**), and γ -tocrotrienol 2,8-dimethyl-2-[(3*E*,7*E*)-4,8,12-trimethyldeca-3,7,11-trienyl]-7-formylchroman-6-ol (**1.107**) (Figure 1.15) [83]. Meanwhile, 1-dotriacontanol (**1.108**), a metabolite isolated from *G. multiflora*, is formed via dihydropyran ring opening of α -tocopherol initiated by enzymatic radical reaction [84].



Figure 1.15. Terpenoid quinone derivatives (1.105–1.113) from *Garcinia* plants.

Coupling reaction of two terpenoid quinones resulted the formation of various dimer products, including δ , δ -bi-O-amplexichromanol (1.109) and δ , γ -biamplexichromanol (1.110) with ether and C-C linkages on the chroman moieties, while linkage of two δ -tocotrienol derivatives via ester and peroxyl ester bonds at the side chains gives δ , δ -biamplexichromanoate A (1.111) and δ , δ -amplexichromanol peroxide (1.112). These four tocotrienol dimers were reported from the stem bark of *G. amplexicaulis*. In addition, a C-C bond formation of two δ -tocotrienol derivatives at methyl group of C-8 generates nigrolineaquinone B (1.113) and its putative

biosynthesis pathway has been proposed by A. Raksat *et. al.*, 2019. Compounds **1.109–1.112** were significantly active to inhibit lipid peroxidation enzyme with IC₅₀ values ranging from 1.7 to 9.9 μ M [85].

1.3.7. Phloroglucinols

From acetate biosynthesis pathway, many polyketide-derived natural products are formed, including benzenetriols, well-known as phloroglucinols, which are constituted from a series of coupled acetyl CoA/malonyl CoA units. 5,7-dihydroxy-2,6,8-methylchromone (1.114) and 2,3-dihydrochromone derivative (1.115), isolated from the pericarps of *G. mangostana*, are two examples of phloroglucinol compounds derived from five and six acetyl CoA/malonyl CoA precursors which are involved in intramolecular cyclization and enolization to form a phloroglucinol skeleton and second cyclization to generate a γ -pyrone moiety. In addition, a chromone glycoside, eucryphin (1.116), and a chromone dimer, paucinervin R (1.117), were obtained from *G. mangostana* and *G. paucinervis* (Figure 1.16) [52, 76, 86].

Three metabolites 4-geranyl-2-(2'-methylpropionyl)-phloroglucinol (1.118), dauphinol F (1.119) and dauphinol C (1.120) isolated from the roots of *G. dauphinensis* are phloroglucinol derivatives containing geranyl side chains, as well as isobutyryl and 2-metylbutyryl units derived from precursor L-valine and isoleucine amino acids. An intramolecular cyclization of geranyl moiety generates dauphinol A (1.121), 3'-methylhyperjovinol B (1.122), empetrikarinol B (1.123), and empetrifranzinan C (1.124) [87].

Garcicowin A (1.125) from *G. cowa* and garsubelone B (1.126) and garsubelone A (1.127) from *G. subelliptica* represent polyprenylated phloroglucinols. Compound 1.127 is the first report of a phloroglucinol constituent possessing an unprecedented 6/6/6/6/6/6 heptacyclic ring system derived from its monomer 1.126. Cytotoxic evaluation showed that compounds 1.118–1.121 and 1.123

inhibited the growth of A2780 ovarian cancer cells with IC₅₀ values of 4.5–16.4 μ M, while **1.126** was cytotoxic against HeLa and HepG2 cancer cells with IC₅₀ values of 6.0 and 7.3 μ M, respectively. In addition, metabolites **1.118–1.122** displayed significant anti-plasmodial activity against *P. falciparum* with IC₅₀ values of 4.7–14.1 μ M [88, 89].



Figure 1.16. Phloroglucinol derivatives (1.114–1.127) from Garcinia plants.

1.3.8. Triterpenoids

Triterpenoid, a mevalonate-derived compound with 30 carbon atoms on the skeleton, is one of major non-phenolic substances that can be found in *Garcinia*

species. Two farnesyl pyrophosphate (FPP) containing C₁₅ building block each incorporate through a series of enzymatic reaction, including electrophilic addition, proton and carbon rearrangements, and hydration to form squalene intermediate. This precursor is subsequently cyclized to generate various triterpenoid structures which mostly feature tetracyclic and pentacyclic ring system [19].



Figure 1.17. Tetracyclic triterpenoids (1.128–1.138) from Garcinia plants.

Lanosta-8,25-en-3 β -ol (1.128), garcinielliptones Q (1.129), and garcihombronane D (1.130), isolated from three *Garcinia* species [90-92], are classified to be a lanostane-type structure which is considered to be a basic parent for other triterpenoid derivatives. The formation of a methylene bridge at C-9 and C-10 gives a cycloartane as in 23-acetoxy-mangiferonic acid (1.131) and mangiferolic acid (1.132)

[93]. An euphane triterpene from *G. eugenifolia* roots, euphadienol (**1.133**), differs from lanostane only in the orientation of methyls C-18 and C-30 [94], while the migration of methyl groups at C-13 and C-14 of lanostane generates a friedolanostane structure, such as (22Z,24E)-9 α -hydroxy-3-oxo-17,13-friedolanosta-12,22,24-trien-26-oic acid (**1.134**), (22Z,24E)-3-oxo-17,14-friedolanosta-8,14,22,24tetraen-26-oic acid (**1.135**), and (22Z,24E)-9 α -hydroxy-3-oxo-13 α ,30-cyclo-17,13friedolanosta-22,24-dien-26-oic acid (**1.136**) from the bark of *G. celebica* [47]. Steroids β -sitosterol (**1.137**) and stigmasterol (**1.138**) are the most common phytochemicals found not only in genus *Garcinia*, but also in other higher plants as modified tetracyclic triterpenoids lacking methyl units at C-4 and C-14 (Figure 1.17).



Figure 1.18. Pentacyclic triterpenoids (1.139–1.143) from Garcinia plants.

Pentacyclic triterpenoid (Figure 1.18) consists five ring system with the new ring E formed via cyclization of the side chain at ring D of tetracyclic triterpenes. Betulin (1.139), betulinic acid (1.140), β -amyrin (1.141), messagenic acid (1.142), and 2β -hydroxy-3 α -O-caffeoyltaraxar-14-en-28-oic acid (1.143) are the examples of pentacyclic lupane and oleanane triterpenoids. The migration of one of methyl groups at C-4 to C-5 and C-20 to C-19 allows the formation of friedelane and ursane types, respectively, as in endodesmiadiol (1.144), canophyllol (1.145), canophyllal (1.146) and α -amyrin (1.147) [87, 95-97].

Cycloartane **1.131** displayed melanin deposition inhibition with IC₅₀ values of 19.4 μ M by downregulated the expression of tyrosinase gene. The cholinesterase inhibition of compounds **1.140** and **1.143** was reported against AChE and BChE enzymes with IC₅₀ values in the range of 10.6–24.2 μ M, while **1.139** selectively inhibited AChE activity with IC₅₀ value of 28.5 μ M. Three friedelanes **1.144–1.146** showed anti-plasmodial activity on *P. folciparum* W2 strain with IC₅₀ values of 13.0– 18.2 μ M. The combination of **1.141** with cisplatin was capable to induce apoptosis and cell cycle arrest, mediate ROS formation in NTUB1 human bladder carcinoma cells, and reduce the side effect and drug resistance of cisplatin. Oleanane glycoside **1.142** showed apoptosis induction of HL-60 human leukemia cell line at a concentration of 4.0 μ M and caused the cell cycle arrest at sub-G1 stage.

1.3.9. Miscellaneous compounds

(-)-Hydroxycitric acid (1.144), a citric acid from *G. cambogia* fruits, was found to block fat accumulation and this substance have been marketed as a weight loss supplement (Figure 1.19) [18]. Two phenolic glycosides, (2*R*)-1-*O*-4-hydroxy-benzoyl-3-*O*- α -d-glucuronosyl glycerol (1.145) and garcinophenylpropanic acid (1.146) and chlorogenic acid (1.147) were obtained from three *Garcinia* species [52, 98, 99]. Angelicoin B (1.148) and 8-hydroxy-6-methoxy-3-pentylisocoumarin (1.149) isolated in the twigs of *G. xanthochymus* the stem bark of *G. dulcis* are classified as isocoumarin derivatives which are constructed through acetate pathway [33, 100]. In some cases, isocoumarin metabolites bearing a phenyl ring at C-3, instead of alkyl chains, were recorded and their biosynthesis is proposed to be similar with that of flavonoid. This class of compound is abundant in microbes, such as fungi, bacteria, and lichens, while it is limited in some higher plant families, including Leguminocease, Asteraceae, Moraceae, and Umbelliferae [101].



Figure 1.19. Miscellaneous compounds (1.144–1.156) from *Garcinia* plants.

A benzofuran glycoside, 2R,3R-2,3-dihydro-2-(4'-hydroxy-3'-methoxyphenyl)-3-(glucosyloxymethyl)-7-methoxy-benzofuran-5-propanol (**1.150**) was reported from the pericarps of *G. mangostana* [81]. Anthraquinones vismiaquinone A (**1.151**) and 3geranylemodin (**1.152**) were isolated from the bark of *G. schomburgkiana* and they shared the same biosynthesis pathway with isocoumarin and phloroglucinol [102]. Two diterpenes, cembrene A (**1.153**) and 2-cyclohexene- γ , η ,2,6,6-pentamethyl-1nonanol (**1.154**), and two megastigmane sesquiterpenes, dihydrophaseic acid (**1.155**) and 4-O-sulpho- β -d-glucopyranosyl abscisate (**1.156**), were identified from the leaves of *G. paucinervis* and the pericarps of *G. mangostana*, respectively [52, 60, 81].

1.4. Cytotoxic evaluation of natural products for preliminary anticancer screening

The MTT assay is one of the most widely used biological tests for preliminary screening of both natural products and synthetic compounds as an anticancer drug candidate. The colorimetric based assay is readily performed on a wide range of cell lines and measure the number of living cells. In this assay, NADH or NADPH produced by mitochondrial dehydrogenase enzymes inside the living cells reduce the tetrazolium salt (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) into a colored formazan product at 37°C. The amount of formazan formed is directly proportional to the number of viable cells present in the culture (Figure 1.20). IC₅₀ value is the concentration of a tested sample that is required to cause the death of 50% of the cells. This value is commonly used to define the degree of cytotoxicity of the substance. The lower the value, the more cytotoxic is the anticancer drug candidate [103, 104].



Figure 1.20. Formazan formation through NADH-catalyzed MTT reduction in live cells.

1.5. Anti-inflammatory evaluation of natural products on COX enzymes inhibition

Several evidences were reported that severe inflammation can induce apoptosis inhibition as well as increase angiogenesis, genome damage, and cellular proliferation which are typical in carcinogenesis. During the chronic symptom, proinflammatory molecules such as cytokines, reactive oxygen species (ROS), and inducible nitric oxide synthase (*i*NOS) are upregulated causing the exponential growth of malignant cells [105]. The use of nonsteroidal anti-inflammatory drugs (NSAIDs) in inflammation and cancer treatment have been reported since significant indications were found that NSAIDs can decrease primary and recurrent cancer incidence. The long term-administration of a NSAID, aspirin, may lower the incidence of esophageal, colorectal, lung, breast, and bladder cancers and the use of other NSAIDs, such as sulindac, ibuprofen, and piroxicam, decreases the risk of existing cancer development [106]. Therefore, eliminating severe inflammation is one of strategies in cancer prevention and the cytokine molecules involved in the process can serve as biomarker for therapy decision.

In this dissertation, we focus on searching anti-inflammatory agent from natural products to inhibit COX enzymes activity. Cyclooxygenase (COX), also known as prostaglandin-endoperoxide synthase (PGHS), is key enzyme comprising two main isoforms, which are COX-1, activated and expressed with constant level by physiological and pathological stimuli, and COX-2, a highly inducible enzyme by pro-inflammatory stimuli [107]. The COX enzymes catalyze the biotransformation of arachidonic acid (AA), a lipid-derived substance released as an intracellular messenger when body tissues get injured or infected, to prostaglandins (PGs) which are responsible as mediators to inflammation reactions in damaged tissues (Figure 1.21) [108]. Nevertheless, uncontrolled biosynthesis of the PG metabolites may also initiate cancer cell development. For example, COX derived prostaglandin E₂ (PGE₂),

a key mediator in acute inflammation, can induce cancer cells growth by binding its receptor and trigger the signal pathway controlling cell proliferation, migration, apoptosis, and angiogenesis and COX-2 expression was found being elevated in many human cancers. Advanced research in regulation of COX enzymes have been conducted to inhibit the enzyme activities (Figure 1.21) [105, 109].



Figure 1.21. Prostaglandins production via arachidonic acid metabolism and a strategy in COX activity inhibition.

The COX inhibition assay is commonly used to screen bioactive compounds from nature or synthetic products as anti-inflammatory agents. The reaction in the assay starts by adding arachidonic acid to the mixtures of COX enzyme, sample, and other reagents. The concentration of PGE₂ generated from the AA transformation by COX enzymes is quantified using prostaglandin E₂ (PGE₂) EIA kit, a competitive immunoassay for quantitative determination of PGE₂ in biological fluids. PGE₂ in the sample mixture is bound by either PGE₂ monoclonal antibody or alkaline phosphatase enzyme in competitive manner [110]. A chromogenic *p*NPP substrate is added to react with conjugated enzyme (alkaline phosphatase—PGE₂) during second incubation. The absorbance of final reaction mixture is measured at 405 nm using UV/Vis spectrophotometer. Sample with high PGE₂ antigen concentration results in lower signal intensity since less substrate react with bound enzyme (blue color) to produce *p*-nitrophenol (yellow color). This implies that the intensity of the yellow color is inversely proportional to the PGE_2 concentration in the standard or samples (Figure 1.22) [111, 112].



Figure 1.22. Reaction between *p*NPP and alkaline phosphatase enzyme to produce *p*-nitrophenol.

1.6. Objectives of the research

In agreement with the above data, exploring phytochemicals from *Garcinia* plants is of high importance to discover and identify novel compounds with promising biological activities, particularly as anticancer candidates. In this work, three *Garcinia* species collected from Indonesia, including *G. cylindrocarpa*, *G. picrorhiza*, and *G. tetrandra*, will be studied since there have been a few reports conducted in terms of their secondary metabolite isolation, with the main objectives of research as follows:

- 1. To isolate the chemical constituents from the stems of *Garcinia cylindrocarpa* and the stem bark of *Garcinia picrorhiza* and *Garcinia tetrandra*;
- To elucidate the chemical structure of isolated compounds by spectroscopic analysis, including UV-Vis, FT-IR, 1D (¹H and ¹³C) and 2D (COSY, HSQC, and HMBC) NMR, and HRMS;
- 3. To evaluate the cytotoxic properties of the isolated compounds against five human cancer cells (KB, HeLa S-3, HT-29, MCF-7, and Hep G2) and antiinflammatory activity towards COX-1 and COX-2 enzymes inhibition.

Chapter II

Phytochemical and biological investigation of the stems of *Garcinia* cylindrocarpa

2.1. Botanical and chemical aspects of G. cylindrocarpa



Figure 2.1. The whole plant, leaves, and fruit of Garcinia cylindrocarpa.

Family	: Clusiaceae
Genus	: Garcinia
Species	: Garcinia cylindrocarpa
Common name	: Kogbirat
Local name	: Kogbirat (in Maluku Island)

Garcinia cylindrocarpa Kosterm is a woody plant and native to Maluku Islands, Indonesia (Figure 2.1). In its place of origin, the species is found in lowland tropical rainforest at altitude of 100 m. It is a medium to large evergreen tree and reaching heights of 15–20 meter with a girth of 20–30 cm. Its leaves are broadly elliptic, and its fruit is turning pink when ripe, which contain 5–6 seeds inside. The ripe fruit is edible but acidic. This species, locally named as Kogbirat, has been used as a traditional medicine for fever remedy [13, 15]. The first report on the chemical constituents of *G. cylindrocarpa* revealed the presence of three pyranoxanthones,

cylindroxanthones A–C (2.1–2.3), from the methanol extract of its stem bark (Figure 2.2). The cytotoxic activity of three compounds was evaluated against five human cancer cell lines (KB, HeLa S-3, HT-29, MCF-7, and Hep G2). The results showed that cylindroxanthone A (2.1) exhibited a potent cytotoxicity against KB cells with IC_{50} value of 2.36 μ M, while the others showed moderate to inactive against the five cancer cells [13].



Figure 2.2. Cylindroxanthones A–C (2.1–2.3) from the stem bark of G. cylindrocarpa.

As a continuation of our research interest to explore bioactive compounds from *G. cylindrocarpa*, we describe herein the isolation and structure elucidation of four new xanthones, cylindroxanthones D–G (**GC1–GC4**), and two new biphenyls, cylindrobiphenyls A and B (**GC5** and **GC6**), along with 28 known compounds from the stems of this plant. The cytotoxic evaluation of the isolated compounds against five human cancer cell lines were also reported.

2.2. Experimental

2.2.1. General Experiment Procedures

Melting points were determined by Fischer-Johns melting point apparatus. Optical rotations were measured on a Jasco (Oklahoma City, OK, USA) P-1010 polarimeter. IR data were obtained using a Nicolet (Thermo Scientific, Waltham, MA, USA) 6700 FT-IR spectrometer using KBr discs. UV—visible absorption spectra were obtained on a Shimadzu (Kyoto, Japan) UV-2550 UV—vis spectrometer. NMR spectra were recorded on a Bruker 400 AVANCE spectrometer (400 MHz for ¹H and 100 MHz for ¹³C) in CDCl₃ and acetone- d_6 . HRESIMS spectra were recorded using a Bruker MICROTOF model mass spectrometer. Silica gel 60G (Merck), silica gel 70–230 mesh (Merck), and Sephadex LH-20 (GE Chemical Corporation) were used for column chromatography. Radial chromatography (Chromatotron model 7924 T, Harrison Research, Palo Alto, California, USA) was carried out on silica gel 60 GF₂₅₄ containing gypsum (Merck). For TLC analysis, precoated silica gel 60 GF₂₅₄ (0.25 mm; Merck) was used.

2.2.2. Plant Material

The stems of *Garcinia cylindrocarpa* were collected from Saumlaki Forest, Southeast West Maluku Islands, Indonesia. The plant was identified by Mrs. Rismita Sari (a botanist at Bogor Botanical Garden, Indonesia). A voucher specimen (No. 630) was deposited at the Herbarium Bogoriense, Bogor Botanical Garden, Indonesia.

2.2.3. Extraction and Isolation

The air-dried stems of *G. cylindrocarpa* (3.0 kg) were ground into powder and extracted by maceration at room temperature with MeOH (3 x 15 L) for three days. The solvent was evaporated under reduced pressure to obtain a residue (31.4 g). The crude extract was then suspended in distilled H₂O and partitioned with CH₂Cl₂ and EtOAc to afford a CH₂Cl₂ fraction (6.04 g), an EtOAc fraction (9.23 g), and the remaining aqueous solution. The CH₂Cl₂-soluble fraction was subjected to VLC on silica gel (125.0 g) using a gradient of hexanes:EtOAc (90:10–0:100) to obtain nine fractions (D1–D9). Fraction D2 (488.8 mg) was chromatographed on Sephadex LH-20 column (50.0 g) eluted with CH₂Cl₂:MeOH (1:1, v/v) to yield subfractions D2.1–D2.3. Subfraction D2.1 (93.2 mg) was separated by repeated Sephadex LH-20 CC (50.0 g) using CH₂Cl₂:MeOH (1:1, v/v) to afford compounds **GC4** (4.3 mg) and **GC7** (4.2 mg).

Compound **GC8** (16.7 mg) was yielded from subfraction D2.2 (70.4 mg) by separation using radial chromatography (chromatotron) with hexanes:EtOAc (95:5) as eluent. Subfraction D2.3 (112.0 mg) was separated by repeated chromatotron eluted with hexanes:EtOAc (95:5) to afford compounds **GC9** (3.1 mg), **GC10** (2.0 mg), **GC11** (2.6 mg), **GC12** (9.3 mg), **GC13** (1.5 mg), and **GC14** (4.4 mg).

Fraction D3 (270.3 mg) was loaded to Sephadex LH-20 column (50.0 g) eluted with CH₂Cl₂:MeOH (1:1, v/v) to provide three subfractions (D3.1–D3.3). Compound **GC1** (2.7 mg) was obtained by purification of subfraction D3.1 (54.6 mg) using chromatotron with hexanes:CH₂Cl₂ (25:75) as eluent. Fraction D4 (229.6 mg) was chromatographed using Sephadex LH-20 column (50.0 g) with CH₂Cl₂:MeOH (1:1, v/v) as eluent to give four subfractions (D4.1–D4.4). Compounds **GC15** (11.6 mg) and **GC16** (1.4 mg) were yielded from subfraction D4.3 (56.5 mg) using chromatotron eluted with hexanes:EtOAc (85:15). Subfraction D4.4 (29.1 mg) was purified by chromatotron using hexanes:CH₂Cl₂ (25:75) as eluent to obtain compound **GC3** (1.2 mg). Fractionation of fraction D5 (202.1 mg) using Sephadex LH-20 CC (50.0 g) with CH₂Cl₂:MeOH (1:1, v/v) as eluent was performed to obtain subfractions D5.1–D5.4. Compound **GC17** (6.0 mg) was yielded from subfraction D5.1 (67.4 mg) using chromatotron eluted with hexanes:EtOAc (85:15), while compound **GC18** (1.7 mg) was obtained from subfraction D5.2 (42.7 mg) using the same technique with hexanes:acetone (75:25) as eluent.

Fraction D6 (581.0 mg) was subjected to Sephadex LH-20 column (50.0 g) using solvent system of CH₂Cl₂:MeOH (1:1, v/v) to obtain subfractions D6.1–D6.6. Compounds **GC5** (7.2 mg), **GC19** (24.7 mg), and **GC20** (3.6 mg) were successfully afforded from subfraction D6.2 (97.2 mg) using chromatotron with CH₂Cl₂:MeOH (97:3) as eluent. Fractionation of subfraction D6.3 (102.1 mg) was conducted using chromatotron with CH₂Cl₂:MeOH (97:3) as eluent to furnish compounds **GC6** (31.1 mg), **GC21** (1.4 mg), and **GC22** (1.7 mg). A separation technique using chromatotron

was employed to obtain compounds **GC23** (7.2 mg) and **GC24** (11.6 mg) from subfraction D6.4 (45.3 mg) with eluent system of CH_2Cl_2 :MeOH (97:3). Subfraction D6.5 (85.4 mg) was applied to chromatotron with CH_2Cl_2 :MeOH (97:3) as eluent to give compounds **GC25** (14.4 mg) and **GC26** (15.5 mg). Compound **GC27** (14.0 mg) was afforded using chromatotron with hexanes:EtOAc (70:30) as eluent from subfraction D6.6 (41.0 mg).

Fraction D7 (204.3 mg) was separated by Sephadex LH-20 CC (50.0 g) with $CH_2Cl_2:MeOH$ (1:1, v/v) as eluent to obtain five subfractions (D7.1–D7.5). A chromatotron technique using eluent system of $CH_2Cl_2:MeOH$ (97:3) was performed to obtain compounds **GC28** (1.4 mg) and **GC29** (1.7 mg) from subfraction D7.1 (17.2 mg) and compound **GC2** (5.8 mg) from subfraction D7.4 (24.5 mg). The same separation technique was conducted to afford compounds **GC30** (20.4 mg) and **GC31** (7.6 mg) from subfraction D7.2 (80.8 mg) using hexanes:EtOAc (60:40) as eluent. Subfraction D7.3 (56.2 mg) was applied on Sephadex LH-20 CC (25.0 g) with $CH_2Cl_2:MeOH$ (1:1, v/v) as eluent to get compound **GC32** (7.7 mg). Finally, fraction D8 (92.1 mg) was separated using Sephadex LH-20 CC (50.0 g) with $CH_2Cl_2:MeOH$ (1:1, v/v) to obtain compounds **GC33** (10.5 mg) and **GC34** (18.9 mg).

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2.2.4. Cytotoxicity Assay

Compounds **GC1–GC34** were tested to *in vitro* cytotoxic evaluation against KB, HeLa S-3, MCF-7, Hep G2, and HT-29 cancer cell lines using MTT colorimetric method as previously described [113]. Doxorubicin was used as the positive control. The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-tetrazolium bromide (Sigma Chemical Co., USA) was dissolved in saline to make a 5 mg/mL stock solution. Cancer cells (3×103 cells) suspended in 100 µg/wells of MEM medium containing 10% fetal calf serum (FCS, Gibco BRL, Life Technologies, NY, USA) were seeded onto a 96-well culture plate (Costar, Corning Incorporated, NY 14831, USA). After 24 h of pre-

incubation at 37 °C in a humidified atmosphere of 5% CO₂/95% air to allow cellular attachment, various concentrations of test solution (10 μ L/well) were added and these were then incubated for 72 h under the above conditions. At the end of the incubation, 10 μ L of tetrazolium reagent was added into each well followed by further incubation at 37 °C for 4 h. The supernatant was decanted, and DMSO (100 μ L/well) was added to allow formazan solubilization. The optical density (OD) of each well was detected using a Microplate reader at 550 nm and for correction at 595 nm. The 50% inhibition concentration (IC₅₀ value) was determined by curve fitting.

2.3. Results and discussion

The CH₂Cl₂-soluble fraction from the stems of G. cylindrocarpa was subjected chromatographic methods to afford four to various new xanthones, cylindroxanthones D-G (GC1-GC4), and two new biphenyls, cylindrobiphenyls A and B (GC5 and GC6), along with 28 previously described xanthones and biphenyls (GC7-GC34). The structures of the new compounds were elucidated using extensive spectroscopic analyses. The known compounds (Figure 2.3) were identified as trapezifolixanthone (GC7) [114], cylindroxanthone A (GC8) [13], 1,2-dihydro-5hydroxy-10-methoxy-1,1,2-trimethyl-6H-furo[2,3-c]xanthen-6-one (GC9) [115], 6hydroxy-11-methoxy-2,2-dimethyl-3H,7H-pyrano[2,3-c]xanthen-7-one (GC10) [115], 6desoxyjacareubin (GC11) [116], 6-desoxyisojacareubin (GC12) [117], 1,6-dihydroxy-5methoxy-6,6-dimethylpyrano[2',3':2,3]-xanthone (GC13) [118], pancixanthone B (GC14) [119], cylindroxanthone B (GC15) [13], osajaxanthone (GC16) [120], 1,3,5trihydroxy-2-prenylxanthone (GC17) [116], α -mangostin (GC18) [121], garciosine A (GC19) [53], garciocine C (GC20) [53], 2,3-dihydro-4,7-dihydroxy-2(1-hydroxy-1methylethyl)-5*H*-furo[3,2-b]xanthen-5-one (GC21) [122], 1,3,5-trihydroxy-4prenylxanthone (GC22) [117], 2-deprenylrheediaxanthone B (GC23) [123], 1,3,5trihydroxy-4-(3'-hydroxy-3'-methylbutyl)xanthone (**GC24**) [117], 4',5'-dihydro-1,6,7trihydroxy-4',4',5'-trimethyl-furano[2',3':3,4]xanthone (**GC25**) [124], neriifolone C (**GC26**) [125], 1,3,5-trihydroxyxanthone (**GC27**) [126], garcimangosxanthone E (**GC28**) [127], 11-hydroxy-1-isomangostin (**GC29**) [128], garcinone D (**GC30**) [129], cratoxylone (**GC31**) [129], 1,3,7-trihydroxy-2-(3-hydroxy-3-methylbutyl)-9*H*-xanthen-9-one (**GC32**) [130], 1,3,5,6-tetrahydroxyxanthone (**GC33**) [128], and 1,3,6,7-tetrahydroxyxanthone (**GC34**) [126] after comparing their ¹H and ¹³C NMR spectroscopic data with previously published data.

2.3.1. Structural elucidation of compound GC1

Physical and spectroscopic properties of cylindroxanthone D (**GC1**): Yellow needles; mp: 264–266 °C; UV (MeOH) λ_{max} : 328, 260, and 213 nm; IR v_{max} (KBr): 3295, 2929, 1657, 1440, and 1255 cm⁻¹; for ¹H (400 MHz, acetone- d_6) and ¹³C (100 MHz, acetone- d_6) NMR spectroscopic data, see Table 2.1; and HRESIMS m/z 311.0937 [M – H] (calcd. for C₁₈H₁₅O₅, 311.0919).

Compound **GC1** was isolated as yellow needles. The molecular formula was deduced as $C_{18}H_{16}O_5$ based on HRESIMS data (m/z = 311.0937 [M - H], calcd. for $C_{18}H_{15}O_5$, 311.0919). The UV spectrum showed absorption bands at λ_{max} 328, 260, and 213 nm, which were characteristic absorbances of a xanthone chromophore. The IR spectrum exhibited strong bands at 3295 and 1657 cm⁻¹, assigned the presence of hydroxy group and conjugated carbonyl group [117]. The ¹H NMR spectrum (Table 2.1) exhibited resonances of two *gem*-dimethyl protons at δ_H 1.39 (each 3H, s, H-4' and H-5'), two sets of methylene protons at δ_H 1.93 (2H, t, J = 6.8 Hz, H-2') and 2.94 (2H, t, J = 6.8 Hz, H-1'), one aromatic proton at δ_H 6.15 (1H, s, H-2), and an intramolecular hydrogen-bonded hydroxy group at δ_H 12.73 (1H, s, 1-OH). Moreover, a characteristic spin system of a 1,2,3-trisubsituted benzene moiety in ring B were observed at δ_H 7.28 (1H, t, J = 8.0 Hz, H-7), 7.38 (1H, d, J = 8.0 Hz, H-6) and 7.69 (1H,

d, J = 8.0 Hz, H-8), which was further corroborated by COSY correlations of H-6/H-7 and H-7/H-8. The ¹³C NMR and HSQC experiments displayed 18 carbon signals, including two methyls at $\delta_{\rm C}$ 27.0 (C-4') and 27.0 (C-5'), two methylenes at $\delta_{\rm C}$ 17.0 (C-1') and 32.4 (C-2'), four methines at $\delta_{\rm C}$ 99.9 (C-2), 116.6 (C-8), 121.8 (C-6), and 125.0 (C-7), nine quaternary carbons at $\delta_{\rm C}$ 77.3 (C-3'), 101.4 (C-4), 104.2 (C-9a), 122.5 (C-8a), 146.5 (C-10a), 147.2 (C-5), 155.6 (C-4a), 162.0 (C-1), and 162.7 (C-3), and one carbonyl carbon at $\delta_{\rm C}$ 181.9 (C-9). The ¹H and ¹³C NMR data of **GC1** were closely related to 6desoxyisojacareubin (**GC12**), except for the double bond of a pyran moiety in **GC12** was reduced to be dihydropyran unit in **GC1**. The COSY correlation of H-1'/H-2' and the HMBC cross-peaks of H-1' with C-3 ($\delta_{\rm C}$ 162.7), C-4 ($\delta_{\rm C}$ 101.4), C-4a ($\delta_{\rm C}$ 155.6), C-2' ($\delta_{\rm C}$ 32.4), and C-3' ($\delta_{\rm C}$ 77.3), H-2' with C-4 and C-1' ($\delta_{\rm C}$ 17.0), and H-4' and H-5' with C-2' and C-3' confirmed the presence of this unit which was fused at C-3 and C-4 with an ether linkage at C-3 (Figure 2.4). Consequently, **GC1** was a dihydro derivative of **GC12**.

2.3.2. Structural elucidation of compound GC2

Physical and spectroscopic properties of cylindroxanthone E (**GC2**): Yellow powder; $[\alpha]_D^{20}$ + 8.3 (c 0.50, MeOH); mp: 276–278 °C; UV (MeOH) λ_{max} : 316, 252, and 206 nm; IR v_{max} (KBr): 3256, 2900, 1630, 1454, and 1250 cm⁻¹; for ¹H (400 MHz, acetone- d_6) and ¹³C (100 MHz, acetone- d_6) NMR spectroscopic data, see Table 2.1; and HRESIMS m/z 327.0895 [M – H] (calcd. for C₁₈H₁₅O₆, 327.0869).

Compound **GC2** was isolated as a yellow powder with a molecular formula of $C_{18}H_{16}O_6$, as suggested by HRESIMS data ($m/z = 327.0895 [M - H]^{-}$, calcd. for $C_{18}H_{15}O_6$, 327.0869). The ¹H NMR data of **GC2** were closely related to **GC1**, except for the replacement of the $-CH_2-CH_2-$ unit in the dihydropyran ring of **GC1** with a $-CH(OH)CH_2-$ moiety in **GC2** as deduced from the resonances of a set of methylene protons at δ_H 2.87 (1H, dd, J = 20.0, 11.2 Hz) and 3.18 (1H, dd, J = 20.0, 5.2 Hz), an oxygenated methine proton at δ_H 3.92 (1H, dd, J = 11.2, 5.2 Hz), and a hydroxy

proton at $\delta_{\rm H}$ 4.48 (1H, d, J = 5.2 Hz) (Table 2.1). The position of the hydroxyl methine proton at C-2' was assigned by HMBC correlations of H-1' to C-2' ($\delta_{\rm C}$ 69.0) and C-3' ($\delta_{\rm C}$ 80.1) and H-4' at $\delta_{\rm H}$ 1.41 (3H, s) and H-5' at $\delta_{\rm H}$ 1.35 (3H, s) to C-2' and C-3' (Figure 2.4). The Mosher's method was applied to confirm the absolute configuration at C-2' by treating **2** with (*S*)-MTPACl and (*R*)-MTPACl [131]. Unfortunately, the NMR spectral data of the MTPA esters of **GC2** showed a mixture of the diastereomers. Therefore, **GC2** was a 2'-hydroxy derivative of **GC1**.

2.3.3. Structural elucidation of compound GC3

Physical and spectroscopic properties of cylindroxanthone F (**GC3**): Yellow powder; UV (MeOH) λ_{max} : 333, 280, and 238 nm; IR v_{max} (KBr): 3280, 2910, 1655, 1438, and 1263 cm⁻¹; for ¹H (400 MHz, acetone- d_6) and ¹³C (100 MHz, acetone- d_6) NMR spectroscopic data, see Table 2.1; and HRESIMS m/z 267.0305 [M – H]⁻ (calcd. for C₁₅H₇O₅, 267.0293).

Compound **GC3** was isolated as a yellow powder with a molecular formula $C_{15}H_8O_5$ determined by HRESIMS ($m/z = 267.0305 [M - H]^-$, calcd. for $C_{15}H_7O_5$, 267.0293). The ¹H and ¹³C NMR spectroscopic data were closely related to the structure of **GC1**, except that **GC3** possessed characteristic signals of a furan ring instead of dihydropyran moiety in **GC3**. In this respect, the 1D NMR (Table 2.1) and HSQC experiments showed resonances of two *cis*-olefinic protons at δ_H 7.38 (d, 1H, J = 1.6 Hz, H-2)/ δ_C 105.3 (C-2') and 7.87 (d, 1H, J = 1.6 Hz, H-1)/ δ_C 145.9 (C-1') [132]. The presence of this furan ring attached at C-3 and C-4 of ring A was further supported by COSY correlation of H-1'/H-2' and HMBC cross-peaks of H-1' with C-3 (δ_C 161.7), C-4 (δ_C 109.9), and C-2' and H-2' with C-3 and C-4 (Figure 2.4).

2.3.4. Structural elucidation of compound GC4

Physical and spectroscopic properties of cylindroxanthone G (**GC4**): Yellow needles; mp: 135–136 °C; UV (MeOH) λ_{max} : 344, 281 and 245 nm; IR v_{max} (KBr): 3327, 2978, 2929, 1640, 1602, 1455 and 1196 cm⁻¹; for ¹H (400 MHz, CDCl₃) and ¹³C NMR (100 MHz, CDCl₃) spectroscopic data, see Table 2.1; and HRESIMS *m/z* 385.1291 [M + H]⁺ (calcd. for C₂₁H₂₁O₇, 385.1287).

Compound **GC4** was obtained as yellow needles. A molecular formula of $C_{21}H_{20}O_7$ was suggested by HRESIMS data ($m/z = 385.1291 [M + H]^+$, calcd. for $C_{21}H_{21}O_7$, 385.1287). The UV and IR spectra indicated a similar pattern as those of xanthone core structures [13]. Comparison of the ¹H and ¹³C NMR data of **GC4** with those of **GC8** [13] showed the replacement of the proton signal of a methoxy group in **GC8** with that of an aromatic proton at δ_H 7.40 (1H, s) in **GC4** which was assigned as H-8 based on its HMBC correlations to C-6 (δ_C 150.3), C-7 (δ_C 148.4), C-8a (δ_C 116.1), C-9 (δ_C 180.2), and C-10a (δ_C 147.4) (Figure 2.4). From the above evidences and comparison with literature data, the structure of **GC4** was determined as cylindroxanthone G.

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2.3.5. Structural elucidation of compound GC5

Physical and spectroscopic properties of cylindrobiphenyl A (**GC5**): Pale yellow gum; $[\alpha]_D^{20}$ –17.2 (c 0.70, CHCl₃); UV (MeOH) λ_{max} : 274 and 213 nm; IR v_{max} (KBr): 3359, 1613, 1510, and 1463 cm⁻¹; for ¹H (400 MHz, CDCl₃) and ¹³C (100 MHz, CDCl₃) NMR spectroscopic data, see Table 2.2; and HRESIMS m/z 269.1190 [M – H]⁻ (calcd. for C₁₇H₁₇O₃, 269.1178).

Compound **GC5** was obtained as pale-yellow gum. Its molecular formula was determined as $C_{17}H_{18}O_3$ by HRESIMS measurement at m/z 269.1190 [M – H]⁻ (calcd. for $C_{17}H_{17}O_3$, 269.1178). The UV absorptions at λ_{max} 274 and 213 nm indicated the presence of benzene chromophore [133]. The IR spectrum showed absorption bands

of hydroxy group (3359 cm⁻¹) and aromatic ring (1613 cm⁻¹). Comparison of the ¹H and ¹³C NMR data with those of **GC20** showed that **GC5** differed from **GC20** in the presence of a 2-hydroxy-3,3-dimethylpyran ring instead of a 2-(1,1-dimethyl-1-hydroxymethyl)dihydrofuran unit, which was characterized by resonances of two methyl protons at $\delta_{\rm H}$ 1.33 (3H, s, H-5")/ $\delta_{\rm C}$ 24.7 and 1.39 (3H, s, H-4")/ $\delta_{\rm C}$ 22.4, a set of methylene protons at $\delta_{\rm H}$ 2.75 and 2.95 (each 1H, dd, J = 17.2, 5.2 Hz, H-1")/ $\delta_{\rm C}$ 26.3, one oxygenated methine proton at $\delta_{\rm H}$ 3.86 (1H, t, J = 5.2 Hz, H-2")/ $\delta_{\rm C}$ 69.5 and a quaternary carbon at $\delta_{\rm C}$ 76.9 (Table 2.2). Attempts to determine the absolute configuration at C-2" using the same Mosher's analysis as that of **GC2** were also unsuccessfully achieved due to a racemic mixture of **GC5**.

2.3.6. Structural elucidation of compound GC6

Physical and spectroscopic properties of cylindrobiphenyl B (**GC6**): Pale yellow powder; mp: 101–102 °C; UV (MeOH) λ_{max} : 270, 231, and 206 nm; IR ν_{max} (KBr): 3364, 1636, 1619, 1500, and 1467 cm⁻¹; for ¹H (400 MHz, CDCl₃) and ¹³C (100 MHz, CDCl₃) NMR spectroscopic data, see Table 2.2; and HRESIMS m/z 245.0827 [M – H] (calcd. for C₁₄H₁₃O₄, 245.0814).

Compound **GC6** was obtained as a pale-yellow powder. Its molecular formula was deduced as $C_{14}H_{14}O_4$ by HRESIMS data (m/z = 245.0827 [M - H], calcd. for $C_{14}H_{13}O_4$, 245.0814). The ¹H NMR spectroscopic data of **GC6** were observed to have similar features with those of **GC19**, except the absence of a methoxy proton signal and the replacement of proton signals of 4-subsituted 1-hydroxy-2-methoxyphenyl ring with those of a para-disubstituted benzene ring at δ_H 6.88 (each 1H, d, J = 8.4 Hz, H-3'/H-5') and 7.42 (each 1H, d, J = 8.4 Hz, H-2'/H-6') in **GC6** (Table 2.2). The COSY spectrum showed correlations of aromatic protons H-2'/H-3' and H-5'/H-6' whereas the HMBC spectrum showed cross-peaks of H-2' and H-6' with C-4' (δ_C 155.3) and H-3' and H-5' with C-1' (δ_C 133.9) and C-4' (Figure 2.4). In agreement with the aforementioned data, the structure of **GC6** was unambiguously assigned as cylindrobiphenyl B.



Figure 2.3. Isolated compounds (**GC1–GC34**) from the stems of *Garcinia cylindrocarpa*.

O+/ 11 .T.7								
	GC1		GC2		GC3		GC4	
	$\delta_{ m H}$ (/ in Hz)	$\delta_{ m C}$	$\delta_{ m H}$ (/ in Hz)	$\delta_{ m C}$	$\delta_{ m H}$ (J in Hz)	$\delta_{ m C}$	$\delta_{ m H}$ (J in Hz)	$\delta_{ m C}$
1		162.0		161.9		161.1		157.6
2	6.15 s	9.9	6.17 s	9.6	6.96 s	95.1		104.9
6		162.7	งา	162.2		161.7		160.7
4		101.4	- asr	100.5	1 Allow	109.9	6.43 s	95.4
4a		155.6		155.8		152.4		157.2
10a		146.5	้ จังจัง โมห	146.4		145.9		147.4
5		147.2	131	147.2		147.3		141.5
9	7.38 d (8.0)	121.8	7.38 dd (8.0, 1.2)	121.8	7.45 d (8.0)	122.2		150.3
7	7.28 t (8.0)	125.0	7.29 t (8.0)	125.1	7.38 t (8.0)	125.8		148.4
80	7.69 d (8.0)	116.6	7.69 dd (8.0, 1.2)	116.6	7.79 d (8.0)	116.8	7.40 s	100.5
8a		122.5		122.5		121.5		116.1
6		181.9		181.9		183.1		180.2
9a		104.2		104.4		106.3		104.2
1'	2.94 t (6.8)	17.0	2.87 dd (20.0, 11.2)	26.2	7.87 d (1.6)	145.9	6.73 d (10.0)	115.6
			3.18 dd (20.0, 5.2)					

Table 2.1. ¹H (400 MHz) and ¹³C (100 MHz) spectroscopic data of **GC1–GC3** in acetone- d_6 (in ppm) and **GC4** in CDCl₃ (δ in ppm)

49

ecz			GC3		GC4	
~	Hz) ć	$\delta_{ m C}$	$\delta_{ m H}$ (/ in Hz)	$\delta_{\scriptscriptstyle \mathbb{C}}$	$\delta_{ m H}$ (/ in Hz)	$\delta_{\scriptscriptstyle \mathbb{C}}$
		30.1				78.4
		25.9			1.48 s	28.6
		21.4			1.48 s	28.6
			12.89 s		13.19 s	
)					4.04 s	62.1
					3.96 s	56.4
		8			4.06 s	61.7

Position	GC5 (in CDCl ₃)		GC6 (in CDCl ₃)	
1 0310011	$\delta_{ m H}$ (J in Hz)	$\delta_{\scriptscriptstyle extsf{C}}$	$\delta_{ extsf{H}}$ (J in Hz)	$\delta_{\scriptscriptstyle extsf{C}}$
1		141.3		137.4
2	6.60 s	106.0	6.63 d (1.6)	103.2
3		154.9		152.6
4		105.8		134.9
5		154.2		149.5
6	6.71 s	108.6	6.78 d (1.6)	106.7
1′	2/11	140.7		133.9
2′	7.51 d (8.0)	127.0	7.42 d (8.4)	128.4
3'	7.38 t (8.0)	128.8	6.88 d (8.4)	115.7
4'	7.31 t (8.0)	127.5		155.3
5′	7.38 t (8.0)	128.8	6.88 d (8.4)	115.7
6'	7.51 d (8.0)	127.0	7.42 d (8.4)	128.4
1″	2.75 dd (17.2, 5.2)	26.3		
	2.95 dd (17.2, 5.2)		ວິຍ	
2″	3.86 t (5.2)	69.5		
3"		76.9	ERSTIY	
4″	1.39 s	22.4		
5″	1.33 s	24.7		
5-OH			5.83 brs	
4'-OH			5.04 brs	
3-OMe			3.92 s	56.1
4-OMe			3.93 s	61.2

Table 2.2. ^1H (400 MHz) and ^{13}C (100 MHz) spectroscopic data of compounds <code>GC5</code> and <code>GC6</code>



Figure 2.4. Key COSY and HMBC correlations of GC1-GC6.

2.3.7. Cytotoxic activities of the isolated compounds

Xanthones and biphenyls isolated from plants were well known to exhibit cytotoxic activities against cancer cells and some of which were reported to have a remarkable cytotoxicity [28, 127, 134]. Moreover, our previous works showed that cylindroxanthone A (**GC8**), 6-desoxyjacareubin (**GC11**), and cylindroxanthone B (**GC15**) isolated from two *Garcinia* species displayed moderate to good cytotoxicity against several human cancer cells [13, 21]. Thus, we evaluated the cytotoxicity of compounds **GC1–GC34** against KB and HeLa S-3 cancer cell lines using modified MTT method [113]. Seven compounds (**GC14**, **GC22**, **GC23**, **GC25**, **GC31**, **GC33**, and **GC34**) with IC₅₀ values lower than 30 μ M against these two cancer cells were selected to further investigate their cytotoxicity against MCF-7, Hep G2, and HT-29 cancer cell lines. The results are summarized in Table 2.3. The tested compounds mostly showed moderate to inactive against KB and HeLa S-3 cells. However, a potent cytotoxicity was observed for compound **GC23** against four human cancer cell lines including KB, HeLa S-3, MCF-7, and Hep G2 with IC₅₀ values of 3.41, 5.04, 2.20, and 6.00 μ M, respectively. In addition, compound **GC25** selectively showed good

cytotoxicity against MCF-7 cells with IC₅₀ value of 8.77 μ M and compound **GC31** exhibited good cytotoxicity against HT-29 cells with IC₅₀ value of 9.18 μ M. Structureactivity relationship studies (Figure 2.3 and Table 2.3) suggested that the presence of hydroxy groups at C-5 and C-6 of furanoxanthones might be improved the cytotoxicity, as inferred from the comparison of their cytotoxicity with compounds **GC9**, **GC14**, **GC23**, and **GC25**. Among linear pyranoxanthones (**GC4**, **GC7**, **GC8**, **GC11**, **GC13**, **GC15**, **GC16**, and **GC28**), only compound **GC8** selectively exhibited potent cytotoxicity against KB cell line with IC₅₀ value of 2.36 μ M due to the methoxy groups which were fully substituted at B-ring.

Compound	$IC_{50} (\mu M) \pm SD$					
compound	KB	HeLa S-3	MCF-7	Hep G2	HT-29	
GC1	47.06 ± 0.29	inactive	NT	NT	NT	
GC2	32.28 ± 1.09	50.79 ± 0.71	NT	NT	NT	
GC3	inactive	inactive	NT	NT	NT	
GC4	inactive	inactive	NT	NT	NT	
GC5	inactive	inactive	ห _ก ริทยาลัย	NT	NT	
GC6	inactive	inactive	NT NIVERSI	NT	NT	
GC7	35.20 ± 0.88	38.08 ± 0.04	NT	NT	NT	
GC8	2.36 ± 0.01	75.12 ± 0.81	98.54 ± 6.16	10.41 ± 0.15	25.51 ± 3.26	
GC9	inactive	inactive	NT	NT	NT	
GC10	73.24 ± 0.43	inactive	NT	NT	NT	
GC11	18.56 ± 0.15	19.69 ± 0.46	17.53 ± 0.06	39.35 ± 1.05	69.29 ± 0.79	
GC12	78.62 ± 0.54	inactive	NT	NT	NT	
GC13	inactive	inactive	NT	NT	NT	
GC14	12.13 ± 0.04	20.73 ± 0.56	15.61 ± 0.32	22.84 ± 0.39	inactive	
GC15	57.24 ± 2.56	71.38 ± 0.81	inactive	59.53 ± 0.49	inactive	

Table 2.3. In vitro cytotoxicity of compounds GC1-GC34

Compound	$IC_{50} (\mu M) \pm SD$					
compound	KB	HeLa S-3	MCF-7	Hep G2	HT-29	
GC16	inactive	inactive	NT	NT	NT	
GC17	36.50 ± 0.44	59.28 ± 0.40	NT	NT	NT	
GC18	36.83 ± 0.46	54.94 ± 0.58	NT	NT	NT	
GC19	89.05 ± 3.11	95.25 ± 2.74	NT	NT	NT	
GC20	inactive	inactive	NT	NT	NT	
GC21	36.06 ± 0.78	32.41 ± 1.11	NT	NT	NT	
GC22	22.65 ± 0.08	29.74 ± 1.22	28.59 ± 0.87	50.27 ± 0.47	39.43 ± 1.16	
GC23	3.41 ± 0.08	5.04 ± 0.04	2.20 ± 0.05	6.00 ± 0.07	46.50 ± 0.45	
GC24	39.93 ± 0.44	40.73 ± 0.48	NT	NT	NT	
GC25	13.37 ± 0.17	16.09 ± 0.11	8.77 ± 0.30	11.65 ± 0.06	58.70 ± 1.08	
GC27	73.15 ± 0.14	79.02 ± 0.17	NT	NT	NT	
GC28	48.95 ± 1.57	43.94 ± 0.82	NT	NT	NT	
GC29	inactive	inactive	NT	NT	NT	
GC30	30.11 ± 0.54	23.91 ± 0.98	NT	NT	NT	
GC31	13.43 ± 0.33	18.90 ± 0.37	32.40 ± 1.32	47.28 ± 1.25	9.18 ± 0.52	
GC32	43.63 ± 0.54	44.06 ± 0.32	NT	NT	NT	
GC33	21.69 ± 0.47	29.66 ± 0.46	20.93 ± 0.38	20.66 ± 0.26	inactive	
GC34	11.31 ± 0.34	26.40 ± 0.28	21.77 ± 0.14	99.69 ± 0.36	77.86 ± 0.57	
Doxorubicin	0.02 ± 0.01	0.13 ± 0.01	0.61 ± 0.06	1.07 ± 0.16	0.34 ± 0.07	

Note: $IC_{50} \le 10 \ \mu M$ = good cytotoxicity, $10 \ \mu M < IC_{50} \le 30 \ \mu M$ = moderate cytotoxicity, $30 \ \mu M < IC_{50} \le 100 \ \mu M$ = weak cytotoxicity, $IC_{50} > 100 \ \mu M$ = inactive, NT = not tested.

Chapter III

Phytochemical and biological investigation of the stem bark of Garcinia tetrandra

3.1. Botanical and chemical aspects of G. tetrandra



Figure 3.1. The whole plant, stem, and stem bark of Garcinia tetrandra.

Family	: Clusiaceae
Genus	: Garcinia
Species	: Garcinia tetrandra
Common name	: Kandis Watu
Local name	: Kandis Watu (Sulawesi Island), Laru (East Nusa Tenggara

Province), Mapau (Maluku Island)

Garcinia tetrandra Pierre originally exist in Sulawesi, East Nusa Tenggara, and Maluku Islands and commonly named as Kandis Watu (Figure 3.1). In its natural habitat, it is found in tropical rainforest from 30–600 m altitudes and grows on clayey soil. The species is a medium-sized evergreen tree and can reach heights of 18 meter with a girth up to 30 cm. Its fruiting season starts from December to February and the woods are commonly utilized for building construction materials [15, 135].

Phytochemical studies on the stem bark of this species led to the isolation of several types of secondary metabolites such as xanthones (**3.1–3.3**), polyprenylated benzophenones (**3.4** and **3.5**), and triterpenoids (**3.6–3.8**) (Figure 3.2) and some of them showed antioxidant and antibacterial activities [136].



Figure 3.2. Isolated compounds (3.1–3.8) from the stem bark of *G. tetrandra*.

As part of our studies to explore novel and bioactive compounds from *Garcinia* species [21, 35, 137], a CH_2Cl_2 extract from *G. tetrandra* stem bark was investigated. Herein, the separation, structural characterization, and cytotoxic evaluation of the isolated compounds (**GT1–GT31**) are described.

3.2. Experimental

3.2.1. General Experimental Procedures

The chromatographic materials and instruments information were the same to those previously reported [137, 138] and as described in Chapter II section 2.2.1. The experimental ECD data were recorded on a JASCO J-815 circular dichroism spectropolarimeter and Semi-preparative HPLC utilized a Waters Delta 600 instrument (Waters Corporation) with a Waters 2996 photodiode array detector using an Apollo C₁₈ 5 μ m, 250 x 10.0 mm column.

3.2.2. Plant Material

Garcinia tetrandra stem bark was collected from Baluran National Park, Banyuwangi, Indonesia (7°55′3″ S 114°23′25″ E) in November 2016. The plant material was identified by Mr. Deden Mudiana, and a voucher specimen of this plant (no. XVII.J.11.22) was deposited at Purwodadi Botanical Garden, Indonesia.

3.2.3. Extraction and Isolation

G. tetrandra stem bark (3.0 kg) was macerated with MeOH (each 15 L x 3 days). The filtrate was concentrated by a rotary evaporator to give a residue (71.0 g). The crude extract was suspended in water and partitioned with CH_2Cl_2 and EtOAc to obtain a CH_2Cl_2 and an EtOAc fractions. The dried CH_2Cl_2 fraction (18.6 g) was separated to a quick column chromatography (QCC) separation on silica gel (500.0 g) with a gradient of hexanes-EtOAc (95:5–0:100) to yield seven fractions (A–G). Fraction B (3.5 g) was subjected to a Sephadex LH-20 column (50% $CH_2Cl_2/MeOH$) to afford subfractions B1–B3. Subfraction B2 (1.1 g) was subjected to a Chromatotron using hexanes-acetone (95:5) to give three subfractions (B2.1–B2.3). Subfraction B2.1 (17.2 mg) was separated by a Chromatotron using hexanes-acetone (95:5) to obtain **GT25** (7.4 mg). Subfraction B2.3 (78.4 mg) was separated on a Chromatotron using hexanes-
chloroform (60:40) to yield compound **GT1** (8.1 mg) and a subfraction that was further purified by semi-preparative HPLC using 100% CH₃CN to give **GT2** (3.6 mg, $t_{\rm R}$ = 23.9 min) and **GT3** (4.6 mg, $t_{\rm R}$ = 22.7 min). Subfraction B3 (41.0 mg) was separated on a Chromatotron with hexanes-CH₂Cl₂ (60:40) to obtain **GT26** (8.4 mg) and **GT28** (4.2 mg).

Fraction C (2.2 g) was separated on a Sephadex LH-20 column (50% CH_2Cl_2 /MeOH) to obtain four subfractions (C1–C4). Subfraction C2 (43.0 mg) was then separated on a Chromatotron with hexanes-CH₂Cl₂ (10:90) to afford compounds GT10 (3.2 mg) and GT11 (3.0 mg). Subfraction C3 (80.9 mg) was separated using a Chromatotron with hexanes-CH₂Cl₂ (20:80) to yield GT6 (2.9 mg) and GT23 (4.1 mg). Subfraction C4 (65.5 mg) was subjected to a Sephadex LH-20 column (50% CH₂Cl₂/MeOH) to obtain GT19 (6.6 mg) and GT30 (5.0 mg). Fraction D (2.4 g) was passed through a Sephadex LH-20 column (50% CH₂Cl₂/MeOH) to give three subfractions (D1-D3). Subfraction D2 (67.1 mg) was purified using a Chromatotron by elution with 100% chloroform to afford GT24 (8.9 mg). Subfraction D3 (482.4 mg) was then separated on a Chromatotron using hexanes-chloroform (1:1) to give subfractions D3.1-D3.4. Compounds GT18 (2.8 mg) and GT27 (2.1 mg) were obtained from subfraction D3.2 (18.9 mg) using a Chromatotron with hexanes-chloroform (1:1). Subfraction D3.3 (227.5 mg) was purified using a Chromatotron with hexanes-CH₂Cl₂ (20:80) to afford GT7 (4.1 mg), GT8 (3.8 mg), GT12 (8.4 mg), GT15 (8.5 mg), and GT31 (1.2 mg).

Fraction E (2.1 g) was separated on silica gel CC using CH_2Cl_2 -MeOH (98:2) to obtain subfractions E1—E5. Compound **GT29** (13.1 mg) was purified from subfraction E1 (59.4 mg) by Sephadex LH-20 column using 50% CH_2Cl_2 /MeOH. Compound **GT4** (5.2 mg) was purified from subfraction E2 (15.5 mg) using a Chromatotron with hexanes-acetone (30:70). Subfraction E3 (34.7 mg) was separated on a Chromatotron

with CH_2Cl_2 -MeOH (30:1) to afford **GT5** (3.7 mg) and **GT9** (4.5 mg). Subfraction E4 (100.6 mg) was purified on a Chromatotron using hexanes-EtOAc (65:35) to obtain **GT13** (8.4 mg), **GT14** (2.9 mg), **GT16** (8.3 mg), and **GT17** (14.5 mg). Fraction F (507.1 mg) was subjected to a Sephadex LH-20 column (50% CH_2Cl_2 /MeOH) to obtain four subfractions (F1—F4). Subfraction F3 (15.9 mg) was separated on a Chromatotron using CH_2Cl_2 -MeOH (30:1) to yield **GT21** (6.1 mg). Compounds **GT20** (3.3 mg) and **GT22** (5.7 mg) were afforded from subfraction F4 (31.1 mg) by a Sephadex LH-20 column (50% $CH_2Cl_2/MeOH$).

3.2.4. Cytotoxicity Assay

The isolated compounds (**GT1–GT31**) were tested for their cytotoxicity against Hep G2, HeLa S3, MCF-7, HT-29, and KB cell lines using an MTT colorimetric method with incubation of cells for 72 h [113], as previously described in Chapter II Section 2.2.4.

3.3. Results and discussion

Nine new xanthones (GT1-GT9) and 22 known analogues (GT10-GT31) were isolated from the CH2Cl2 extract of G. tetrandra stem bark through a series of chromatographic techniques (Figure 3.3). Comparison of NMR spectroscopic data with reported data led to the identification of the previously described xanthones as isocowanin (GT10) [139], parvifolixanthone C (GT11) [51], α -mangostin (GT12) [140], garcinone D (GT13) [129], xanthochymone A (GT14) [100], rubraxanthone (GT15) [139]. parvixanthone G (GT16) [36], butyraxanthone D (GT17) [141]. nigrolineaxanthone E (GT18) [142], euxanthone (GT19) [143], gentisein (GT20) [143], 1,3,6,7-tetrahydroxyxanthone (GT21) [126], 1,3,5,6-tetrahydroxyxanthone (GT22) [126], nitidaxanthone (GT23) [118], cylindroxanthone B (GT24) [13], cylindroxanthone A (GT25) [13], 9-hydroxycalabaxanthone (GT26) [144], macluraxanthone (GT27) [145],

3-isomangostin (GT28) [146], 3-isomangostin hydrate (GT29) [146], dihydroosajaxanthone (GT30) [147], and toxyloxanthone B (GT31) [148].



Figure 3.3. Structures of isolated xanthones (**GT1–GT31**) from the stem bark of *G. tetrandra*.

3.3.1. Structural elucidation of compound GT1

Physical and spectroscopic properties of tetrandraxanthone A (**GT1**): Yellow gum; UV (MeOH) λ_{max} (log ε) 316 (4.26), 245 (4.15), 205 (4.19) nm; IR (KBr) v_{max} 3427,

2922, 1635, 1620, 1579, 1462, 1296, 1159 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 545.2922 [M – H] (calcd for C₃₄H₄₁O₆, 545.2903).

The (–)-HRESIMS analysis of tetrandraxanthone A (GT1) indicated a $[M - H]^{-1}$ peak at m/z 545.2922, assigning to the molecular formula $C_{34}H_{42}O_6$. The ¹H NMR spectrum indicated signals for a hydrogen-bonded hydroxy group at $\delta_{\rm H}$ 13.68 (1H, s, OH-1), an aromatic methine proton at $\delta_{\rm H}$ 6.85 (1H, s, H-5), and a methoxy group at $\delta_{\rm H}$ 3.81 (3H, s, OCH₃-7) (Table 3.1). A geranyl unit was identified from the resonances of two olefinic protons at $\delta_{\rm H}$ 5.24 (1H, t, J = 6.8 Hz, H-2") and 5.05 (1H, t, J = 6.4 Hz, H-6"), three methylene protons at $\delta_{\rm H}$ 3.51 (2H, d, J = 6.8 Hz, H-1"), 2.09 (2H, m, H-5"), and 2.03 (2H, m, H-4"), and three methyl groups at $\delta_{\rm H}$ 1.86 (3H, s, H-10"), 1.64 (3H, s, H-8"), and 1.57 (3H, s, H-9"), which was also corroborated by the observed HMBC data (Figure 3.4). The remaining signals of two olefinic protons at $\delta_{\rm H}$ 5.28 (1H, t, J = 6.8 Hz, H-2') and 5.26 (1H, t, J = 6.0 Hz, H-2"'), two methylene protons at $\delta_{\rm H}$ 4.10 (2H, d, J = 6.0 Hz, H-1") and 3.45 (2H, d, J = 6.8 Hz, H-1), and four methyl groups at $\delta_{\rm H}$ 1.84 (3H, s, H-5"), 1.83 (3H, s, H-5'), 1.76 (3H, s, H-4'), and 1.68 (3H, s, H-4") suggested the presence of two prenyl moieties. The 1D NMR data of GT1 indicated that it has the same xanthone skeleton as cowagarcinone A [149], with the major differences being in the positions of geranyl and two prenyl units. The HMBC cross-peaks of H-1' with $\delta_{\rm C}$ 158.7 (C-1), 109.0 (C-2), and 160.1 (C-3) and the deshielded H-1" with $\delta_{\rm C}$ 142.7 (C-7), 137.2 (C-8), and 112.2 (C-8a) confirmed unambiguously the positions of the two prenyl moieties at C-2 and C-8 (Figure 3.4). The attachment of the geranyl unit at C-4 in the A-ring was determined from the cross-peaks of H-1" with $\delta_{\rm C}$ 160.1 (C-3), 104.5 (C-4), and 152.2 (C-4a). The configuration of the $\Delta^{2^{",3^{"}}}$ alkene unit was determined as (2''E) according to ¹³C NMR data comparison with the geranyl unit of previously reported xanthones [150], wherein in GT1, the resonance of the methylene carbon at C-4" was more deshielded ($\Delta\delta_{\rm C}$ +8.0), and that of the methyl carbon at C-10" was more shielded ($\Delta\delta_{\rm C}$ –6.9), than those of garcinianone B (with Z configuration).

Compound **GT1** was named tetrandraxanthone A, as shown in Figure 3.3. It is worth noting that the trivial name "tetrandraxanthone" has been used previously for a compound isolated from the title plant [136].

3.3.2. Structural elucidation of compound GT2

Physical and spectroscopic properties of tetrandraxanthone B (**GT2**): Brown gum; UV (MeOH) λ_{max} (log ε) 329 (4.31), 244 (4.33), 212 (4.16) nm; IR (KBr) v_{max} 3399, 2926, 1651, 1600, 1575, 1464, 1286, 1176 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 475.2161 [M – H]⁻ (calcd for C₂₉H₃₁O₆, 475.2121).

The (–)-HRESIMS analysis of tetrandraxanthone B (**GT2**) indicated a [M – H] peak at m/z 475.2161, assigning to the molecular formula $C_{29}H_{32}O_6$. The ¹H and ¹³C NMR data of **GT2** were nearly identical to those of isocowanin (**GT10**), except that the signals for the prenyl moiety in **GT10** were absent. The 1D NMR data combined with an HSQC experiment indicated signals for *cis*-olefinic protons at δ_{H}/δ_{C} 6.77 (1H, d, J = 10.0 Hz, H-1)/115.3 and 5.57 (1H, d, J = 10.0 Hz, H-2)/127.0, *gem*-dimethyl groups at δ_{H}/δ_{C} 1.47 (6H, s, H-4' and H-5)/28.4, and an oxygenated sp³ carbon at δ_{C} 78.1 (C-3), revealing the presence of a dimethyl pyran unit. The HMBC correlations of H-1' with C-3 (δ_{C} 160.1) and H-2' with C-4 (δ_{C} 100.5) indicated that this unit is fused at its ether linkage C-3 and C-4 in the A-ring (Figure 3.4).

3.3.3. Structural elucidation of compound GT3

Physical and spectroscopic properties of tetrandraxanthone C (**GT3**): Brown gum; $[\alpha]_D^{20}$ +5.8 (c 0.72, CHCl₃); UV (MeOH) λ_{max} (log ε) 321 (4.27), 258 (4.04), 210 (4.11) nm; IR (KBr) v_{max} 3400, 2926, 1649, 1579, 1465, 1278, 1217, 1157 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 475.2134 [M – H]⁻ (calcd for C₂₉H₃₁O₆, 475.2121). The (-)-HRESIMS analysis of tetrandraxanthone C (**GT3**) provided a [M – H] peak at m/z 475.2134, indicating to the molecular formula C₂₉H₃₂O₆. Analysis of the 1D NMR data suggested that **GT3** has a scaffold similar to parvifolixanthone C (**GT11**). The notable differences were in the presence of *cis*-olefinic protons at δ_{H}/δ_{C} 6.81 (1H, d, J = 10.0 Hz, H-1)/115.7 and 5.52 (1H, d, J = 10.0 Hz, H-2)/125.9 and an oxygenated sp³ carbon at δ_{C} 80.6 (C-3') in **GT3** instead of the C-1' methylene, C-2' olefinic methine, and C-3' quaternary sp² carbon signals in **GT11**. The key COSY cross-peaks of H-1/H-2' and H-4/H-5' and HMBC cross-peaks from H-1' to C-3 (δ_{C} 160.5) and C-3', H-2' to C-4 (δ_{C} 100.3) and C-3', H-4' to C-2' and C-5' (δ_{C} 22.8), and H-10' to C-3' and C-4' (δ_{C} 41.8) indicated the geraryl unit at C-4 to be cyclized (Figure 3.4). The experimental ECD spectrum showed no Cotton effects, indicating that **GT3** was obtained as a racemic mixture.

3.3.4. Structural elucidation of compound GT4

Physical and spectroscopic properties of tetrandraxanthone D (**GT4**): Brown gum; UV (MeOH) λ_{max} (log ε) 315 (4.26), 243 (4.34), 209 (4.10) nm; IR (KBr) v_{max} 3392, 2920, 1645, 1612, 1579, 1462, 1429, 1296, 1153 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 495.2425 [M – H] (calcd for C₂₉H₃₅O₇, 495.2383).

The (–)-HRESIMS analysis of tetrandraxanthone D (**GT4**) indicated a [M – H] peak at m/z 495.2425, attributing to the molecular formula C₂₉H₃₆O₇. Comparison of the 1D NMR data of **GT4** with those of **GT11** suggested the principal difference to be the absence of resonances attributed to a prenyl unit in **GT11**. The signals of two methylene groups at δ_{H}/δ_{C} 3.46 (2H, m, H-1")/23.3 and 1.75 (2H, m, H-2")/45.8, two methyl groups at δ_{H}/δ_{C} 1.30 (6H, s, H-4" and H-5")/29.3, and one oxygenated sp³ carbon at δ_{C} 70.6 (C-3") were observed instead. The COSY cross-peak of H-1"/H-2" and key HMBC correlations from H-1" to C-7 (δ_{C} 144.7), C-8 (δ_{C} 140.1), C-8a (δ_{C} 111.9), and C-2", H-2" to C-3", and H-4"/H-5" to C-3" revealed the presence of a 3-hydroxy-3-

methylbut-1-yl fragment at C-8 (Figure 3.4). Thus, **GT4** was assigned as a hydrated derivative of **GT11**.

3.3.5. Structural elucidation of compound GT5

Physical and spectroscopic properties of tetrandraxanthone E (**GT5**): Brown gum; $[\alpha]_D^{20}$ +1.2 (c 0.40, MeOH); UV (MeOH) λ_{max} (log ε) 320 (4.37), 241 (4.12), 208 (4.28) nm; IR (KBr) v_{max} 3395, 2922, 1643, 1616, 1580, 1465, 1292, 1150 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 495.2415 [M – H]⁻ (calcd for C₂₉H₃₅O₇, 495.2383).

The (-)-HRESIMS analysis of tetrandraxanthone E (**GT5**) provided a $[M - H]^{-1}$ peak at m/z 495.2415, indicating to the molecular formula C₂₉H₃₆O₇. The close NMR data similarities between **GT5** and **GT10** were evident, and these compounds differed only in the chemical shifts corresponding to the geranyl side chain at C-8. A methine group at $\delta_{\rm H}$ 5.18 (1H, t, J = 6.8 Hz, H-6") and four methylene groups at $\delta_{\rm H}$ 3.44 (2H, m, H-1"), 2.19 (2H, m, H-5"), 1.77 (2H, m, H-2"), and 1.58 (2H, m, H-4") were observed in the ¹H NMR spectrum, indicating that one of the double bonds in the geranyl unit is hydrated in **GT5**. The COSY correlations of H-1"/H-2" and key HMBC correlations from H-1" to C-7 ($\delta_{\rm C}$ 144.7), C-8 ($\delta_{\rm C}$ 140.2), and C-2" ($\delta_{\rm C}$ 43.7), H-4" with an oxygenated sp³ carbon C-3" ($\delta_{\rm C}$ 72.5), and H-10" with C-2", C-3", and C-4" ($\delta_{\rm C}$ 42.8) suggested that the double bond at position C-2" was also hydrated (Figure 3.4). The ECD spectrum showed no Cotton effects, indicating that **GT5** was obtained to be racemic.

3.3.6. Structural elucidation of compound GT6

Physical and spectroscopic properties of tetrandraxanthone F (**GT6**): Brown gum; UV (MeOH) λ_{max} (log ε) 316 (4.22), 244 (4.14), 210 (4.13) nm; IR (KBr) v_{max} 3423,

2924, 1641, 1610, 1583, 1458, 1280, 1162 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (-)-HRESIMS m/z 409.1682 [M - H] (calcd for C₂₄H₂₅O₆, 409.1651).

The (–)-HRESIMS analysis of tetrandraxanthone F (**GT6**) gave a $[M - H]^{-}$ peak at m/z 409.1682, assigning to the molecular formula $C_{24}H_{26}O_6$. The NMR spectroscopic data confirmed that **GT6** is a positional isomer of α -mangostin (**GT12**), in which the prenyl moiety of **GT6** is attached at C-4. Confirmation for this hypothesis was obtained from the HMBC spectrum, wherein the methylene proton at $\delta_{\rm H}$ 3.52 (2H, d, J = 6.4 Hz, H-1') correlated with a quaternary carbon at $\delta_{\rm C}$ 104.4 (C-4) and two oxygenated sp² carbons at $\delta_{\rm C}$ 161.1 (C-3) and 154.0 (C-4a) (Figure 3.4).

3.3.7. Structural elucidation of compound GT7

Physical and spectroscopic properties of tetrandraxanthone G (**GT7**): Brown gum; $[\alpha]_D^{20}$ +3.2 (c 0.35, MeOH); UV (MeOH) λ_{max} (log ε) 311 (4.05), 243 (4.16), 211 (4.25) nm; IR (KBr) v_{max} 3429, 2922, 1643, 1606, 1579, 1467, 1298, 1161 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 409.1680 [M – H]⁻ (calcd for C₂₄H₂₅O₆, 409.1651).

The (–)-HRESIMS analysis of tetrandraxanthone G (**GT7**) gave a [M – H] peak at m/z 409.1680, attributing to the molecular formula $C_{24}H_{26}O_6$. The 1D NMR data of **GT7** resembled those of rubraxanthone (**GT15**), with two *meta*-coupled aromatic protons at δ_H 6.28 (1H, d, J = 2.0 Hz, H-4) and 6.25 (1H, d, J = 2.0 Hz, H-2), implying the presence of a 1,3-disubstituted A-ring. Key differences between **GT7** and **GT15** occurred for the resonances of the geranyl side chain at C-8. For **GT7**, 1D NMR data combined with HSQC correlations (Table 3.1 and 3.2) indicated characteristic signals for a terminal methylene unit at δ_H/δ_C 4.31 and 3.84 (each 1H, s, H-10)/109.9, a benzylic methylene group at δ_H/δ_C 2.24 (1H, dd, J = 11.6, 3.6 Hz, H-2)/55.0, and three methylene groups at δ_H/δ_C 2.36 (1H, td, J = 13.2, 5.2 Hz, H-4)/31.2, 1.93 (1H, d, J = 13.6 Hz, H-4)/31.2, 1.59 (1H, m, H-5)/23.4, 1.48 (1H, tt, J = 12.8, 4.0 Hz, H-5)/23.4, 1.66 (1H, m, H-6)/35.2, and 1.29 (1H, m, H-6)/35.2, wherein the methylene group signals were mutually coupled according to the COSY cross-peaks of H-4/H-5' and H-5'/H-6'. The remaining resonances attributed to two methyl groups were observed at δ_{H}/δ_{C} 1.21 (3H, s, H-9)/28.3 and 0.90 (3H, s, H-8)/28.1. The above data, supported by HMBC cross-peaks of H-1' with C-8 (δ_{C} 138.3), H-2' with C-1', C-3' (δ_{C} 149.4), and C-7' (δ_{C} 34.9), H-8' and H-9' with C-2', C-6', and C-7', and H-10' with C-2', C-3', and C-4', indicated a 6,6-dimethyl-2-methylenecyclohexylmethyl unit (Figure 3.4). The presence of this unit was also corroborated by the similarity of the NMR data to those reported in a flavonoid named ugonin J [151]. The ECD spectrum of **GT7** presented no apparent Cotton effects, suggesting its occurrence as a racemate. Compound **GT7** was thus elucidated as a new xanthone bearing a cyclized geranyl unit, as shown in Figure 3.3.

3.3.8. Structural elucidation of compound GT8

Physical and spectroscopic properties of tetrandraxanthone H (GT8): Brown gum; UV (MeOH) λ_{max} (log ε) 309 (4.12), 243 (4.16), 207 (4.38) nm; IR (KBr) v_{max} 3425, 2921, 1640, 1600, 1573, 1466, 1295, 1168 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (–)-HRESIMS m/z 409.1671 [M – H] (calcd for C₂₄H₂₅O₆, 409.1651).

The (–)-HRESIMS data of tetrandraxanthone H (**GT8**) displayed a $[M - H]^{-1}$ peak at m/z 409.1671, assigning to the molecular formula $C_{24}H_{26}O_6$. Hence, this compound was found to be isomeric with **GT7**. Detailed analysis of the 1D NMR data for **GT8** revealed the absence of the methine signal at C-2' and the resonances attributed to the $\Delta^{3',10'}$ alkene unit in **GT7**, which were replaced by resonances of a methyl group at δ_H/δ_C 1.27 (3H, s, H-10')/20.9 and two quaternary sp² carbons at δ_C 135.5 (C-2') and δ_C 126.7 (C-3'). In addition, the signal of the methylene group at C-1' in **GT7** changed to a singlet at $\delta_H 4.32/\delta_C$ 28.4 in **GT8**. The HMBC cross-peaks (Figure

3.4) of H-1' with C-2' and C-3', H-8'/H-9' with C-2', and H-10' with C-2', C-3', and C-4' ($\delta_{\rm C}$ 34.8) indicated that the exocyclic double bond of **GT7** was replaced by an endocyclic double bond in **GT8** as depicted in Figure 3.3.

3.3.9. Structural elucidation of compound GT9

Physical and spectroscopic properties of tetrandraxanthone I (**GT9**): Pale yellow powder; $[\alpha]_D^{20}$ +3.0 (*c* 0.65, MeOH); UV (MeOH) λ_{max} (log ε) 313 (4.22), 245 (4.20), 210 (4.16) nm; IR (KBr) v_{max} 3392, 2920, 1645, 1612, 1579, 1462, 1429, 1296, 1153 cm⁻¹; 1D NMR data (Tables 3.1 and 3.2); and (-)-HRESIMS *m/z* 427.1797 [M - H]⁻ (calcd for C₂₄H₂₇O₇, 427.1757).

The (-)-HRESIMS analysis of tetrandraxanthone I (GT9) provided a $[M - H]^{-1}$ peak at m/z 437.1210, attributing to the molecular formula $C_{24}H_{26}O_6$. The 1D NMR data confirmed the same xanthone moiety as for GT7, except for a modification of the side chain at C-8. The 1D NMR data showed signals of five methylene groups at $\delta_{\rm H}/\delta_{\rm C}$ 3.43 (2H, m, H-1)/22.9, 1.79 and 1.75 (each 1H, m, H-5)/17.5, 1.72 (2H, m, H-2')/45.9, 1.64 and 1.50 (each 1H, m, H-4')/35.4, and 1.50 and 1.40 (each 1H, m, H-6)/37.7, and three methyl groups at $\delta_{\rm H}/\delta_{\rm C}$ 1.34 (3H, s, H-10)/27.5, $\delta_{\rm H}/\delta_{\rm C}$ 1.22 (3H, s, H-8')/32.5 and 1.19 (3H, s, H-9')/29.9. The COSY cross-peak of H-1'/H-2' indicated that the two methylene groups at C-1' and C-2' were coupled. The presence of a dihydropyran ring substituted with gem-dimethyl groups at C-7' and a methyl group at C-3' was suggested by the COSY cross-peaks of H-4'/H-5' and H-5'/H-6', along with HMBC cross-peaks of H-4' with an oxygenated sp³ carbon C-3' (δ_c 74.0), H-8' and H-9' with C-6' and an oxygenated sp³ carbon C-7' ($\delta_{\rm C}$ 71.6), and H-10' with C-3' and C-4' (Figure 3.4). The ring was connected to C-2', as deduced from the HMBC correlation of H-10' with C-2'. The experimental ECD spectrum showed no Cotton effects, indicating that GT9 was obtained as a racemic mixture.

Table 3.1. ¹ F	H NMR spectro	oscopic data ((400 MHz) of (compounds G1	Г 1—GT9 (δ in	ppm, J in Hz)			
position	GT1 [°]	GT2 ^a	GT3 ⁰	GT4 ^b	GT5 ⁶	GT6 ^a	GT7 ^a	GT8 ⁰	GT9 ^b
1									
2		6.21, s	6.20, s	6.29, s	6.29, s	6.24, s	6.25, d (2.0)	6.22, d (2.0)	6.19, d (2.4)
ŝ									
4			ຊີ ຈູ ທ [.] 10L	6			6.28, d (2.0)	6.28, d (2.0)	6.31, d (2.4)
4a			าลง ALO			Ella .			
10a			ุ่ กร NG						
5	6.85, s	6.88, s	6.87, s	6.90, s	6.92, s	6.87, s	6.78, s	6.84, s	6.82, s
6			เหา เหา			Mund .			
7			วิท ม						
80			ได้ ยาส์ VER						
8a			้ เย SIT	2					
6									
9a									
1'	3.45, d (6.8)	6.77, d	6.81, d	3.47, d (7.2)	3.47, d (7.2)	3.52, d (6.4)	3.81°	4.32, s	3.43, m
		(10.0)	(10.0)				3.32, t (11.6)		
2'	5.28, t (6.8) ^c	5.57, d	5.52, d	5.29, t (7.2)	5.27, t (7.2)	5.26, t (6.4)	2.24, dd		1.72, m ^c
		(10.0)	(10.0)				(11.6, 3.6)		

position	GT1 ^a	GT2 ⁰	GT3 ⁰	GT4 ^b	GT5 ^b	GT6 ^a	GT7 ^a	GT8 ⁰	GT9 ^b
3'									
4,	1.76, s	1.47, s	1.77, m	1.97, m	1.65, s	1.75, s	2.36, td	1.92, t (6.0)	1.64, m ^c
							(13.2, 5.2)		
							1.93, d		1.50, m ^c
			ະ ຈຸນ HUL	C.			(13.6)		
5,	1.83, s	1.47, s	2.10, m	2.07, m	1.87, s	1.87, s	1.59, m ^c	1.60, m	1.79, m ^c
			งกร DNG				1.48, tt		1.75, m ^c
			าณ์ม KOF				(12.8, 4.0)		
6'			5.09, t (7.2)	5.01, t (6.8)		11111	1.66, m ^c	1.48, m	1.50, m ^c
			วิท ม				1.29, m ^c		1.40, m
۲,			ยาล VER						
8'			1.66, s	1.52, s	2		0.90, s	1.09, s	1.22, s
,6			1.58, s	1.50, s			1.21, s	1.09, s	1.19, s
10′			1.44, s	1.88, s			4.31, s	1.27, s	1.34, s
							3.84, s		
1"	3.51, d (6.8)	4.09, d (6.0)	4.09, d (6.4)	3.46, m	3.44, m	4.09, d (6.0)			
2"	5.24, t (6.8) ^c	5.25, t (6.0)	5.26, t (6.4)	1.75, m	1.77, m	5.26, t (6.0)			
3"									

position	$GT1^{a}$	$GT2^{a}$	$GT3^{a}$	$GT4^{b}$	$GT5^{b}$	GT6 ⁰	GT7 ^a	GT8 ⁰	GT9 ⁶
4"	2.03, m	2.02, m	1.70, s	1.30, s	1.58, m	1.70, s			
5 "	2.09, m	2.04, m	1.83, s	1.30, s	2.19, m	1.83, s			
6"	5.05, t (6.4)	5.02, t (6.4)			5.18, t (6.8)				
۲"									
8″	1.64, s	1.60, s	า จุฬ HUL	90	1.67, s				
	1.57, s	1.54, s	าล _` AL(1.87, s	E AL			
10"	1.86, s	1.82, s	งกร DNG		1.30, s				
1‴	4.10, d (6.0)		ัณ์ม KOF						
2‴	5.26, t (6.0) ^c		เหา เหา เก ไ			11111			
3‴			ีวิท ปทเ						
ď‴t	1.68, s		ยาล VER						
5‴	1.84, s		ลัย SIT	Ð	2				
0H-1	13.68, s	13.48, s	13.46, s	13.43, s	13.41, s	13.36, s	13.63, s	13.52, s	13.58, s
OH-3	6.32, s					6.07, brs			
9-HO	6.40, brs	6.40, brs	6.37, brs			6.37, brs			
OCH ₃ -7	3.81, s	3.81, s	3.81, s	3.86, s	3.86, s	3.81, s	3.78, s	3.79, s	3.86, s
^a ln CDCl ₃ . ^b ln	acetone-d ₆ . ^c Ov	rlapping signe	IIS.						

position	$GT1^a$	$GT2^{a}$	GT3 ^a	GT4 ^b	$GT5^b$	GT6 ^a	$GT7^{a}$	GT8 ^a	$GT9^b$
1	158.7	163.5	163.5	162.5	162.6	161.8	164.2	164.1	165.1
2	109.0	99.3	99.1	98.4	98.5	98.6	98.3	98.4	98.9
3	160.1	160.1	160.5	162.9	163.0	161.1	162.4	162.4	165.6
4	104.5	100.5	100.3	106.4	106.5	104.4	93.4	93.4	93.9
4a	152.2	151.2	151.1	155.2	155.2	154.0	157.1	157.1	158.1
10a	154.6	154.8	154.8	156.6	156.7	154.8	154.4	154.7	156.5
5	101.7	101.7	101.8	102.8	102.9	101.7	101.4	102.0	102.7
6	155.9	155.8	155.8	157.7	157.7	156.0	156.1	155.9	157.7
7	142.7	143.0	143.0	144.7	144.7	142.9	144.0	143.9	144.8
8	137.2	137.4	137.4	140.1	140.2	137.3	138.3	139.5	140.6
8a	112.2	112.4	112.4	111.9	112.0	112.2	112.6	113.6	112.3
9	182.5	182.2	182.2	183.2	183.3	182.5	182.1	182.4	182.9
9a	103.7	103.9	103.8	104.0	104.1	104.2	104.1	104.2	103.9
1′	21.8	115.3	115.7	22.1	22.3	21.8	26.1	28.4	22.9
2′	122.0	127.0	125.9	123.6	123.6	121.6	55.0	135.5	45.9
3'	134.9	78.1	80.6	135.3	132.6	134.8	149.4	126.7	74.0
4'	26.0	28.4	41.8	40.5	26.0	26.0	31.2	34.8	35.4
5′	18.1	28.4	22.8	27.4	17.9	18.1	23.4	19.6	17.5
6'			124.0	125.1			35.2	41.1	37.7
7'			132.1	131.7			34.9	36.0	71.6
8'			26.0	25.8			28.1	28.8	32.5
9'			17.8	17.7			28.3	28.8	29.9
10'			27.2	16.5			109.9	20.9	27.5
1″	21.9	26.7	26.7	23.3	23.0	26.7			
2″	122.0	123.3	123.3	45.8	43.7	123.3			
3"	137.6	135.8	132.4	70.6	72.5	132.4			

Table 3.2. ¹³C NMR spectroscopic data (100 MHz) of compounds **GT1–GT9** (δ in ppm)

position	GT1 ^a	GT2 ^a	GT3 ^a	GT4 ^b	GT5 ^b	GT6 ^a	GT7 ^a	GT8 ^a	GT9 ^b
4″	39.9	39.8	25.8	29.3	42.8	26.0			
5″	26.6	26.7	18.4	29.3	23.5	18.4			
6"	124.1	124.4			126.4				
7"	131.9	131.4			131.7				
8″	25.8	25.6			26.0				
9″	17.8	17.8			17.9				
10″	16.4	16.7			27.6				
1‴	26.7				1220				
2‴	123.4		100000			5			
3‴	132.2		1	11					
4‴	26.0		_///	600					
5‴	18.4		_///	AQK	8 }				
OCH ₃ -7	62.2	62.2	62.2	61.7	61.8	62.3	62.0	61.5	61.7
^a In CDCl ₃ .	^b In acetor	ne-d ₆ .	1 al		N OF				
		3 14		รณ์แห	าวิทยา	อ๊ย			
		Сни			Unive				



Figure 3.4. Key ${}^{1}H - {}^{1}H$ COSY (blue line) and HMBC (red arrow) correlations of tetrandraxanthones A–I (**GT1–GT9**).

3.3.10. Cytotoxic activities of the isolated compounds

Previous reports have revealed that prenylated and geranylated xanthones isolated from plants, especially in the genus Garcinia, possess potent cytotoxic properties against several cancer cell lines [152-155]. Therefore, the isolated compounds, except for GT12, GT13, GT18, and GT21-GT25, which have been evaluated in this manner previously [13, 21, 137], were evaluated for their cytotoxic activity against HeLa S3 and KB cells. The active compounds (GT5, GT10, GT11, GT26, and GT27) with IC₅₀ values lower than 10 μ M toward these two cancer cell lines were further evaluated against the other three cell lines, comprising Hep G2, HT-29, and MCF-7 cells. The results are shown in Table 3.3. Compound GT26 significantly exhibited cytotoxic activities against five cancer cells with IC₅₀ values in the range of $1.64-3.39 \ \mu$ M. Compound **GT10** and its isomer **GT11** were cytotoxic toward HeLa S3, MCF-7, and KB cells with IC₅₀ values of 4.33–4.90 μ M. Compound GT5 exhibited selective cytotoxicity toward two cancer cells, MCF-7 and KB, with IC_{50} values of 8.80 and 7.06 μ M, whereas **GT27** was cytotoxic toward HeLa S3 and KB, with IC_{50} values of 6.74 and 7.41 μ M. Based on a preliminary structure-activity relationship (SAR) analysis, the hydration of pyran and prenyl units in the linear pyranoxanthones GT28 and GT29 decreased their cytotoxic properties compared to GT26. Among the 1,3,6,7-tetraoxygenated xanthone derivatives with a geranyl side chain at C-8 (GT2, GT5, GT7–GT10, and GT15–GT17), only compound GT10 was cytotoxic against three cancer cell lines (Table 3.3) due to the presence of unmodified geranyl units at C-8 and unmodified prenyl units at C-4.

compound	$IC_{50} \pm SD (\mu M)$				
compound	KB	HeLa S3	MCF-7	Hep G2	HT-29
GT1	16.52 ± 0.37	18.35 ± 0.82	NT	NT	NT
GT2	51.91 ± 1.83	60.95 ± 1.87	NT	NT	NT
GT3	>100	>100	NT	NT	NT
GT4	55.36 ± 6.63	>100	NT	NT	NT
GT5	7.06 ± 0.44	15.43 ± 2.75	8.80 ± 0.65	21.91 ± 0.39	47.39 ± 1.40
GT6	14.28 ± 0.93	19.06 ± 1.74	NT	NT	NT
GT7	43.71 ± 0.26	64.03 ± 2.57	NT	NT	NT
GT8	>100	>100	NT	NT	NT
GT9	47.71 ± 2.74	79.40 ± 1.21	NT	NT	NT
GT10	7.26 ± 0.49	9.03 ± 0.13	6.99 ± 0.84	19.38 ± 0.57	46.07 ± 0.76
GT11	4.90 ± 0.33	5.79 ± 0.48	4.33 ± 0.61	13.87 ± 0.14	36.84 ± 0.99
GT12	NT	NT	NT	NT	NT
GT13	NT	NT	NT	NT	NT
GT14	43.16 ± 5.82	>100	NT	NT	NT
GT15	16.01 ± 0.92	34.40 ± 1.12	NT	NT	NT
GT16	25.52 ± 1.44	73.61 ± 5.79	NT	NT	NT
GT17	13.51 ± 0.85	46.33 ± 3.00	NT	NT	NT
GT18	NT	NT	NT	NT	NT
GT19	62.59 ± 9.74	87.27 ± 9.98	NT	NT	NT
GT20	45.03 ± 2.91	>100	NT	NT	NT
GT21	NT	NT	NT	NT	NT
GT22	NT	NT	NT	NT	NT
GT23	NT	NT	NT	NT	NT
GT24	NT	NT	NT	NT	NT
GT25	NT	NT	NT	NT	NT

Table 3.3. Cytotoxicity data of compounds $GT1-GT31^{a}$

compound	$IC_{50} \pm SD (\mu M)$				
compound	KB	HeLa S3	MCF-7	Hep G2	HT-29
GT27	7.41 ± 0.36	6.74 ± 0.28	13.64 ± 1.29	18.58 ± 0.13	47.17 ± 2.39
GT28	14.84 ± 0.27	16.19 ± 0.53	NT	NT	NT
GT29	12.41 ± 1.39	19.41 ± 1.17	NT	NT	NT
GT30	95.05 ± 3.48	>100	NT	NT	NT
GT31	>100	>100	NT	NT	NT
Doxorubicin ^b	0.01 ± 0.002	0.12 ± 0.134	0.49 ± 0.008	1.43 ± 0.182	0.34 ± 0.039

 a Cytotoxicity was expressed as the mean values of three experiments ± SD; b Doxorubicin was

used as the positive control.



CHULALONGKORN UNIVERSITY

Chapter IV

Phytochemical and biological investigation of the stem bark of Garcinia picrorhiza

4.1. Botanical and chemical aspects of G. picrorhiza



Figure 4.1. The whole plant, branches, and leaves of *Garcinia picrorhiza*.

	A Greeced approximation of
Family	: Clusiaceae
Genus	: Garcinia
Species	: Garcinia picrorhiza
Common name	: Sesoot
Local name	: Kayu Ambong, Laloi Ritek (Sulawesi Island), Daun Limon, Kayu
	Lemon, Sesoot (Maluku Island)

Garcinia picrorhiza Miq. is a woody plant that widely grow in Sulawesi, Maluku, and Papua Islands (Figure 4.1), with its origin in Hitu mountain and Laitimor Island, Maluku region. It is an evergreen tree distributed in lower and upper rainforest from 300 up to 2,100 m elevation and thrives on clayey and black soils. The species can reach heights of 18 meter with a girth up to 85 cm. The flowering and fruiting season start from February to August and September to January, respectively, and its wood is commonly utilized for building construction materials. Local people called this species as "Sesoot" and they usually use a decoction of its roots as energy drinking and supplement [15, 156, 157]. Previous phytochemical investigations on the roots and the stem bark of this plant revealed the existence of several types of secondary metabolites, including 3-hydroxy-isonicotinic acid (4.1), two polyprenylated benzophenones (4.2 and 4.3), and three triterpenoids (4.4–4.6) (Figure 4.2) [156, 158-160].



Chulalongkorn University

In the continuing search for novel bioactive substances in *Garcinia* plants [13, 35, 135, 137], the stem bark of *G. picrorhiza* was investigated leading to the isolation of known metabolite **GP9** and new picrorhizones A–H (**GP1–GP8**), along with three known benzoylphloroglucinols (**GP10–GP12**). Herein, the isolation and structural elucidation of the isolated compounds, as well as an *in vitro* cytotoxic evaluation of five human cancer cell lines using the MTT viability method and anti-inflammatory activity using COX-1 and COX-2 inhibitory assays are described.

4.2. Experimental

4.2.1. General Experimental Procedures

The chromatographic materials and instruments information were the same to those previously reported [137, 138] and as described in Chapter III section 3.2.1.

4.2.2. Plant Material

Garcinia picrorhiza stem bark material was collected from Bogor Botanical Garden, Bogor, Indonesia (6°35′51″ S 106°47′55″ E) in July 2016. The plant material was identified by Dr. Rismita Sari and a voucher specimen (No. VI.A.26) was deposited at Bogor Botanical Garden, Indonesia.

4.2.3. Extraction and Isolation

The air-dried stem bark of *G. picrorhiza* (3.0 kg) was extracted with MeOH (15 L each x 3 days) at room temperature. The solvent was evaporated under reduced pressure to obtain a residue (92.0 g). The crude MeOH extract was suspended in distilled water and partitioned with CH_2Cl_2 and EtOAc to afford two organic extracts. The dried CH_2Cl_2 extract (54.1 g) was subjected to silica gel column chromatography and eluted with hexanes/EtOAc (95:5–0:100) to afford fractions A–R. Fraction D (560.0 mg) was chromatographed on a Sephadex LH-20 column with $CH_2Cl_2/MeOH$ (1:1) to obtain subfractions D1–D4. Subfraction D2 (51.6 mg) was subjected to a Chromatotron that was eluted with hexanes/ CH_2Cl_2 (90:10) to yield **GP3** (9.2 mg) and **GP12** (1.8 mg). Compound **GP9** (5.6 g) was isolated as a major component from fractions G (4.2 g) and H (5.1 g) after recrystallization using acetone/hexanes (1:2). Fractions I (708.3 mg) and J (556.0 mg) were subjected separately to a Sephadex LH-20 column with a $CH_2Cl_2/MeOH$ (1:1) solvent system to give subfractions I1–I4 and J1–J3, respectively. Subfraction I1 (49.2 mg) was purified by a Chromatotron that was eluted with 100% CH_2Cl_2 to give **GP2** (11.5 mg), while subfraction I2 (55.8 mg) was

chromatographed on an RP–C₁₈ silica gel column eluted with 70% CH_3CN/H_2O to afford **GP6** (5.7 mg). Subfraction J1 (10.3 mg) was purified on semipreparative HPLC using 93% MeOH/H₂O to yield **GP10** (4.3 mg) and **GP11** (3.2 mg).

Fraction Q (3.3 g) was subjected to silica gel CC eluted with hexanes/EtOAc (75:25–20:80) to give three subfractions (Q1–Q3). Subfraction Q2 (640.2 mg) was separated on silica gel CC using 30% acetone/hexanes to give subfractions Q2.1 and Q2.2. Subfraction Q2.2 (67.5 mg) was separated on a Chromatotron eluted with $CH_2Cl_2/MeOH$ (15:1) to afford **GP1** (14.6 mg). Subfraction Q3 (61.2 mg) was separated on a Chromatotron with $CH_2Cl_2/MeOH$ (20:1) to afford **GP4** (8.7 mg) and **GP5** (4.2 mg). Fraction R (3.7 g) was subjected to silica gel CC that was eluted with hexanes/EtOAc (70:30–20:80) to give five subfractions (R1–R5). Compound **GP8** (6.0 mg) was purified from subfraction R2 (71.4 mg) using an RP–C₁₈ silica gel column that was eluted with 70% CH_3CN/H_2O , while **GP7** (6.7 mg) was obtained from subfraction R5 (52.6 mg) using Sephadex LH-20 with $CH_2Cl_2/MeOH$ (1:1).

4.2.4. X-ray crystallography of GP9

Light yellow single crystals of **GP9** were recrystallized from a mixture of acetone/hexanes (1:2, v/v) in the orthorhombic space group $P2_12_12_1$ (no. 19) with the following unit cell parameters: a = 9.0232(4), b = 18.4443(9), c = 23.7543(11) Å, V = 3953.3(3) Å³ for Mo-K α radiation; a = 9.0225(3), b = 18.3922(6), c = 23.8517(7) Å, V = 3958.0(2) Å³ for Cu-K α radiation; Z = 4; $D_{calc} = 1.012$ g cm⁻³. Two diffraction data sets were collected from two crystals at 296(2) K on Bruker APEXII and PROSPECTOR CCD area-detector diffractometers (Mo-K α radiation, $\lambda = 0.71073$ Å; Cu-K α radiation, $\lambda = 1.54178$ Å) using the APEX2 Software Suite [161]. The data were processed according to a standard procedure, including integration with SAINT+, multi-scan absorption correction, and scaling by SADABS and then merging by XPREP, implemented in APEX2 Software Suite [161], yielding 7314 and 7201 unique reflections with $R_{int} =$

0.0878 and 0.0571, respectively. Two structures were solved by the intrinsic phasing method with SHELXTL XT [162] and refined anisotropically by full matrix leastsquares on F^2 with SHELXTL XLMP [163]. The disordered solvents without direct interaction with GP9 in large intermolecular interstices were removed using PLATON SQUEEZE procedure [164]. The structure refinement converged to final $R_1 = 0.0612$ for 3657 data and $R_1 = 0.0571$ for 6034 data with $F^2 > 2\sigma(F^2)$ for the respective Mo-K α and Cu-K α radiations. The Flack parameter [165] of -0.02(12) was estimated using copper radiation. Data have been deposited with the Cambridge Crystallographic Data Centre (CCDC nos. 1945943 and 1978321). Copies of these data can be obtained, free of charge, on application the CCDC via to www.ccdc.cam.ac.uk/conts/retrieving.html (or 12 Union Road, Cambridge CB2 1EZ, UK, fax: +44 1223 336033, e-mail: deposit@ccdc.cam.ac.uk).

4.2.5. ECD calculation of GP6 and GP9

To determine the absolute configurations of **GP1–GP9**, ECD calculations using the DFT method were carried out, and the calculated ECD spectra were compared with those derived experimentally. Conformers of **GP6** and **GP9** were optimized by semiempirical PM3 calculation and were fully reoptimized with the DFT method at the B3LYP/6-31G(d) level. Subsequently, the ECD calculations were carried out using the TD-DFT method at the B3LYP/6-31+G(d,p) level with a PCM solvent model (MeOH) and n states = 50. All the calculations were performed with GAUSSIAN 09 [166] on a DELL PowerEdge T430 server. The ECD spectra were generated using SpecDis 1.71 [167] with σ = 0.5.

4.2.6. Cytotoxicity Assay

The *in vitro* cytotoxic evaluation of compounds **GP1–GP12** against the KB, HeLa S3, HT-29, MCF-7, and Hep G2 cancer cell lines was performed using an MTT colorimetric method by incubating cells for 72 h, as described previously [137]. The results are expressed as the mean values of three independent experiments. Doxorubicin was used as the positive control.

4.2.7. Cyclooxygenase-1 and -2 Inhibition Assays

COX-1 and COX-2 inhibitory assays were performed in 96-well plate format using purified PGHS-1 from ram seminal vesicles for COX-1 and human recombinant PGHS-2 for COX-2 (Cayman Chemical Co., Ann Arbor, MI, USA), as reported previously [168]. Briefly, the incubation mixture contained 180 μ L of 0.1 M TRIS/HCl-buffer (pH 8.0), 50 μ M Na₂EDTA (only for COX-2), 18mM L-epinepherine bitartrate, 5 μ M porcine hematin, and COX-1 or COX-2 enzymes (0.2 U/well). The tested compounds (GP1-GP11) dissolved in DMSO with a final concentration of 20 μ M were added and the mixture was pre-incubated for 5 mins at room temperature. After incubation, 10 µL of 5 μ M arachidonic acid was added to the mixture and incubated for 20 mins at 37° C. The reaction was subsequently stopped by adding 10 μ L of 10% (v/v) formic acid. The concentration of PGE₂ generated in the reaction was quantitatively measured by a competitive PGE₂ ELISA kit (Enzo Life Sciences Inc., Farmingdale, NY, USA), according to the manufacturer's protocol and previous procedure [168]. The assays were performed in duplicates and repeated over two independent experiments. Indomethacin (1.25 μ M) was dissolved in EtOH and used as the positive control for COX-1. Celecoxib (8.8 μ M), which was used as the positive control for COX-2, was dissolved in DMSO [169].

4.3. Results and discussion

The CH_2Cl_2 -soluble fraction from a crude MeOH extract of stem bark from *G. picrorhiza* was separated to yield 12 compounds, including eight new benzoylphloroglucinols (**GP1–GP8**) bearing a cyclobutyl-containing side chain and four known analogues (**GP9–GP12**). The structures of the known compounds were identified as garcinopicrobenzophenone (**GP9**) [158], 30-*epi*-cambogin (**GP10**) [170], isoxanthochymol (**GP11**) [171], and (+)-30-*epi*-13,14-didehydroxyisogarcinol (**GP12**) [172] through an analysis of their NMR spectroscopic data and comparison with literature data.



Figure 4.3. Benzoylphloroglucinols (GP1–GP12) from the stem bark of *G. picrorhiza*.

4.3.1. Structural elucidation of compound GP1

Physical and spectroscopic properties of picrorhizone A (*GP1*): Yellow gum; [*α*]_D²⁰ +27 (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} (Δε) 218 (–8.3), 259 (+13.3), 303 (– 5.2), 326 (–2.4), 348 (–1.7) nm; ¹H (400 MHz, methanol-*d*₄) and ¹³C NMR (100 MHz, methanol-*d*₄) spectroscopic data, see Table 4.1; and HRESIMS *m/z* 641.3455 [M + Na]⁺ (calcd for C₃₈H₅₀O₇Na, 641.3454). The molecular formula of picrorhizone A (**GP1**) was assigned as $C_{38}H_{50}O_7$ according to the sodium adduct ion at m/z 641.3455 [M + Na]⁺ in the HRESIMS, which indicated 14 indices of hydrogen deficiency. The ¹H NMR data showed signals for eight methyls at $\delta_{\rm H}$ 0.84–1.72, an oxygenated methylene at $\delta_{\rm H}$ 3.96 (2H, s), two terminal olefinic protons at $\delta_{\rm H}$ 4.52 (1H, s) and 4.74 (1H, s), and two olefinic protons at $\delta_{\rm H}$ 4.91 (1H, overlap) and 5.35 (1H, br t, J= 8.8 Hz). The proton signals for a 1,3,4-trisubstituted benzene ring [$\delta_{\rm H}$ 6.76 (1H, d, J= 8.4 Hz), 7.13 (1H, dd, J= 8.4, 1.6 Hz), and 7.14 (1H, d, J= 1.6 Hz)] were also observed (Table 4.1). The characteristic peaks of a methylene at $\delta_{\rm C}$ 41.8 (C-7), a methine at $\delta_{\rm C}$ 48.0 (C-6), three sp³ quaternary carbons at $\delta_{\rm C}$ 49.0 (C-5), 60.6 (C-8), and 68.9 (C-4), a conjugated carbonyl carbon at $\delta_{\rm C}$ 196.7 (C-10), an unconjugated carbonyl carbon at $\delta_{\rm C}$ 210.6 (C-9), and an enolized 1,3-diketo group at $\delta_{\rm C}$ 195.3 (C-1), 118.4 (C-2), and 193.7 (C-3) were assigned from the combined ¹³C NMR and HSQC data, suggesting that this metabolite was a benzoylphloroglucinol derivative featuring a bicyclo[3.3.1]nonane skeleton [86, 173-175].

Comprehensive analysis of 1D and 2D NMR data established the 2D structure of GP1 (Figure 4.4), which closely resembled garcinopicrobenzophenone (GP9) [158], a major compound of this plant. The only difference between compounds GP1 and GP9 was that the resonance for one of the methyl groups on the prenyl unit in GP9 was replaced by an oxygenated methylene group at $\delta_{\rm H}$ 3.96 (2H, s, H-20)/ $\delta_{\rm C}$ 68.9 in GP1, indicating the oxidation of that methyl group to a hydroxymethyl moiety. The COSY correlation of H₂-17/H-18 and multiple HMBC correlations from H₂-17 to C-3, C-4, C-9, and C-19, H₂-20 to C-18, C-19, and C-21, and H₃-21 to C-18, C-19, and C-20 confirmed a 4-hydroxy-3-methyl-2-butenyl side chain at C-4 in GP1 (Figure 4.4). The NOESY experiment (Figure 4.5) showed ¹H—¹H interactions of H-18/H₂-20 and H-17a/H₃-21, indicating an *E*-configuration of the $\Delta^{18(19)}$ double bond.

position	${\sf GP1}^a$		GP2 ^a		GP3 ^b	
position	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (J in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)
1	195.3		195.3		193.5	
2	118.4		118.5		116.2	
3	193.7		194.6		197.6	
4	68.9		68.9		69.1	
5	49.0		49.5	1122	49.0	
6	48.0	1.49, m	47.7	1.50, m	46.7	1.41, m
7eq	41.8	2.22, d (13.6)	41.8	2.21, d (14.0)	40.8	2.33, d (14.0)
7ax		2.03, dd (13.6, 6.8)	///]]]	2.03, dd (14.0, 6.0)		2.08 ^c
8	60.6		59.6 ^d		58.2	
9	210.6		209.5		208.7	
10	196.7		198.3		196.9	
11	130.5		140.2	······	136.9	
12	117.6	7.14, d (1.6)	116.1	7.00, d (1.6)	129.3	7.57, d (7.2)
13	145.8		158.6		128.0	7.38, t (7.2)
14	152.2		120.7	6.98, dd (7.6, 1.6)	132.9	7.52, t (7.2)
15	115.5	6.76, d (8.4)	130.0	7.19, t (7.6)	128.0	7.38, t (7.2)
16	124.4	7.13, dd (8.4, 1.6)	121.1	6.96, dd (7.6, 1.6)	129.3	7.57, d (7.2)
17a	26.8	2.76, dd (13.2, 8.8)	27.0	2.69, dd (12.8, 8.8)	26.7	2.71, dd (12.4,
						9.6)
17b		2.59, br d (13.2)		2.57, br d (12.8)		2.50, br d (12.4)
18	122.6	5.35, br t (8.8)	120.7	4.99, br d (8.8)	120.3	5.09, br t (9.6)
19	138.4		136.0		135.0	
20	68.9	3.96, s	26.4	1.71, s	26.3	1.82, s
21	14.2	1.72, s	18.3	1.67, s	18.3	1.67, s
22	23.2	1.19, s	23.0	1.19, s	22.7	1.16, s
23	27.4	1.00, s	27.3	1.00, s	27.1	0.97, s

Table 4.1. ¹H (400 MHz) and ¹³C (100 MHz) NMR spectroscopic data of compounds GP1- GP3 (δ in ppm)

	${\sf GP1}^a$		GP2 ^b		GP3 ^b	
position	$\delta_{\scriptscriptstyle ext{C}}$	$\delta_{\scriptscriptstyle extsf{H}}$ (J in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)
24a	30.3	2.16 ^c	30.3	2.13 ^c	29.0	2.14 ^c
24b		2.11 ^c		2.07 ^c		2.05 ^c
25	125.8	4.91 ^c	125.4	4.88 ^c	124.2	4.87, m
26	133.6		133.8		133.0	
27	26.0	1.67, s	26.0	1.67, s	26.0	1.67, s
28	18.2	1.52, s	18.2	1.51, s	18.0	1.50, s
29a	33.4	1.94, d (13.2)	33.4	1.96, d (13.6)	32.3	2.03 ^c
29b		1.64 ^c	Ellon 1	1.62 ^c		1.79 ^c
30	39.8	1.82 ^c	40.0	1.77 ^c	38.4	1.82 ^c
31	43.7		43.8		43.0	
32	51.1	2.25 ^c	51.1	2.27, dd (11.6, 4.8)	50.2	2.24 ^c
33a	30.4	1.75 ^c	30.4	1.75 ^c	29.7	1.76 ^c
33b		1.67 ^c	A DO	1.65 ^c		1.72 ^c
34	30.6	1.21, s	30.7	1.22, s	30.3	1.23, s
35	16.5	0.84, s	16.5	0.85, s	16.2	0.85, s
36	146.6		146.6		146.0	
37a	109.6	4.74, s	109.7	4.74, s	109.2	4.75, s
37b		4.52, s		4.53, s		4.53, s
38	23.4	1.63, s	23.4	1.63, s VERSITY	23.2	1.64, s

^{*a*}Recorded in CD₃OD. ^{*b*}Recorded in CDCl₃. ^{*c*}Overlapping signals. ^{*d*}Data were extracted from the HMBC spectrum.

4.3.2. Structural elucidation of compound GP2 and GP3

Physical and spectroscopic properties of picrorhizone B (GP2): Pale brown gum; $[\alpha]_D^{20}$ +12 (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} (Δε) 218 (–4.2), 261 (+8.2), 305 (–4.4), 326 (–3.1), 348 (–2.3) nm; ¹H (400 MHz, methanol-*d*₄) and ¹³C NMR (100 MHz, methanol-*d*₄) spectroscopic data, see Table 4.1; and HRESIMS *m/z* 585.3532 [M – H] (calcd for C₃₈H₄₉O₅, 585.3580).

Physical and spectroscopic properties of picrorhizone C (GP3): White gum; $[\alpha]_D^{20}$ +15 (*c* 0.10, CHCl₃); ECD (*c* 0.001, MeOH) λ_{max} (Δε) 212 (+20.0), 241 (-3.1), 272 (+8.5), 301 (-15.0) nm; ¹H (400 MHz, CDCl₃) and ¹³C NMR (100 MHz, CDCl₃) spectroscopic data, see Table 4.1; and HRESIMS *m/z* 593.3587 [M + Na]⁺ (calcd for C₃₈H₅₀O₄Na, 593.3607).

The molecular formulas of picrorhizones B (GP2) and C (GP3) were determined to be $C_{38}H_{50}O_5$ and $C_{38}H_{50}O_4$ based on an [M – H]⁻ ion at m/z 585.3532 and an $[M + Na]^+$ ion at m/z 593.3587 in the HRESIMS data, respectively. A comparison of the ¹H and ¹³C NMR data of GP2 and GP3 with those of GP9 showed that the structural differences were restricted to the benzoyl unit. The 1D NMR and HSQC data displayed resonances corresponding to a 1,3-disubsituted benzoyl ring in **GP2** at $\delta_{\rm H}/\delta_{\rm C}$ 6.96 (1H, dd, J= 7.6, 1.6 Hz)/121.1, 6.98 (1H, dd, J= 7.6, 1.6 Hz)/120.7, 7.00 (1H, d, J= 1.6 Hz)/116.1, and 7.19 (1H, t, J= 7.6 Hz)/130.0 (Table 4.1) [176]. The COSY correlations of H-14/H-15 and H-15/H-16 and HMBC cross-peaks from H-12 to C-10 ($\delta_{\rm C}$ 198.3), C-13 ($\delta_{\rm C}$ 158.6), C-14, and C-16, H-14 to C-12, C-13, and C-16, H-15 to C-11 ($\delta_{\rm C}$ 140.2), C-13, and C-16, and H-16 to C-10, C-12, and C-14 confirmed the presence of the ring in GP2 (Figure 4.4). In the ¹H NMR spectrum of GP3 (Table 4.1), the resonances of five aromatic methine protons appeared in the region of $\delta_{\rm H}$ 7.33– 7.56, representing an unsubstituted benzoyl unit [177]. The ${}^{1}H$ - ${}^{1}H$ correlations of the contiguous spin system from H-12 to H-16 in the COSY spectrum and the cross-peaks of H-12/H-16 with C-10, C-12, C-14, and C-16, H-13/H-15 with C-11, C-13, and C-15, and H-14 with C-12 and C-16 in the HMBC spectrum completed the assignment of the unit (Figure 4.4).

4.3.3. Structural elucidation of compound GP4

Physical and spectroscopic properties of picrorhizone D (GP4): Yellow gum; $[\alpha]_D^{20}$ +34 (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} (Δε) 234 (–0.9), 265 (+6.0), 303 (– 5.2), 326 (-1.5), 348 (-1.2) nm; ¹H (400 MHz, methanol- d_4) and ¹³C NMR (100 MHz, methanol- d_4) spectroscopic data, see Table 4.2; and HRESIMS m/z 643.3610 [M + Na]⁺ (calcd for C₃₈H₅₂O₇Na, 643.3611).

Based on the observed $[M + Na]^+$ peak at m/z 643.3610 in the HRESIMS data, the molecular formula of picrorhizone D (**GP4**) was established as C₃₈H₅₂O₇. The ¹H NMR data of **GP4** closely resembled those of **GP9**, except for the lack of terminal olefinic proton signals in **GP4**. Two methyl resonances observed at δ_{H}/δ_{C} 1.03 (s, H₃-37)/29.7 and 1.16 (s, H₃-38)/27.9 in the NMR data of **GP4** (Table 4.2) indicated the hydration of the propylene moiety, which was supported by its HRESIMS data analysis indicating one less index of hydrogen deficiency and one more oxygen atom than **GP9**. This interpretation was confirmed by the HMBC correlation of H₃-37/H₃-38 with C-32 (δ_{C} 54.9) and C-36 (δ_{C} 72.6), suggesting that the two methyl groups were located at the oxygenated C-36 (Figure 4.4).

4.3.4. Structural elucidation of compound GP5

Physical and spectroscopic properties of picrorhizone E (*GP5*): Yellow gum; $[\alpha]_D^{20}$ +5 (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} (Δε) 224 (+7.0), 238 (+1.5), 263 (+8.5), 305 (-14.0), 327 (-1.2), 348 (-1.2) nm; ¹H (400 MHz, methanol-*d*₄) and ¹³C NMR (100 MHz, methanol-*d*₄) spectroscopic data, see Table 4.2; and HRESIMS *m/z* 627.3300 [M + Na]⁺ (calcd for C₃₇H₄₈O₇Na, 627.3298).

Picrorhizone E (**GP5**) was determined to have the molecular formula $C_{37}H_{48}O_7$ based on an [M + Na]⁺ ion at m/z 627.3300 in the HRESIMS data. A comparison of the ¹H and ¹³C NMR data of **GP4** and **GP5** indicated that the two compounds differed only at the C-32 substituent of the cyclobutyl unit. The resonances for a methyl group and an oxygenated carbon (δ_c 72.6) in **GP4** were replaced by a carbonyl carbon resonating at δ_c 211.3 in **GP5** (Table 4.2), which was also supported by its HRESIMS data analysis indicating 16 amu units less than **GP4**. The above data, which was corroborated by HMBC correlations (Figure 4.4) of methine H-32 [$\delta_{\rm H}$ 2.84 (1H, dd, J= 9.4, 7.2 Hz)] to C-36 ($\delta_{\rm C}$ 211.3) and H₃-37 ($\delta_{\rm H}$ 2.02) with C-32 ($\delta_{\rm C}$ 55.8) and C-36, suggested that C-36 was oxidized to a carbonyl carbon.

position	GP4 ^a		GP5 ^a	
position	$\delta_{\scriptscriptstyle C}$	$\delta_{ m H}$ (J in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{ m H}$ (J in Hz)
1	194.8		194.5	2
2	118.4		119.1	
3	193.5		193.7	
4	68.8		68.7	
5	49.3		49.1	
6	47.9	1.49, m	48.1	1.46, m
7eq	41.7	2.21, d (14.4)	41.7	2.18 ^c
7ax		2.02, dd (14.4, 6.4)	N/48/21 ²	1.98, dd (14.0, 6.4)
8	60.4 ^d		60.9 ^d	
9	210.0		211.3	
10	196.4		197.0	
11	130.0		131.4	
12	117.3	7.20, d (1.6)	117.4	7.19, d (1.6)
13	146.2		146.1	
14	152.5		151.9	
15	115.2	6.72, d (8.4)	115.2	6.71, d (8.4)
16	125.0	7.03, dd (8.4, 1.6)	124.6	7.04, dd (8.4, 1.6)
17a	27.1	2.70, dd (12.8, 9.2)	27.2	2.67, dd (12.8, 8.0)
17b		2.54, br d (12.8)		2.54, br d (12.8)
18	121.1	5.01, br t (9.2)	121.6	5.00, br t (8.0)

Table 4.2. ¹H (400 MHz) and ¹³C (100 MHz) NMR spectroscopic data of compounds **GP4** and **GP5** (δ in ppm)

position	GP4 ^a		GP5 ^a	
position	$\delta_{\scriptscriptstyle C}$	$\delta_{ extsf{H}}$ (J in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{ extsf{H}}$ (/ in Hz)
19	135.8		135.1	
20	26.4	1.72, s	26.4	1.70, s
21	18.3	1.69, s	18.3	1.68, s
22	23.1	1.19, s	23.3	1.19, s
23	27.3	0.99, s	27.4	0.97, s
24a	30.2	2.13 ^c	30.3	2.18 ^c
24b		2.09 ^c		2.13 ^c
25	125.6	4.89 ^c	125.9	4.91, t (8.0)
26	133.7		133.4	
27	26.0	1.67, s	26.0	1.65, s
28	18.2	1.52, s	18.2	1.51, s
29a	33.2	1.96, d (12.8)	33.5	1.92, d (12.0)
29b		1.64 ^c	(\$)	1.60 ^c
30	40.7	1.64 ^c	40.3	1.88 ^c
31	44.1		45.8	
32	54.9	1.70 ^c	55.8	2.84, dd (9.4, 7.2)
33a	29.0	1.70 ^c	27.5	1.95 ^c
33b		1.65 ^c		1.71 ^c
34	31.8	1.16, s	30.2	1.34, s
35	17.8	1.13, s	17.2	0.94, s
36	72.6		211.3	
37a	29.7	1.03, s	30.2	2.02, s
37b				
38	27.9	1.16, s		

^aRecorded in CD₃OD. ^bRecorded in CDCl₃. ^cOverlapping signals. ^dData were extracted from HMBC spectrum.

4.3.5. Structural elucidation of compound GP6

Physical and spectroscopic properties of picrorhizone F (**GP6**): Pale yellow gum; $[\alpha]_D^{20}$ +11 (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} ($\Delta \varepsilon$) 219 (+33.1), 265 (-28.7), 306 (+6.0), 332 (+2.8), 344 (+2.6) nm; ¹H (400 MHz, methanol- d_4) and ¹³C NMR (100 MHz, methanol- d_4) spectroscopic data, see Table 4.3; and HRESIMS *m/z* 619.3637 [M + H]⁺ (calcd for C₃₈H₅₁O₇, 619.3635) and 641.3468 [M + Na]⁺ (calcd for C₃₈H₅₀O₇Na, 641.3454).

The molecular formula of picrorhizone F (**GP6**) was determined to be $C_{38}H_{50}O_7$ according to the HRESIMS data at m/z 619.3637 [M + H]⁺ and 641.3468 [M + Na]⁺. The NMR data analysis showed that **GP6** and **GP9** were structurally related. The major difference was that the resonances associated with the prenyl unit in **GP9** were absent in **GP6** and replaced by signals for two methyls at δ_H/δ_C 1.04 (s, H₃-20)/25.4 and 1.08 (s, H₃-21)/25.5, a methylene at δ_H/δ_C 2.25 (dd, J= 13.6, 6.8 Hz, H-17b) and 2.65 (dd, J= 13.6, 10.0 Hz, H-17a)/27.4, an oxygenated methine at δ_H/δ_C 4.61 (dd, J= 10.0, 6.8 Hz, H-18)/94.1, and an oxygenated tertiary carbon at δ_C 71.9 (C-19) (Table 4.3). The shielding of one of carbonyl carbons at δ_C 177.9 (C-3) in **GP6** compared to that of **GP9** (δ_C 193.8) indicated that one oxygen of the enolic system was enolized. The COSY correlation of H₂-17/H-18 and HMBC cross-peaks of H₂-17 with C-3, C-4, C-5, C-9, C-18, and C-19, H-18 with C-3, C-20 and C-21, H₃-20 with C-18 and C-21, and H₃-21 with C-19 and C-20 showed cyclization of the C-4 prenyl unit to C-3 via an ether bridge to form the 2-(2"-hydroxyprop-2"-yl)tetrahydrofuran ring (Figure 4.4) [178].

4.3.6. Structural elucidation of compound GP7

Physical and spectroscopic properties of picrorhizone G (**GP7**): Pale yellow gum; $[\alpha]_D^{20}$ +8 (c 0.10, MeOH); ECD (c 0.001, MeOH) λ_{max} ($\Delta \varepsilon$) 220 (+27.6), 266 (-24.8), 303 (+6.3), 326 (+2.8), 343 (+1.4) nm; ¹H (400 MHz, methanol- d_4) and ¹³C NMR

(100 MHz, methanol- d_4) spectroscopic data, see Table 4.3; and HRESIMS m/z 655.3245 [M + Na]⁺ (calcd for C₃₈H₄₈O₈Na, 655.3247).

The HRESIMS data analysis of picrorhizone G (**GP7**) at m/z 655.3245 [M + Na]⁺ indicated a molecular formula of C₃₈H₄₈O₈ with a hydrogen deficiency index of 15. Its ¹H and ¹³C NMR data were similar to those of **GP6**. The differences involved the tetrahydrofuran ring in which signals for a methine at δ_{H}/δ_{C} 2.56 (m, H-19)/46.1 and a carbonyl carbon at δ_{C} 176.3 (C-21) in **GP7** were observed instead of the resonances for the *gem*-dimethyl group and oxygenated tertiary carbon in **GP6** (Table 4.3). The carbonyl carbon was part of a hydroxycarbonyl group according to its carbon chemical shift and HRESIMS analysis. The above argument, which was supported by the COSY correlations of H₂-17/H-18 and H-18/H-19, and the HMBC correlations of H₂-17 with C-3, C-4, C-9, C-18, and C-19, H-18 with C-20 and C-21, H-19 with C-18, C-20, and C-21, and H₃-20 with C-18, C-19, and C-21, suggested a modified C-18 substituent as shown in Figure 4.4.

4.3.7. Structural elucidation of compound GP8

Physical and spectroscopic properties of picrorhizone H (**GP8**): Pale yellow gum; $[\alpha]_D^{20} - 7$ (*c* 0.10, MeOH); ECD (*c* 0.001, MeOH) λ_{max} ($\Delta \varepsilon$) 226 (+25.2), 259 (-32.8), 303 (-2.5), 320 (+3.7), 350 (+2.9) nm; ¹H (400 MHz, methanol- d_4) and ¹³C NMR (100 MHz, methanol- d_4) spectroscopic data, see Table 4.3; and HRESIMS *m/z* 601.3526 [M + H]⁺ (calcd for C₃₈H₄₉O₆, 601.3529) and 623.3343 [M + Na]⁺ (calcd for C₃₈H₄₈O₆Na, 623.3349).

The molecular formula of picrorhizone H (**GP8**) was deduced to be $C_{38}H_{48}O_6$ according to its $[M + H]^+$ ion at m/z 601.3526 and $[M + Na]^+$ ion at m/z 623.3343 in the HRESIMS, which was indicative of one more index of hydrogen deficiency than in the known metabolite **GP9**. The ¹H and ¹³C NMR spectra were identical to the portions of **GP9** possessing a phloroglucinol skeleton with two prenyl and one 3-

isopropenyl-2,2-dimethylcyclobutylmethyl units. Unlike compound **GP9**, two aromatic singlets at $\delta_{\rm H}$ 6.95 (H-13) and 7.44 (H-16) were identified in the ¹H NMR data of **GP8**. The HMBC spectrum exhibited cross-peaks of H-13 with C-11 ($\delta_{\rm C}$ 118.1) and three oxygenated carbons C-12 ($\delta_{\rm C}$ 147.1), C-14 ($\delta_{\rm C}$ 154.9), and C-15 ($\delta_{\rm C}$ 151.3), as well as H-16 with C-12, C-14, C-15, and carbonyl carbon C-10 ($\delta_{\rm H}$ 174.4), making it assignable to the 1,2,4,5-tetrasubstituted benzoyl ring in **GP8**. The ¹³C NMR data of **GP8** displayed the characteristic signals of an enolized 1,3-diketo group at $\delta_{\rm C}$ 194.9, 119.2, and 178.8. The HMBC correlations from H₂-17 to C-3 ($\delta_{\rm C}$ 178.8), C-4 ($\delta_{\rm C}$ 64.6), and C-9 ($\delta_{\rm C}$ 207.8) and H₂-29 to C-1 ($\delta_{\rm C}$ 194.9), C-7 ($\delta_{\rm C}$ 41.5), C-8 ($\delta_{\rm C}$ 66.2), and C-9 permitted construction of the other linkage between C-3 and C-12 (Figure 4.4) [179, 180].



Figure 4.4. Key COSY (blue line) and HMBC (red arrow) correlations of picrorhizones A–H (GP1–GP8).
n a siti a n	GP6 ^a		GP7 ^a		GP8 ^a	
position	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (J in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (J in Hz)
1	197.8		197.5		194.9	
2	119.3		119.3		119.2	
3	177.9		177.4		178.8	
4	69.0		68.9		64.6	
5	47.9		47.8	1122	49.5	
6	46.9	1.66, m	46.9	1.66, m	47.8	1.51, m
7eq	41.1	2.09, d (14.0)	41.1	2.08, d (13.6)	41.5	2.25, d (14.0)
7ax		2.02, dd (14.0,	////	2.03, dd (13.6, 6.8)		2.03, dd (14.0, 5.6)
		6.4)	//þ§			
8	61.9		62.0		66.2	
9	207.8		207.7		207.8	
10	193.2	y .	192.8		174.4	
11	130.6	0	130.5	All and a start of the start of	118.1	
12	116.7	7.29, d (1.6)	116.5	7.25, d, (1.6)	147.1	
13	146.7	-1311	146.6		104.0	6.95, s
14	152.9	จุฬาลง	152.9	เหาวทยาลย	154.9	
15	115.6	6.75, dd (8.0)	115.8	6.78, d (8.0)	151.3	
16	124.9	7.16, dd (8.0, 1.6)	124.6	7.14, dd (8.0, 1.6)	109.6	7.44, s
17a	27.4	2.65, dd (13.6,	31.0	2.48 ^c	27.6	2.91, dd (13.6, 9.2)
		10.0)				
17b		2.25, dd (13.6,				2.86, d (13.6)
		6.8)				
18	94.1	4.61, dd (10.0,	88.9	4.89 ^c	119.7	4.62 ^c
		6.8)				
19	71.9		46.1	2.56, m	136.1	
20	25.4	1.04, s	13.6	1.03, d (6.8)	26.0	1.42, s

Table 4.3. ¹H (400 MHz) and ¹³C (100 MHz) NMR spectroscopic data of compounds GP6– GP8 (δ in ppm)

position	GP6 ^a		GP7 ^a		GP8 ^a	
position	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)	$\delta_{\scriptscriptstyle C}$	$\delta_{\scriptscriptstyle extsf{H}}$ (/ in Hz)
21	25.5	1.08, s	176.3		18.5	1.73, s
22	24.2	1.25, s	24.2	1.24, s	22.5	1.27, s
23	27.5	1.11, s	27.5	1.10, s	26.9	1.12, s
24a	30.8	2.51, dd (14.0,	30.8	2.53 ^c	30.4	2.03, d (14.0)
		9.2)				
24b		2.42, d (14.0)		2.41, d (14.8)		1.78 ^c
25	126.0	4.88 ^c	125.9	4.88 ^c	124.7	4.90 ^c
26	134.0		134.1	311/2	134.3	
27	26.0	1.67, s	26.0	1.67, s	26.0	1.64, s
28	18.4	1.58, s	18.4	1.59, s	18.0	1.41, s
29a	33.6	1.82, d (13.6)	33.5	1.81, d (14.0)	31.8	1.84 ^c
29b		1.57 ^c		1.58 ^c		1.71 ^c
30	39.7	1.72 ^c	39.6	1.75, dd (10.0, 8.8)	39.3	1.84 ^c
31	43.6	1	43.5		43.5	
32	51.2	2.19 ^c	51.2	2.19, dd (10.0, 8.0)	51.4	2.25 ^c
33a	30.3	1.59 ^c	30.2	1.58 ^c	30.5	1.78 ^c
33b		1.51 ^c		1.53 ^c		1.76 ^c
34	30.5	1.19, s	30.4	1.19, s	30.6	1.18, s
35	16.4	0.79, s HULALO	16.4	0.78, s VERSIT	16.4	0.84, s
36	146.6		146.6		146.6	
37a	109.7	4.70, s	109.7	4.69, s	109.7	4.77, s
37b		4.46, s		4.45, s		4.61, s
38	23.3	1.60, s	23.3	1.60, s	23.4	1.64, s

^aRecorded in CD₃OD. ^bRecorded in CDCl₃. ^cOverlapping signals.

4.3.8. Determining relative and absolute configurations of GP1-GP9

The relative configuration of **GP1** was deduced on the basis of the ${}^{1}H^{-1}H$ coupling constants, carbon chemical shifts, and NOESY experiments. The coupling

constant between H-6 and H-7ax (${}^{3}J_{6,7ax}$ = 6.8 Hz) and chemical shifts at δ_{C} 48.0 (C-6) and 27.4 (C-23) in the ¹³C NMR spectrum of **GP1** favored an axial orientation for the prenyl unit at C-6 (Table 4.1), since the equatorial position of a C-6 substituent showed ${}^{3}J_{6.7ax} \approx 13.0$ Hz and chemical shifts at δ_{C} 40–42 (C-6) and 16–18 (C-23) [174, 178, 181]. The NOESY correlations of H-17b/H₃-23, H₃-23/H-6, H-7ax/H₃-23, and H₃-22/H-24a in GP1 suggested equatorial orientations for CH₂-17 and CH₃-22 and axial orientations for CH3-23 and CH2-24. Although the correlation of H2-29 with its neighboring protons was not observed, CH₂-29 had to be equatorially oriented since the bridged bicyclic ring system of GP1 required that the side chains at C-4 and C-8 be equatorially oriented (Figure 4.5). In addition, the ${}^{1}H-{}^{1}H$ NOESY interactions of H-30/H-32, H-30/H₃-34, and H-32/H₃-34 indicated that methines H-30 and H-32 in the cyclobutyl ring were cofacial (Figure 4.5). The chemical shifts of the gem-dimethyl groups at $\delta_{\rm C}$ 30.6 (C-34) and 16.5 (C-35) supported the same orientation of H-30 and H-32, since the opposite orientation of two methine protons in the cyclobutyl unit was indicated by the presence of the gem-dimethyl group resonances at δ_c 23.6– 25.1, as reported in caloinophyllin A and brasiliensophyllic acids A and B [182, 183].

The relative configurations of **GP2–GP9** in the bicyclo[3.3.1]nonane core resembled **GP1** based on the ${}^{3}J_{6,7ax}$ value, the C-6 and C-23 chemical shifts, and the NOESY data. The β -orientation of H-18 in **GP6** was confirmed by the NOE correlations of H-17b/H-18 and H-18/H₃-22 (Figure 4.5) and by comparing its NMR data with those of nujiangefolin C and thorelione B regarding the dihydrofuran moiety [178, 184]. The orientation of H-18 in **GP7** was also assigned to be the same as **GP6** based on its NOE interactions, except for the relative configuration at C-19 of **GP7**, which remained undetermined.



Figure 4.5. Key NOE correlations (blue dashed line) of compounds GP1 and GP6.

The absolute configurations of GP1 and GP9 were determined by two methods. The absolute configurations were established by X-ray diffraction using Cu- $K\alpha$ radiation combined with ECD data. The calculated ECD spectrum of GP9a was in good agreement with the experimental ECD spectrum of GP9, indicating a (45,65,85,305,32R) absolute configuration of GP9 (Figure 4.6). The effect of opposite configurations of the cyclobutyl ring on the ECD spectra was also assessed. The calculated spectra of GP9a and its diastereomer (45,65,85,30R,325)-GP9b were highly similar, suggesting that the ECD spectra were marginally influenced by different configurations at C-30 and C-32 (Figure C63, Supporting Information), although the optimized structure model GP9a was more energetically favorable than GP9b. The X-ray diffraction data analysis of GP9 using Cu-K α radiation unequivocally confirmed the absolute configuration (Figure 4.6). The (45,65,85,305,32R) absolute configuration of 1 was defined based on the fact that the experimental ECD spectra of GP1 and GP9 were similar. The absolute configurations of GP2-GP5 were determined as (4*S*,6*S*,8*S*,30*S*,32*R*), (4*S*,6*S*,8*S*,30*S*,32*R*), (45,65,85,305,325), (45,65,85,305,325), respectively, based on the similarity of their experimental ECD spectra with those of GP1 and GP9 (Figure C63, Supporting Information).





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The calculated ECD curve of **GP6a** was in a good agreement with the experimental ECD spectrum of **GP6** (Figure 4.6), indicating a (4*R*,6*S*,8*R*,19*S*,30*S*,32*R*) absolute configuration. The (4*R*,6*S*,8*R*,19*S*,30*S*,32*R*) and (4*R*,6*S*,8*R*,30*S*,32*R*) absolute configurations of **GP7** and **GP8**, respectively, were defined by comparison of their experimental ECD data with those of **GP6**. It is worth noting that compounds **GP6**—**GP8** displayed antipodal experimental ECD spectra relative to those of **GP1**—**GP5** and **GP9**, although the corresponding compounds shared the same side chain orientations of the bicyclo[3.3.1]nonane moiety. This finding could imply that the keto-enol tautomerism in the core structure might affect the molecular conformation

and hence the ECD curves (Figure 4.6), as explained by a previous study reporting the opposite ECD spectra of two methylated (+)-guttiferone K derivatives with a different enolized position [174]. The above-mentioned data suggested that **GP1–GP5** and **GP9** mainly exist in C-1 enol and C-3 keto forms, while compounds **GP6–GP8** are present in C-1 keto form and enolized C-3 via ether ring closure.

4.3.9. Cytotoxic and anti-inflammatory activities of the isolated compounds

Several polyprenylated benzoylphloroglucinols from the genus *Garcinia* have been previously reported to have anticancer and anti-inflammatory activities [18, 67, 86, 173, 185, 186]. For example, (–)-garcimultin D, a benzoylphloroglucinol featuring a tricyclo[4.3.1.0^{3,7}]decane core, induces apoptosis in human erythroleukemia cells and G1-phase cell-cycle arrest and inhibits leukemia oncogene expression [67]. Another benzoylphloroglucinol-type compound, garcinol, shows anti-inflammatory effects by inhibiting *i*NOS and COX-2 expressions and NF- κ B and JAK/STAT-1 activation in LPS-activated macrophages [18]. The above evidence prompted us to evaluate the isolated compounds (**GP1–GP12**) for their cytotoxic properties against KB, HeLa S3, HT-29, MCF-7, and Hep G2 cancer cell lines using an MTT viability assa and for anti-inflammatory activity on COX-1 and COX-2 enzymes. Compound **GP12** was not tested for cyclooxygenase assays due to its small sample size.

Based on the cytotoxicity results (Table 4.4), most of the compounds (**GP3**, **GP5**, **GP6**, **GP8**–**GP11**) were cytotoxic against KB cancer cells, with IC₅₀ values ranging from 5.2 to 9.9 μ M. Surprisingly, only compound **GP6** showed cytotoxicity against MCF-7 and Hep G2 cancer cells with IC₅₀ values of 9.3 and 9.4 μ M, respectively, when compared with the other compounds (**GP1**–**GP5**, **GP7**–**GP9**) bearing a similar cyclobutyl moiety. Among compounds **GP1**–**GP5** and **GP9** tested against HeLa S3 cancer cells, compounds **GP1** and **GP9** possessed cytotoxicity with IC₅₀ values of 8.0

and 7.2 μ M, respectively, indicating that the presence of both 3,4-dihydroxybenzoyl moiety and terminal double bond at the cyclobutyl unit may be required to enhance cytotoxicity. Compounds **GP10** and **GP11** were cytotoxic against MCF-7 cancer cells with IC₅₀ values of 5.9 and 7.2 μ M, respectively, while no cytotoxicity was found for **GP12**. These results suggest that the absence of the 3,4-dihydroxybenzoyl moiety may reduce the cytotoxic properties against those cancer cells.

Table 4.4. Cytotoxic activity^a of compounds **GP1**, **GP3**, **GP5**, **GP6**, and **GP8–GP11** after 72 h of treatment

Compound	$IC_{50} \pm SD (\mu M)$						
compound -	КВ	HeLa S3	MCF-7	Hep G2	HT-29		
GP1	>10	8.0 ± 0.9	NT	NT	NT		
GP2	9.9 ± 0.1	>10	>10	>10	>10		
GP5	7.4 ± 0.7	>10	>10	>10	>10		
GP6	5.9 ± 0.4	6.0 ± 0.4	9.3 ± 1.7	9.4 ± 0.8	>10		
GP8	5.2 ± 0.7	6.2 ± 0.1	>10	>10	>10		
GP9	7.0 ± 0.2	7.2 ± 0.4	>โถยาลัย	>10	>10		
GP10	6.4 ± 0.7	>10 GKORN	5.9 ± 0.1	>10	>10		
GP11	5.3 ± 0.1	>10	7.2 ± 0.1	>10	>10		
Doxorubicin ^b	0.02 ± 0.01	0.15 ± 0.02	1.29 ± 0.02	1.00 ± 0.17	0.59 ± 0.03		

^aResults are expressed as the mean values of three experiments \pm SD; the other isolated compounds were inactive (IC₅₀ > 10 μ M); NT: not tested. ^bDoxorubicin was used as the positive control.

The COX-1 and COX-2 assays (Table 4.5) revealed that only compounds **GP6–GP8** displayed selective inhibitory activity (>10% inhibition) towards COX-1 at a concentration of 20 μ M. The highest activity was found for **GP8** (35.2 ± 9.6%

inhibition). None of compounds led greater than 10% COX-2 inhibition at 20 μ M. Preliminary structure-activity relationship studies suggest that the presence of the keto group at C-1 and enolized C-3 in compounds **GP6–GP8** seems to be important for increasing the COX-1 inhibitory activity.

Compound	% inhibition ± SD				
compound	COX-1	COX-2			
GP1	<10	<10			
GP2	<10	<10			
GP3	<10	<10			
GP4	<10	<10			
GP5	<10	<10			
GP6	10.7 ± 8.8	<10			
GP7	11.9 ± 1.8	<10			
GP8	35.2 ± 9.6	<10			
GP9	<10	<10			
GP10	<10 จุฬาล	<10 ณ์มหาวิทยาลัย			
GP11	<10 GHULAL	0<10 korn Universi			

Table 4.5. COX-1 and COX-2 inhibitory activity^a of compounds **GP1–GP11**.

^aThe experiment was performed in duplicates and repeated twice. The final concentration of compounds used in the assay was 20 μ M. Indomethacin (1.25 μ M) and celecoxib (8.8 μ M) were used as positive control with % inhibition of 78.4 ± 4.1 and 83.5 ± 4.8, respectively. NT: not tested.

4.4. Chemotaxonomic study

Polyprenylated benzoylphloroglucinols (PPBPs) are commonly found in specific genera belonging to the Cluciaceae family, including *Garcinia*, *Clusia*, and *Rheedia*. The PPBPs that have been hitherto isolated mostly possess a

bicyclo[3.3.1]nonane backbone connected to the benzoyl unit and some structural modifications occur at the prenyl or geranyl side chains, which attach to the bicyclic ring system [24]. The new compounds, picrorhizones A–H (**GP1–GP8**), are unique PPBPs that share a cyclobutyl moiety derived from a geranyl unit which are rarely found in plants. Garcinopicrobenzophenone and eugeniaphenone were the only PPBPs bearing this moiety reported previously from *G. picrorhiza* and *G. eugeniaefolia* [158, 187], while the PPBPs with ring expansion on the modified geranyl unit (cyclopentyl and cyclohexyl moieties), such as thorelione A and coccinone F, were isolated from *Calophyllum thorelii* and *Moronobea coccinea* [179, 184]. The cytotoxicity results suggest that the PPBP compound class could be used as a lead compounds in cancer research. Previous reports also support the ability of PPBPs to inhibit the growth of various cancer cells through different mechanism of actions, such as cell cycle arrest and ROS generation induction leading to autophagy and apoptosis [24, 67, 185].

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Chapter V Conclusion

A total of 70 secondary metabolites, including 23 previously unreported structures and 47 known analogues, were successfully isolated and characterized from three Indonesian Garcinia species, G. cylindrocarpa, G. tetrandra, and G. picrorhiza. These Garcinia phytochemicals are identified as xanthone and biphenyl derivatives bearing simple oxygenated functional groups and prenyl and geranyl side polyprenylated benzoylphloroglucinols. Interestingly, the new chains and metabolites tetrandraxanthones G-H contain uncommon cyclized geranyl units at C-8 of the xanthone skeleton and the new picrorhizone derivatives possess an unusual cyclobutyl-containing side chain attached to the core structure. Cytotoxic activity study showed that 9-hydroxycalabaxanthone, a pyranoxanthone isolated from G. tetrandra, was significantly active against all five human cancer cell lines (HeLa S3, Hep G2, HT-29, MCF-7, and KB) with IC₅₀ values lower than 3.5 μ M, while 2deprenylrheediaxanthone B from G. cylindrocarpa and picrorhizone F from G. picrorhiza displayed cytotoxic effects against four cancer cells, except HT-29, with IC_{50} values lower than 10 μ M. In addition, picrorhizone H possessed the highest inhibitory effect towards COX-1 (35.2 \pm 9.6% inhibition at 20 μ M).

The outcome of this work is expected to enrich information about chemical diversity of the genus *Garcinia*. It also provides an insight about lead compounds which have promising cytotoxic properties for further chemical modification and pharmacological study to develop anticancer drug candidates in future, not only showing remarkable activity against cancer cells, but also having minor adverse effects in human body.

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Figure A1. ¹H NMR spectrum of cylindroxanthone D (**GC1**) in acetone- d_6



Figure A2. ¹³C NMR spectrum of cylindroxanthone D (GC1) in acetone- d_6



Figure A4. HSQC spectrum of cylindroxanthone D (GC1) in acetone- d_6


Figure A5. HMBC spectrum of cylindroxanthone D (GC1) in acetone- d_6

		Mass	Spectru	ım List Re	eport			
Analysis Info Analysis Name Method Sample Name Comment	D:∖Data\Data Service\180316_neg_GCD-18.d NV_neg_0.3min_profile_1segment_lowNubulizerDrygas(2).m 180316_neg_GCD-18				Acquisition Date Operator Instrument / Ser#	3/16/2018 12:25:31 PM CU. micrOTOF-Q II 10335		
Acquisition Para Source Type Focus Scan Begin Scan End	ameter ESI Not active 50 m/z 1500 m/z	lon Pola Set Capi Set End Set Colli	rity Ilary Plate Offset sion Cell RF	Negative 4000 ∨ -500 ∨ 150.0 ∨pp	Set Nebulize Set Dry Heat Set Dry Gas Set Divert Va	er Ive	0.4 Bar 200 ℃ 4.0 I/min Waste	
Intens x105 1.5 1.0 0.5 0.5	311. 200 	09366 409.16689 400 #(13-17)	623.19050 600	9 800	1000	1200	1400	m/z

Figure A6. HRESIMS spectrum of cylindroxanthone D (GC1) in MeCN



Figure A8. ¹³C NMR spectrum of cylindroxanthone E (**GC2**) in acetone- d_6

100 90 f1 (ppm)

. . . .

160 150

180 170

120 110



Figure A10. HSQC spectrum of cylindroxanthone E (GC2) in acetone- d_6



Figure A11. HMBC spectrum of cylindroxanthone E (GC2) in acetone- d_6

		Mass Spectru	ım List Re	eport			
Analysis Info Analysis Name Method Sample Name Comment):\Data∖Data Service\180316_neg_R8-2.d JV_neg_0.3min_profile_1segment_lowNubulizerDrygas(2).m 80316_neg_R8-2			Acquisition Date Operator Instrument / Ser#	3/16/2018 12:29:17 CU. micrOTOF-Q II 10:	- 12:29:17 PM F-Q II 10335	
Acquisition Para Source Type Focus Scan Begin Scan End	ameter ESI Not active 50 m/z 1500 m/z	Ion Polarity Set Capillary Set End Plate Offset Set Collision Cell RF	Negative 4000 ∨ -500 ∨ 150.0 ∨pp	Set Nebulizer Set Dry Heate Set Dry Gas Set Divert Va	r 0.4 Bar er 200 ℃ 4.0 l/min lve Waste		
Intens ×105 1.0 0.8 0.6 0.4 0.4 0.2 0.0	327.0 200	655.18: 655.18: 400 600	208 L	1000	1200 1400	m/z	

Figure A12. HRESIMS spectrum of cylindroxanthone E (GC2) in MeCN



Figure A14. ¹³C NMR spectrum of cylindroxanthone F (**GC3**) in acetone- d_6



Figure A16. HSQC spectrum of cylindroxanthone F (GC3) in acetone- d_6



Figure A18. HRESIMS spectrum of cylindroxanthone F (GC3) in MeCN



Figure A19. $^1\mathrm{H}$ NMR spectrum of cylindroxanthone G (GC4) in CDCl_3



Figure A20. ¹³C NMR spectrum of cylindroxanthone G (GC4) in CDCl₃



Figure A21. COSY spectrum of cylindroxanthone G (GC4) in $CDCl_3$



Figure A22. HSQC spectrum of cylindroxanthone G (GC4) in $CDCl_3$



Figure A23. HMBC spectrum of cylindroxanthone G (GC4) in $CDCl_3$

		F					
		Mass Spec	ctrum List	Report			
Analysis Info Analysis Name Method Sample Name	OSCUHF590307 MKE_tune_wide_ GCy-2 GCy-2	'001.d _20130204.m		Acquisition Date Operator Instrument	3/7/2016 1:00:09 PM Administrator micrOTOF 72		
Acquisition Par Source Type Scan Range Scan Begin Scan End	ameter ESI n/a 50 m/z 3000 m/z	lon Polarity Capillary Exit Hexapole RF Skimmer 1 Hexapole 1	Positive 100.0 V 150.0 V 45.0 V 24.3 V	Set Correcto Set Pulsar F Set Pulsar F Set Reflecto Set Flight Tu Set Detector	r Fill 50 V rull 337 V Push 337 V r 1300 V ube 9000 V TOF 2295 V		
Intens_ x10 ⁶ 0.8 0.6 0.4 0.2		274 2718	385.1291		+MS, 0.2-0.3min #(14-1		
L	100	······································	· · · · · · · · · · · · · · · · · · ·				

Figure A24. HRESIMS spectrum of cylindroxanthone G (GC4) in CH_2Cl_2



Figure A26. ¹³C NMR spectrum of cylindrobiphenyl A (GC5) in CDCl₃



Figure A28. HSQC spectrum of cylindrobiphenyl A (GC5) in $CDCl_3$



Figure A29. HMBC spectrum of cylindrobiphenyl A (GC5) in CDCl₃



Figure A30. HRESIMS spectrum of cylindrobiphenyl A (GC5) in MeCN



Figure A31. ¹H NMR spectrum of cylindrobiphenyl B (GC6) in CDCl₃



Figure A32. ^{13}C NMR spectrum of cylindrobiphenyl B (**GC6**) in CDCl_3

143



Figure A34. HSQC spectrum of cylindrobiphenyl B (GC6) in $CDCl_3$



Figure A35. HMBC spectrum of cylindrobiphenyl B (GC6) in $CDCl_3$

		SALAR SA	A CONTRACTOR	ħ			
		Mass Spectr	um List Re	eport			
Analysis Info Analysis Name Method Sample Name Comment	D:\Data\Data Service\ NV_neg_0.3min_profi 180316_neg_R5-1-5-2	180316_neg_R5-1-5.c le_1segment_lowNub 2	d ulizerDrygas(2).m	Acquisition Date Operator Instrument / Ser#	3/16/20 CU. micrO ⁻	018 12:41:39 TOF-Q II 103	РМ 35
Acquisition Para Source Type Focus Scan Begin Scan End	ameter ESI Not active 50 m/z 1500 m/z	lon Polarity Set Capillary Set End Plate Offset Set Collision Cell RF	Negative 4000 V -500 V 150.0 Vpp	Set Nebulizer Set Dry Heater Set Dry Gas Set Divert Valve		0.4 Bar 200 ℃ 4.0 I/min Waste	
Intens_ x104 0.8 0.6 0.4 0.4 0.2 0.0	245.08273	583.30876 400 600	800	1000	1200	1400	m/z

Figure A36. HRESIMS spectrum of cylindrobiphenyl B (GC6) in MeCN





Figure B2. ^{13}C NMR spectrum of tetrandraxanthone A (GT1) in CDCl_3



Figure B4. HSQC spectrum of tetrandraxanthone A (GT1) in $CDCl_3$





Figure B6. HRESIMS spectrum of tetrandraxanthone A (GT1) in MeCN



Figure B7. ¹H NMR spectrum of tetrandraxanthone B (GT2) in CDCl₃



Figure B8. 13 C NMR spectrum of tetrandraxanthone B (GT2) in CDCl₃



Figure B10. HSQC spectrum of tetrandraxanthone B (GT2) in $CDCl_3$



Figure B11. HMBC spectrum of tetrandraxanthone B (GT2) in CDCl₃



Figure B12. HRESIMS spectrum of tetrandraxanthone B (GT2) in MeCN



Figure B13. ¹H NMR spectrum of tetrandraxanthone C (GT3) in $CDCl_3$



Figure B14. ^{13}C NMR spectrum of tetrandraxanthone C (GT3) in CDCl_3



Figure B16. HSQC spectrum of tetrandraxanthone C (GT3) in CDCl_3





Figure B18. HRESIMS spectrum of tetrandraxanthone C (GT3) in MeCN



Figure B19. ¹H NMR spectrum of tetrandraxanthone D (GT4) in acetone– d_6



Figure B20. ¹³C NMR spectrum of tetrandraxanthone D (GT4) in acetone– d_6



Figure B22. HSQC spectrum of tetrandraxanthone D (GT4) in acetone– $d_{\rm 6}$



Figure B23. HMBC spectrum of tetrandraxanthone D (GT4) in acetone- d_6



Figure B24. HRESIMS spectrum of tetrandraxanthone D (GT4) in MeCN



Figure B26. ¹³C NMR spectrum of tetrandraxanthone E (GT5) in acetone– d_6



Figure B28. HSQC spectrum of tetrandraxanthone E (GT5) in acetone– d_6



Figure B29. HMBC spectrum of tetrandraxanthone E (GT5) in acetone– d_6



Figure B30. HRESIMS spectrum of tetrandraxanthone E (GT5) in MeCN



Figure B31. ¹H NMR spectrum of tetrandraxanthone F (GT6) in CDCl₃



Figure B32. ^{13}C NMR spectrum of tetrandraxanthone F (**GT6**) in CDCl_3



Figure B34. HSQC spectrum of tetrandraxanthone F (GT6) in \mbox{CDCl}_3



Figure B35. HMBC spectrum of tetrandraxanthone F (GT6) in CDCl₃



Figure B36. HRESIMS spectrum of tetrandraxanthone F (GT6) in MeCN



Figure B37. ¹H NMR spectrum of tetrandraxanthone G (GT7) in $CDCl_3$



Figure B38. ¹³C NMR spectrum of tetrandraxanthone G (GT7) in CDCl₃



Figure B40. HSQC spectrum of tetrandraxanthone G (GT7) in CDCl₃


Figure B41. HMBC spectrum of tetrandraxanthone G (GT7) in $CDCl_3$

Generic Display Report			
Analysis Info Analysis Name Method Sample Name	D:\Data\Data Service\180316_neg_GTH-25.d NV_neg_0.3min_profile_1segment_lowNubulizerDrygas(2).m 180316_neg_GTH-25	Acquisition Date Operator Instrument	3/16/2018 12:23:54 PM CU. micrOTOF-Q II
Comment Intens			-MS, 0.1-0.2min #(5-10)
2.0-	409.16800		
1.5-			
1.0-			
0.5-			
0.0	841.31548 200 400 600 800	1000	1200 1400 m/z

Figure B42. HRESIMS spectrum of tetrandraxanthone G (GT7) in MeCN



Figure B43. ¹H NMR spectrum of tetrandraxanthone H (GT8) in $CDCl_3$



Figure B44. ¹³C NMR spectrum of tetrandraxanthone H (GT8) in CDCl₃





Figure B46. HSQC spectrum of tetrandraxanthone H (GT8) in $CDCl_3$



Figure B47. HMBC spectrum of tetrandraxanthone H (GT8) in CDCl₃



Figure B48. HRESIMS spectrum of tetrandraxanthone H (GT8) in MeCN



Figure B49. ¹H NMR spectrum of tetrandraxanthone I (GT9) in acetone– d_6



Figure B50. ¹³C NMR spectrum of tetrandraxanthone I (**GT9**) in acetone– d_6





Figure B52. HSQC spectrum of tetrandraxanthone I (GT9) in acetone– d_6



Figure B53. HMBC spectrum of tetrandraxanthone I (GT9) in acetone $-d_6$



Figure B54. HRESIMS spectrum of tetrandraxanthone I (GT9) in MeCN



Figure C1. ¹H NMR spectrum of picrorhizone A (GP1) in methanol- d_4



Figure C2. ¹³C NMR spectrum of picrorhizone A (**GP1**) in methanol- d_4



Figure C4. HSQC spectrum of picrorhizone A (GP1) in methanol- d_4



Figure C6a. NOESY spectrum of picrorhizone A (GP1) in methanol- d_4

5.5

5.0

4.5

4.0 3.5 f2 (ppm) 3.0

2.5

2.0

1.5

1.0

7.0

6.5

6.0

- 5.5 - 6.0 - 6.5 - 7.0

0.5



Figure C6b. Expanded NOESY spectrum of picrorhizone A (GP1) in methanol- d_4



Figure C7. HRESIMS spectrum of picrorhizone A (GP1) in methanol



Figure C8. ¹H NMR spectrum of picrorhizone B (GP2) in methanol- d_4



Figure C9. ¹³C NMR spectrum of picrorhizone B (**GP2**) in methanol- d_4



Figure C11. HSQC spectrum of picrorhizone B (GP2) in methanol- d_4



Figure C13. NOESY spectrum of picrorhizone B (GP2) in methanol- d_4



Figure C14. HRESIMS spectrum of picrorhizone B (GP2) in methanol- d_4







Figure C16. ^{13}C NMR spectrum of picrorhizone C (GP3) in CDCl_3



Figure C18. HSQC spectrum of picrorhizone C (GP3) in $CDCl_3$



Figure C20. NOESY spectrum of picrorhizone C (GP3) in $CDCl_3$



Figure C21. HRESIMS spectrum of picrorhizone C (GP3) in MeCN





Figure C22. ¹H NMR spectrum of picrorhizone D (GP4) in methanol- d_4



Figure C23. 13 C NMR spectrum of picrorhizone D (**GP4**) in methanol- d_4



Figure C25. HSQC spectrum of picrorhizone D (GP4) in methanol- d_4



Figure C27. NOESY spectrum of picrorhizone D (GP4) in methanol- d_4



Figure C28. HRESIMS spectrum of picrorhizone D (GP4) in methanol





Figure C29. ¹H NMR spectrum of picrorhizone E (GP5) in methanol- d_4



Figure C30. ¹³C NMR spectrum of picrorhizone E (GP5) in methanol- d_4



Figure C32. HSQC spectrum of picrorhizone E (GP5) in methanol- d_4



Figure C34. NOESY spectrum of picrorhizone E (GP5) in methanol- d_4









Figure C36. ¹H NMR spectrum of picrorhizone F (GP6) in methanol- d_4



Figure C37. ¹³C NMR spectrum of picrorhizone F (**GP6**) in methanol- d_4



Figure C39. HSQC spectrum of picrorhizone F (GP6) in methanol- d_4





Figure C41. NOESY spectrum of picrorhizone F (GP6) in methanol- d_4









Figure C43. ¹H NMR spectrum of picrorhizone G (GP7) in methanol- d_4



Figure C44. ¹³C NMR spectrum of picrorhizone G (GP7) in methanol- d_4



Figure C46. HSQC spectrum of picrorhizone G (GP7) in methanol- d_4



Figure C47. HMBC spectrum of picrorhizone G (GP7) in methanol- d_4



Figure C48. NOESY spectrum of picrorhizone G (GP7) in methanol- d_4









Figure C50. ¹H NMR spectrum of picrorhizone H (**GP8**) in methanol- d_4



Figure C51. ¹³C NMR spectrum of picrorhizone H (**GP8**) in methanol- d_4


Figure C53. HSQC spectrum of picrorhizone H (GP8) in methanol- d_4



Figure C55. NOESY spectrum of picrorhizone H (GP8) in methanol- d_4









Figure C57. ¹H NMR spectrum of garcinopicrobenzophenone (GP9) in methanol- d_4



Figure C58. ¹³C NMR spectrum of garcinopicrobenzophenone (**GP9**) in methanol- d_4



Figure C59. COSY spectrum of garcinopicrobenzophenone (GP9) in methanol- d_4



Figure C60. HSQC spectrum of garcinopicrobenzophenone (GP9) in methanol- d_4



Figure C61. HMBC spectrum of garcinopicrobenzophenone (GP9) in methanol- d_4



Figure C62. NOESY spectrum of garcinopicrobenzophenone (GP9) in methanol- d_4



Figure C63. (a) Calculated (6a, 6b, 9a, 9a enantiomer, 9b) and experimental ECD spectra (1-9) of GP1-GP9 and (b) Optimized structures and molecular energies (E) from ECD calculation using TD-DFT method for diastereomeric models **9a** and **9b**, as well as **6a** and **6b**.

(a)

position	GP9	
	$\delta_{\scriptscriptstyle C}$	$\delta_{ m H}$ (/ in Hz)
1	194.8	
2	118.3	
3	193.8	
4	68.8	
5	49.3	SAM 112 .
6	47.8	1.48, overlap
7eq	41.7	2.20, d (14.0)
7ах		2.01, dd (14.0, 6.8)
8	61.3	
9	209.9	
10	196.3	
11	129.9	A Constant of the second secon
12	117.3	7.20, d (1.6)
13	146.3	
14	152.5	
15	115.2	6.72, d (8.4)
16	125.0	7.03, dd (8.4, 1.6)
17a	27.1	2.70, dd (13.2, 9.2)
17b		2.55, br d (13.2)
18	121.1	5.01, br t (9.2)
19	135.8	
20	26.4	1.72, s
21	18.3	1.69, s
22	23.1	1.19, s
23	27.3	0.99, s
24a	30.2	2.10, overlap
24b		2.03, overlap

Table C1. $^1\mathrm{H}$ (400 MHz) and $^{13}\mathrm{C}$ (100 MHz) NMR spectroscopic data of **GP9** in methanol- d_4

25	125.6	4.88, overlap
26	133.7	
27	26.0	1.67, s
28	18.2	1.52, s
29a	33.4	1.95, d (14.8)
29b		1.61, overlap
30	40.0	1.77, overlap
31	43.8	
32	51.1	2.25, m
33a	30.4	1.74, overlap
33b		1.67, overlap
34	30.7	1.21, s
35	16.5	0.85, s
36	146.6	
37a	109.4	4.74, s
37b		4.53, s
38	23.4	1.63, s

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